Columbia Exhibit 2126 Illumina, Inc. v. The Trustees of Columbia University in the City of New York IPR2018-00291, 318, 322, 385 & 797

Illumina, Inc. vs. The Trustees of Columbia University in the City of New York

Videotaped Deposition of FLOYD ROMESBERG, PH.D. September 19, 2018 Volume I



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1	UNITED STATES PATENT AND I	Page 1 RADEMARK OFFICE	1	UNITED	STATES PATENT ANI	TRADEMARK OFFIC	Page
2	BEFORE THE PATENT TRIAL A	ND APPEAL BOARD	2	BEFORI	THE PATENT TRIAL	AND APPEAL BOARI	C
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4		. IPR2018-00291	4	ILLUMINA, INC		No. IPR2018-00293	
5	Petitioner, v.	U.S. Patent 9,718,852	5	Pet: v.	tioner,	U.S. Patent 9	9,718,85
c		IPR2018-00318				IPR2018-00318	
6	THE TRUSTEES OF COLUMBIA UNIVERSITY IN THE CITY OF	U.S. Patent 9,719,139	6	THE TRUSTEES (UNIVERSITY IN		U.S. Patent 9	9,719,13
7	NEW YORK,	IPR2018-00322	7	NEW YORK,		IPR2018-00322	2
8	Patent Owner.	U.S. Patent 9,708,358	8	Pater	nt Owner.	U.S. Patent 9	9,708,35
9		IPR2018-00385 U.S. Patent 9,725,480	9			IPR2018-00389 U.S. Patent 9	
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11	VIDEOTAPED DEPOSITION OF FLOY	D ROMESBERG, PH.D.	11	Videotaped der	osition of Floyd	Romesberg, Ph D	
12	VOLUME I		12		en on behalf of th	-	
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L4 L5	SEPTEMBER 19,	2018	14	beginning at 9	9:04 a.m. and end	ng at 3:48 p.m.,	on
L6			15	September 19,	2018, before PATH	RICIA Y. SCHULER,	
17			16	Certified Shore	thand Reporter No	o. 11949.	
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	PATRICIA Y. SCHULER		19 20				
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1	APPEARANCES:	-	1	WIT IIIN III C C	I-N-D-E-X		
2	FOR PETITIONER:		3	WITNESS: FLOYD ROMESBE	RG, PH.D.	EAAMII	NATION
3	KNOBBE MARTENS - SAN DI	EGO, CA	4		BY MR. SCHWARTZ		8
4	BY: NATHANAEL R. LUMAN	, ESQ.	5				
5	12790 El Camino Real		7		E-X-H-I-B-I-	T-S	
6	San Diego, California	92130	8	PATENT OWNER'	S Declaration of F	loud Pomosborg	PAGE 12
7	(858) 707-4000		9	EXHIDIC 1012		t of Petition fo:	
8 9	nate.luman@knobbe.com - and -		10		inter Partes Rev Patent No. 9,718		
		CTTON DC	11	Exhibit 1013	International Ap	plication	15
10	KNOBBE MARTENS - WASHIN		12		Published Under	the Patent	
11	BY: WILLIAM R. ZIMMERM	AN, ESQ.	13		Cooperation trea Sequencing"	ty, Title: "DNA	
12	1717 Pennyslvania Avenu	e	14	Exhibit 2007		loyd Romesberg,	16
L3	Suite 900		1.5		Ph.D.		
14	Washington, DC 20006		15	Exhibit 2035	Video Deposition	Transcript of	21
15	bill.zimmerman@knobbe.c	om	16		Floyd Romesberg,	Ph.D. dated July	
16	FOR PATENT OWNER:		17		8, 2014		
17	FITZPATRICK, CELLA, HAR	PER & SCINTO		Exhibit 1015	Copy of US Pater	t No. 5,547,839	25
18	BY: ROBERT S. SCHWARTZ	, ESQ.	18	Dubibit 0000	Widee Demoit!	The second second second	~~
19	BY: ZACHARY L. GARRETT	, ESQ.	19	EXNIBIT 2080	Video Deposition Floyd Romesberg,	-	32
20	1290 Avenue of the Amer				January 14, 2014		
21	New York, New York 101		20	Exhibit 2082	Video Deposition	Transcript of	73
22	Also Present: Dr. Steven Menchen		21	DAULUIU 2002	Floyd Romesberg,	-	13
<u> </u>					April 10, 2014		
	John White, Esq.		22	Exhibit 2083	Substitute Decla	ration of Floyd	75
23	John Murnane, Esq.		23	2	Romesberg, Ph.D.	-	, , ,
24	Videographer:				Patent Owner's M	otion to Amend	
25	Daniel Bermudez		24				

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25 Page 7 ? ? ? Page 7 ? And at the end is John Murnane from Fitzpatrick. ? <td?< td=""> ? ? <td?< t<="" td=""><td></td><td>Exhibit 2057</td><td></td><td>190</td><td></td><td></td></td?<></td?<>		Exhibit 2057		190		
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29.04 a.m.3THE VIDEOGRAPHER: Good morning. We are3THE VIDEOGRAPHER: Good morning. We are4Now on the record. Today's date is September 19,552018, and the time is 9.04 a.m. This is the video6deposition of Floyd Romesberg Ph.D. being taken in47the matter of Illumina Inc. versus Trustees of6Marcus Burch of Illumina.8Columbia University and the City of New York6THE VIDEOGRAPHER: The court reporter is9pending in the United States Patent and Trademark6THE VIDEOGRAPHER: The court reporter is10Office before the Patent Trial and Appeal Board,10FLOYD ROMESBERG, PH.D.,11case numbers IPR2018-00322, IPR2018-00385.10FLOYD ROMESBERG, PH.D.,12IPR2018-00322, IPR2018-00385.12testified as follows:13We are at Knobbe Martens, San Diego,131414EXAMINATION15BY MR. SCHWARTZ:15of Aptus Court Reporting located at 600 West15BY MR. SCHWARTZ:16Broadway, Suite 300, San Diego, California 92101.16Q. Good morning, Dr. Romesberg. As I said,17Will counsel please identify yourselves18and state whon you represent.19MR. SCHWARTZ: I'm an attorney with Fitzpatrick, Cella,19Columbia University in these IPRs.20Would you please state your full name and21a home address for the record.21University, and with me is my colleague Zack22A. Floyd Romesberg, and my home address is	1					Page 8
3THE VIDEOGRAPHER: Good morning. We are4now on the record. Today's date is September 19,52018, and the time is 9:04 a.m. This is the video6deposition of Floyd Romesberg Ph.D. being taken in7the matter of Illumina Inc. versus Trustees of8Columbia University and the City of New York9pending in the United States Patent and Trademark10Office before the Patent Trial and Appeal Board,11case numbers IPR2018-00322, IPR2018-00385.13We are at Knobbe Martens, San Diego,1412790 El Camino Real. My name is Daniel Bermudez15of Aptus Court Reporting located at 600 West16Broadway, Suite 300, San Diego, California 92101.17Will coursel please identify yourselves18and state whom you represent.19MR. SCHWARTZ: My name is Robert10Schwartz. I'm an attorney with Fitzpatrick, Cella,11Harper & Scinto in New York. I represent Columbia12University, and with me is my colleague Zack13Garert also from Fitzpatrick. And sitting next to14Land State youn Muthem and then sitting next to15Broy Que have been deposed before, have				19, 2016		-
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52018, and the time is 9:04 a.m. This is the video5Marcus Burch of Illumina.6deposition of Floyd Romesberg Ph.D. being taken in7THE VIDEOGRAPHER: The court reporter is7the matter of Illumina Inc. versus Trustees of7Patricia Schuler, and she may now swear in the8Columbia University and the City of New York8witness.9pending in the United States Patent and Trademark910Office before the Patent Trial and Appeal Board,10FLOYD ROMESBERG, PH.D.,11case numbers IPR2018-00321, IPR2018-00318,11having been administered an oath, was examined and12IPR2018-00322, IPR2018-00385.12testified as follows:13We are at Knobbe Martens, San Diego,131412790 El Camino Real. My name is Daniel Bermudez14EXAMINATION15of Aptus Court Reporting located at 600 West15BY MR. SCHWARTZ:16Broadway, Suite 300, San Diego, California 92101.16Q. Good morning, Dr. Romesberg. As I said,17Will counsel please identify yourselves17my name is Robert Schwartz, I'm usually referred to18and state whom you represent.18as Bob Schwartz. I am an attorney with Fitzpatrick, Cella,20Schwartz. I'm an attorney with Fitzpatrick, Cella,20Would you please state your full name and21Harper & Scinto in New York. I represent Columbia21a home address for the record.22A. Floyd Romesberg, and my home address is23Garrett also from Fitzpatrick.						
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	24				-	
	25		-			-
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		yu Komesberg, Fil.D.		University in the City of New TOTK
	1	Page 9 A. I have.	1	Page 10 A. Yes.
	2	Q. And the rule that we follow in	2	Q. I am only asking for a yes or no.
	3	depositions is that unless your counsel instructs	3	A. Yes.
	4	you not to answer a question, you should attempt to	4	Q. What are the names of the attorneys you
	5	answer that question to the best of your ability.	5	met with?
	6	You understand that, correct?	6	A. Kerry and Nate.
	7	A. Yes, I do.	7	Q. How many times did you meet with those
	8	Q. And if at any time you don't understand	8	attorneys in preparation for your deposition today?
	9	my question, please let me know. If you do not say	9	A. In person?
	10	otherwise I'll assume that if you answered my	10	
	11	question that you understood that question, okay?	11	A. I didn't count, but I believe it would
	12	A. Yes.	12	
	13	Q. And you've heard this before, but the	13	
	14	other rule we try to follow it's broken,	14	C C
	15	unfortunately, all the time is that we try not	15	
	16	to talk over one because the court reporter it	16	
	17	becomes impossible for the court reporter to take	17	
	18	down questions and answers. So we'll try to talk	18	
	19	one at a time, okay?	19	A. I am going to guess about the same, three
	20	A. I will try.	20	
	21	Q. And the other rule is that we give	21	Q. And when did those meetings take place?
	22	audible answers, so head shaking or nodding is not	22	- · ·
	23	acceptable, okay?	23	•
	24	Did you meet with your attorneys in	24	pretty early on, and then one in the middle maybe a
	25	preparation for your deposition today?	25	
H			<u> </u>	
	1	Page 11 Q. And I quess maybe my question was not	1	Page 12 A. I believe so. I don't. unfortunately.
		Q. And I guess maybe my question was not	1	A. I believe so. I don't, unfortunately,
	1 2 3	Q. And I guess maybe my question was not clear, but I was specifically referring in	1 2 3	A. I believe so. I don't, unfortunately, recognize the numbers, but they are the four
	2	Q. And I guess maybe my question was not clear, but I was specifically referring in preparation for the deposition, so when did the	2	A. I believe so. I don't, unfortunately, recognize the numbers, but they are the four nucleotide cases, yes.
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110	a Romesberg, Ph.D.		University in the City of New York
1	Page 13 BY MR. SCHWARTZ:	1	Page 14 A. To the best of my knowledge, yes.
2	Q. Just look at the end of that declaration.	2	Q. And did you do any of your own literature
3	I just want to confirm that that's your signature.	3	searching independent of the materials that were
4	MR. ZIMMERMAN: Just so we have a clear	4	provided to you by the attorneys?
5	record, we're looking at Exhibit 1012 in the 291	5	A. Yes, I did.
6		6	Q. And are all of those materials listed in
7	BY MR. SCHWARTZ:	7	your reports?
8	Q. Correct. That is what we're looking at	8	A. No, they are not.
9	now.	9	Q. How did you decide what to list and what
10	Is that your signature?	10	
11	A. That is my signature, yes.	11	A. I chose ones that I thought were most
12	Q. Do you recall signing the three expert	12	5
13	reports in these cases as well?	13	articles that I examined.
14	A. Yes.	14	Q. Did you rely on any materials to form the
15	Q. And what did you do to prepare your	15	opinions in any of your reports that are not
16	expert declarations in this case?	16	listed?
17	A. A lot of reading, discussions with my	17	A. Yes, in a general way. For several of
18	lawyers, and a lot of writing and editing.	18	the issues I looked at many papers, many more than
19	Q. Did you receive any materials such as	19	you would have wanted me to list and bring here.
20	scientific references or patents from your	20	And I just wanted to make sure that my
21	attorneys that you used to prepare any of your	21	recollections of the time period were correct, my
22	declarations?	22	impressions were correct. And the ones that I
23	A. I did, yes.	23	
24	Q. And are all of those materials listed in	24	
25	your reports?	25	
	Jean opened		
1	Page 15 provide us any materials that you relied on to form	1	Page 16 MR_SCHWARTZ ⁻ Lam going to put in front
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2	provide us any materials that you relied on to form the opinions that are in your report that are not	2	MR. SCHWARTZ: I am going to put in front of it is marked as Exhibit 1013. It is entitled
2 3	provide us any materials that you relied on to form the opinions that are in your report that are not listed. So I'm making that request on the record,	2 3	MR. SCHWARTZ: I am going to put in front of it is marked as Exhibit 1013. It is entitled "DNA Sequencing."
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	/d Romesberg, Ph.D.		University in the City of New York
1	Page 17 case. It's an expert declaration in IPR2013-00517.	1	Page 18 three sentences in paragraph 36: "One of ordinary
2	Dr. Romesberg, this is one of the expert	2	skill in the art would have understood these
3	declarations that you submitted on behalf of	3	teachings of Tsien as stating that Tsien's
4	Illumina in the IPR challenges made by Intelligent	4	invention was nucleotide sequenced determination
5	Bio-Systems against Illumina; is that correct?	5	using DNA polymerase, and in particular, sequencing
6	A. It appears to be, yes.	6	by synthesis."
7		7	
	Q. And the Illumina patents at issue in	-	That is what you wrote, correct?
8	those IPRs related to SBS, correct?	8	
9	A. Yes.	9	Q. And it is your opinion that a person of
10	Q. Would you please go to Page 14 of your	10	ordinary skill in the art understood that Tsien's
11	declaration. And on Page 14, towards the bottom of	11	invention was SBS, correct?
12	the page there is a header entitled "PCT	12	,
13	publication" W-zero I'm sorry. WO91/00678 to	13	
14	Tsien. Do you see?	14	
15	A. It's 06678.	15	Q. I am referring to a person of ordinary
16	Q. 00678? Thank you.	16	skill as you defined it in your declarations.
17	A. One zero.	17	MR. ZIMMERMAN: And, again, objection.
18	Q. To Tsien, right?	18	Vague as to time.
19	A. Yes.	19	BY MR. SCHWARTZ:
20	Q. That's the same Tsien reference as	20	Q. Well, in your declaration you say as of
21	Exhibit 1013 in the present IPRs, correct?	21	2000, October 2000. So it's your understanding
22	A. Yes, it is.	22	that a person of ordinary skill and again,
23	Q. Please draw your attention to the last	23	whenever I use that term I am referring to how you
24	three sentences in paragraph 36. And in paragraph	24	define it in October 2000 understood that
25	36 you write and this is on Page 15, the last	25	Tsien's invention was SBS?
1	Page 19 A. Yes.	1	Page 20 multiple serial cycles, correct?
1 2		1 2	
	A. Yes.		multiple serial cycles, correct?
2	 A. Yes. Q. And I'll specifically, again, for the 	2	Multiple serial cycles, correct?
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 A. Yes. Q. And I'll specifically, again, for the record refer to Dr. Romesberg's definition of POSA in paragraph 24 of his declarations in these cases. So whenever I use the word person of ordinary skill or if I abbreviate it as POSA, we'll understand we're using your definition as you have it in your declarations as of October 2000, correct? A. Okay. Q. Can you please go back to the Tsien reference, 1013. And again, this is the same Tsien reference that you write about in Exhibit 2007, the declaration that we just looked at, correct? A. Yes. Q. Please turn to Page 16 of Tsien and draw your attention to lines 21 to 23. Are you there? A. I am. Q. Tsien writes "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." Do you see that? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 multiple serial cycles, correct? A. Yes. Q. And earlier I asked you about whether you had been A. I would have to think that. I believe it requires serial reactions. I guess I'm not 100 percent positive that simply reading a single nucleotide, a sequence of one I would have to check to see if that was covered in this patent or not. Q. No, I am just saying Tsien writes. When Tsien writes that "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." A person reading that understood that Tsien was referring to a method that requires multiple serial reactions, correct? A. That is what he wrote, and that is what a POSA would have understood. Q. And that's in relationship to his SBS methods? A. Yes. Q. And earlier I asked you about whether you
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24	 A. Yes. Q. And I'll specifically, again, for the record refer to Dr. Romesberg's definition of POSA in paragraph 24 of his declarations in these cases. So whenever I use the word person of ordinary skill or if I abbreviate it as POSA, we'll understand we're using your definition as you have it in your declarations as of October 2000, correct? A. Okay. Q. Can you please go back to the Tsien reference, 1013. And again, this is the same Tsien reference that you write about in Exhibit 2007, the declaration that we just looked at, correct? A. Yes. Q. Please turn to Page 16 of Tsien and draw your attention to lines 21 to 23. Are you there? A. I am. Q. Tsien writes "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." Do you see that? A. I do. Q. So a reader of Tsien, a POSA reading 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24	 multiple serial cycles, correct? A. Yes. Q. And earlier I asked you about whether you had been A. I would have to think that. I believe it requires serial reactions. I guess I'm not 100 percent positive that simply reading a single nucleotide, a sequence of one I would have to check to see if that was covered in this patent or not. Q. No, I am just saying Tsien writes. When Tsien writes that "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." A person reading that understood that Tsien was referring to a method that requires multiple serial reactions, correct? A. That is what he wrote, and that is what a POSA would have understood. Q. And that's in relationship to his SBS methods? A. Yes. Q. And earlier I asked you about whether you had been deposed in the Illumina versus Intelligent
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 A. Yes. Q. And I'll specifically, again, for the record refer to Dr. Romesberg's definition of POSA in paragraph 24 of his declarations in these cases. So whenever I use the word person of ordinary skill or if I abbreviate it as POSA, we'll understand we're using your definition as you have it in your declarations as of October 2000, correct? A. Okay. Q. Can you please go back to the Tsien reference, 1013. And again, this is the same Tsien reference that you write about in Exhibit 2007, the declaration that we just looked at, correct? A. Yes. Q. Please turn to Page 16 of Tsien and draw your attention to lines 21 to 23. Are you there? A. I am. Q. Tsien writes "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." Do you see that? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 multiple serial cycles, correct? A. Yes. Q. And earlier I asked you about whether you had been A. I would have to think that. I believe it requires serial reactions. I guess I'm not 100 percent positive that simply reading a single nucleotide, a sequence of one I would have to check to see if that was covered in this patent or not. Q. No, I am just saying Tsien writes. When Tsien writes that "One obvious potential shortcoming of the present invention is that it employs a long sequence of serial reactions." A person reading that understood that Tsien was referring to a method that requires multiple serial reactions, correct? A. That is what he wrote, and that is what a POSA would have understood. Q. And that's in relationship to his SBS methods? A. Yes. Q. And earlier I asked you about whether you had been deposed in the Illumina versus Intelligent

	a Romesberg, Ph.D.	University in the City of New Yor
1	Page 21 Do you remember that?	Page 22 1 A. It is.
2	A. I do.	2 Q. Please turn to page 179 of the
3	Q. And you testified you had been deposed,	3 transcript. And please draw your attention to
4	correct?	4 line 2 on page 179. And starting at line 2 the
5	A. Correct.	5 transcript reads "Question: Dr. Romesberg, you
6	Q. And you testified that there were	6 testified that approximately 25 base pairs would be
7	deposition transcripts from those depositions,	7 needed for sequencing by synthesis. Do you recal
8	correct?	8 that."
9	A. Correct.	9 And you answered, "Yes, I do."
10	Q. Did you review any of those deposition	10 That was your testimony, correct?
11	transcripts in preparation for your deposition	11 A. That is what is written, yes.
12	today?	12 Q. It is your opinion that SBS requires
13	A. I looked over them, yes.	13 sequencing about 25 base pairs, correct?
14	Q. I am ready to mark my next exhibit, which	14 MR. ZIMMERMAN: Objection. Vague as to
15	is exhibit let me get this to the court	15 time.
16	reporter.	16 THE WITNESS: I would have to also
17	(Exhibit 2035 was marked for	17 BY MR. SCHWARTZ:
18	identification.)	18 Q. And I'm referring always to the
19	BY MR. SCHWARTZ:	19 October 2000 time frame, as I mentioned earlier,
20	Q. I am putting in front of Dr. Romesberg	20 whenever I use the word POSA.
21	Exhibit 2035, which is entitled video deposition of	
22	Floyd Romesberg Ph.D. dated July 8, 2014. And	22 refresh my memory to see the context of which I was
23	please turn to the last page of Exhibit 235, I	23 describing this, but it would not surprise me that
24	believe it is page 190. And is that your signature	24 I would say that. I would imagine I was talking
25	on page 190?	25 about, as it goes on to explain, the amount of
23	on page 190 :	
-	Dave 00	Page 24
	Page 23	
1	nucleotide reads that he would have to assemble to	1 to, in a practical way, sequence a genome or
2	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case
23	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome.	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable
2	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record,	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer.
23	 nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of 	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ:
2 3 4 5 6	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right?	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your
2 3 4 5	 nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of 	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript
2 3 4 5 6	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right?	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your
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2 3 4 5 6 7 8	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for
2 3 4 5 6 7 8 9	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis.
2 3 4 5 6 7 8 9 10	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right?
2 3 4 5 6 7 8 9 10	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else.	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes.
2 3 4 5 6 7 8 9 10 11 12	 nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give 	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes. Q. So when Tsien states in this patent
2 3 4 5 6 7 8 9 10 11 12 13	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes. Q. So when Tsien states in this patent application that a long sequence of serial
2 3 4 5 6 7 8 9 10 11 12 13 14	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes. Q. So when Tsien states in this patent application that a long sequence of serial reactions is required for his SBS method, that's
2 3 4 5 6 7 8 9 10 11 12 13 14 15	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes. Q. So when Tsien states in this patent application that a long sequence of serial reactions is required for his SBS method, that's
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent?
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay?	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form. 19 MR. SCHWARTZ: You can just object.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a little bit careful to define what you mean by	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form. 19 MR. SCHWARTZ: You can just object. 20 THE WITNESS: Given the caveat that I
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a little bit careful to define what you mean by sequencing by synthesis, but you can read a	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form. 19 MR. SCHWARTZ: You can just object. 20 THE WITNESS: Given the caveat that I 21 just described, but I believe that that statement
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a little bit careful to define what you mean by sequencing by synthesis, but you can read a sequence, and you can read a sequence of one. I	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form. 19 MR. SCHWARTZ: You can just object. 20 THE WITNESS: Given the caveat that I 21 just described, but I believe that that statement 22 is correct.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a little bit careful to define what you mean by sequencing by synthesis, but you can read a sequence, and you can read a sequence of one. I assume that what I was talking about here I'd	 to, in a practical way, sequence a genome or sequence an unknown piece of DNA, in which case something like 25 would seem like a reasonable answer. BY MR. SCHWARTZ: Q. But did you testify just your testimony I'm just reading it from your transcript is approximately 25 base pairs would be needed for sequencing by synthesis. That is what your testimony was, right? A. That's what is written, yes. Q. So when Tsien states in this patent application that a long sequence of serial reactions is required for his SBS method, that's consistent with your opinion that SBS requires sequencing about 25 base pairs, correct? It's MR. ZIMMERMAN: Object as to form. MR. SCHWARTZ: You can just object. THE WITNESS: Given the caveat that I just described, but I believe that that statement a BY MR. SCHWARTZ:
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	nucleotide reads that he would have to assemble to be able to reassemble a genome and uniquely place that sequence within the genome. Q. So, again, just so I have a clean record, it is your opinion that SBS requires sequencing of about 25 base pairs. That's what you said, right? MR. ZIMMERMAN: Same objection. MR. SCHWARTZ: And, again, vague objections are actually not allowed in patent office proceedings, just so you know. You can just say "objection," but you can't say anything else. MR. ZIMMERMAN: I'm just trying to give you an opportunity to correct the question so we are clear as to what time frame you are talking about. MR. SCHWARTZ: I am always talking about the time frame of October 2000, okay, as he defines what a POSA would understand, okay? THE WITNESS: I think you have to be a little bit careful to define what you mean by sequencing by synthesis, but you can read a sequence, and you can read a sequence of one. I	 1 to, in a practical way, sequence a genome or 2 sequence an unknown piece of DNA, in which case 3 something like 25 would seem like a reasonable 4 answer. 5 BY MR. SCHWARTZ: 6 Q. But did you testify just your 7 testimony I'm just reading it from your transcript 8 is approximately 25 base pairs would be needed for 9 sequencing by synthesis. 10 That is what your testimony was, right? 11 A. That's what is written, yes. 12 Q. So when Tsien states in this patent 13 application that a long sequence of serial 14 reactions is required for his SBS method, that's 15 consistent with your opinion that SBS requires 16 sequencing about 25 base pairs, correct? It's 17 consistent? 18 MR. ZIMMERMAN: Object as to form. 19 MR. SCHWARTZ: You can just object. 20 THE WITNESS: Given the caveat that I 21 just described, but I believe that that statement 22 is correct.

	a Romesberg, Ph.D.		University in the City of New York
1	Page 25 Thank you.	1	Page 26 that is long and of practical use, the sort of way
2	(Exhibit 1015 was marked for	2	people would have wanted to use this, then yes.
3	identification.)	3	Q. So practical use of SBS would require
4	BY MR. SCHWARTZ	4	that about 25 base pairs be sequenced?
		-	
5	Q. I put in front of you Exhibit 1015, which	5	A. As long as that practical use required
6	is patent No. 5547839. And again, this is from one	6	determining the sequence of a large piece of DNA
7	of the IPRs, but I believe they are all the same.	7	that you don't know, yes.
8	This is actually Illumina's exhibit, so it was used	8	Q. And that's consistent with your testimony
9	in all of your declarations.	9	that about 25 base pairs is required for SBS?
10	The first named inventor is Dower.	10	A. Yes. The reason I'm qualifying this is
11	Exhibit 1015 is the same Dower patent that you rely	11	because there was a lot of interest, especially at
12	on in your expert declarations in the present IPRs,	12	this time, in determining single nucleotide
13	correct?	13	polymorphisms. And that would be a specific single
14	A. It is.	14	nucleotide. And so it is entirely possible that
15	Q. And the Dower patent relates to SBS,	15	someone could have wanted to use these sorts of
16	correct?	16	things for that, but that would not be the de novo
17	A. That is correct.	17	sequencing that we're talking about.
18	Q. And so, like Tsien's SBS method, Dower's	18	Q. But that would still require about 25
19	SBS method, since it's about SBS, requires	19	base pairs of sequencing to do snips, right?
20	sequencing about 25 base pairs. That's your	20	A. Well, it depends on how you would do it.
21	opinion?	21	If you had a small piece of DNA and you just wanted
22	A. Again, it is a little bit dependent on	22	to know what the nucleotide was at position at
23	what your goal of sequencing by synthesis would be.	23	the position immediately following a primer, then
24	But if it is to determine the unique to uniquely	24	no.
25	determine the sequence of an unknown piece of DNA	25	Q. Let's go back to your deposition
		-	
1	Page 27 transcript that was Exhibit 2035. And go back to	1	Page 28 Q. And that is in relation to your testimony
	transcript that was Exhibit 2035. And go back to	1	Q. And that is in relation to your testimony
1 2 3	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we	2	Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base
2	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you		Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct?
2 3	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be	2 3 4	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct.
2 3 4 5	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your	2 3 4 5	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS
2 3 4 5 6	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right?	2 3 4 5 6	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about
2 3 4 5 6 7	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes.	2 3 4 5 6 7	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct?
2 3 4 5 6 7 8	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about	2 3 4 5 6 7 8	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second
2 3 4 5 6 7 8 9	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you	2 3 4 5 6 7 8 9	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So
2 3 4 5 6 7 8 9 10	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism	2 3 4 5 6 7 8 9 10	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA
2 3 4 5 6 7 8 9 10 11	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what	2 3 4 5 6 7 8 9 10 11	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the
2 3 4 5 6 7 8 9 10 11 12	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then	 2 3 4 5 6 7 8 9 10 11 12 	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is
2 3 4 5 6 7 8 9 10 11 12 13	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was	2 3 4 5 6 7 8 9 10 11 12 13 13	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was
2 3 4 5 6 7 8 9 10 11 12 13 14	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can	2 3 4 5 6 7 8 9 10 11 12 13 14	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would
2 3 4 5 6 7 8 9 10 11 12 13 14 15	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer.	2 3 4 5 6 7 8 9 10 11 12 13 14 15	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application,
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the sequence within the genome to know or not a	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the sequence within the genome to know or not a specific nucleotide is a SNP. So the SNP is	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I think we're going to spend most of our time
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome to know or not a specific nucleotide is a SNP. So the SNP is determined after you assemble the genome, so it	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I think we're going to spend most of our time discussing here, you would need read lengths that
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the sequence within the genome to know or not a specific nucleotide is a SNP. So the SNP is determined after you assemble the genome, so it doesn't change anything."	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I think we're going to spend most of our time discussing here, you would need read lengths that were longer than one.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the sequence within the genome to know or not a specific nucleotide is a SNP. So the SNP is determined after you assemble the genome, so it doesn't change anything." That was your testimony, correct?	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 3	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I think we're going to spend most of our time discussing here, you would need read lengths that were longer than one. Q. So for a practical application of SBS,
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	transcript that was Exhibit 2035. And go back to that page 179 we actually looked at before. And we were looking at the top of the page where you testified that approximately 25 base pairs would be needed for sequencing by synthesis. That was your testimony, right? A. Yes. Q. And would you look just down to about line 12. And you were asked a question "If you were looking for a single nucleotide polymorphism using a sequencing by synthesis method, what minimum read length would you need?" And then there was you started to answer that. There was an objection. And then you were instructed you can answer. And then you said your answer was: "It would not change, because the SNP identification comes from the assembled genome, so you still need to assemble the genome, you still need to place the sequence within the genome to know or not a specific nucleotide is a SNP. So the SNP is determined after you assemble the genome, so it doesn't change anything."	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 Q. And that is in relation to your testimony above that SBS requires sequencing of 25 base pairs, correct? A. That is correct. Q. So even if you are sequencing using SBS for SNPs, your testimony is it still requires about 25 base pairs of sequencing, correct? A. So again, as I tried to explain a second ago, it would depend on how you were doing it. So if you simply had a genome or a large piece of DNA and you wanted to identify all the differences, the SNPs, then what I said in this testimony is correct. The reason I caveated it a minute ago was because I could also imagine cases where one would have a small piece of DNA and one would need to simply know the identity of a nucleotide at a given position, in which case then you would not need a read length of 25. So it really depends on the application, but for the sort of practical applications that I think we're going to spend most of our time discussing here, you would need read lengths that were longer than one.

FIO	yu Romesberg, Fli.D.		University in the City of New Fork
1	Page 29 it is your opinion that you need to sequence about	1	Page 30 Q. And that's a practical application of
2	25 base pairs?	2	SBS, is that what you are saying?
3	MR. ZIMMERMAN: Objection. Asked and	3	A. No. In the particular practical
4	answered.	4	application that I was talking about in this
5	MR. SCHWARTZ: You can't make that	5	testimony, yes. But as I just said, I can imagine
6	objection. It's just objection.	6	there would be other practical applications I
7	BY MR. SCHWARTZ	7	don't know all practical applications that people
8	Q. I'm just looking at your testimony. I	8	might be interested in where you would need a
	just want to make sure that we understand what your	9	read length of one.
	testimony is.	10	Q. What practical application would you need
11	A. What I was referring to here was a	11	a read length of one?
12	certain sort of application. I think it's very	12	A. As I described a minute ago, I'll try to
13	possible that people would have other applications	13	be clear now. You might have a piece of DNA. And
	in mind where they would not need read lengths of	14	you might say I want to know what the nucleotide is
15		15	at position 28. You could design a primer of 27
16	BY MR. SCHWARTZ:	16	length or a primer that terminates immediately
17	Q. But you did testify that for SBS you need	17	before the nucleotide in the template that you're
18	a read length of 25. That is your testimony?	18	interested in, and you could get a read length of
19	A. The context of what I was talking about	19	one and determine what that nucleotide was.
20	at the time, which I hope I made clear, would	20	Now, that was not the applications that I
21	require read lengths of 25, yes.	21	was describing in the context that we're
22	Q. And that would be the same if it was for	22	discussing. But it would seem to be an application
23	SNPs?	23	that someone might be interested in. I'm just
24	A. In the context of what I was describing	24	trying to be clear about what people might have
25	here, which I hope I have made clear.	25	read Tsien or Prober and been interested in doing.
			-
1	Page 31 And lots of people are interested in lots of	1	Page 32 Q. I am going to mark another exhibit.
2	different things.	2	MR. SCHWARTZ: Court reporter, before
3	Q. But you were asked specifically about	3	you hand that over, this is an exhibit that has not
4	SBS, and your testimony was that 25 base pairs	4	yet been marked. Could you please mark this as
5	would be required.	5	Exhibit 2080.
6	A. Everything that we just talked about is	6	(Exhibit 2080 was marked for
7	SBS. SBS in the context of my testimony here was	7	identification.)
8	in a specific situation where you would need a read	8	MR. SCHWARTZ: You might want to put the
9	length of 25. I can certainly imagine other	9	numbers on the bottom in case we have to come back
10	applications of SBS where you would not need that.	10	to it. This is going to be 2080.
11	Q. And the practical applications that you	11	BY MR. SCHWARTZ:
12	need 25 when you're sequencing for purposes of	12	Q. And this is a document entitled "Video
13	determining a gene, how it fits into the genome, is	13	Deposition of Floyd Romesberg Ph.D." dated
14	that something like that?	14	January 14, 2014.
15	A. Yeah. So if you don't know anything	15	Do you see that?
16	about the piece of DNA that you're sequencing, for	16	A. I do.
17	example, and you need to assemble it, you need	17	Q. And this is the transcript of your
18	certain overlap with other sequences to be able to	18	deposition in IPR2013-00128, correct?
19	do that, and so you would need longer read lengths	19	A. It would seem to be, yes.
20	there.	20	Q. Again, the Illumina patents that were
21	Q. Going back to the Tsien patent	21	being challenged in this IPR relate to SBS,
22	application. Tsien doesn't disclose any sequencing	22	correct?
23	data using a 30 allyl nucleotide, correct?	23	A. Yes.
24	A. He does not have data for a three prime	24	Q. Please turn to page 101 of the
25	allyl nucleotide, that is correct.	25	transcript. Please draw your attention to line 6.

	yu Romesberg, Fli.D.		
1	Page 33 And starting at line 6 of the transcript it reads.	1	Page 34 moiety, correct?
2	"Question: And would you agree with me	2	"Answer: Yes, correct.
3	that in 2002 it was known that you could cleave a	3	"Question: And that is a fluorescent
4	linker and a protecting group under identical	4	label attached to the base of the nucleotide,
5	conditions?	5	correct?
6	"Answer: No, I would not agree with	6	"Answer: yes.
7	that.	7	"Question: Okay. And then the next
8	"Question: So if we can turn to start	8	sentence says the tag may be chemically cleaved,
9	Exhibit 1012, that is the Tsien reference. And in	9	either separately or simultaneously with the
10			
11	particular, if can you turn to page 28 of Tsien."	10 11	deblocking step. Do you see that? "Answer: I do.
	Now I'd like to stop here, because I just	12	
12	want to clarify something about the transcript.		"Question: And the deblocking step that
13	In the transcript, Tsien, the T-S-I-E-N	13	Tsien is referring to there is the three prime OH
14	paper is misspelled as C-H-E-N, correct? That is	14	
15	just a misspelling, correct?	15	"Answer: Yes.
16	A. I believe so, yes.	16	"Question: So then would you" strike
17	Q. So what you're actually the question	17	
18	and answer in the section relates to the Tsien,	18	"Question: So then you would agree that
19	T-S-I-E-N, reference that is the same Tsien that we	19	the concept that it was known in the art that you
20	have been looking at in these IPRs, correct?	20	could cleave a base labeled, a cleavable linker
21	A. I believe so, yes.	21	attaching a label to a base in a three prime
22	Q. Let's jump to line 25 of the same page,	22	
23	and starting at line 25, the transcript reads, "So	23	"Answer: Just to be clear, I guess what I
24	you see there that Tsien says the one method	24	5 5
25	involves the use of a tag attached to the base	25	whether with here is whether the suggestion had
	Page 35		Page 36
4		1	
	been made or whether it was known that you could.	1	A. You read it correctly.
2	been made or whether it was known that you could. I see that Tsien is suggesting that, and that was	2	A. You read it correctly.Q. And in your deposition testimony in
2 3	been made or whether it was known that you could. I see that Tsien is suggesting that, and that was known in, and that had been suggested in the	23	 A. You read it correctly. Q. And in your deposition testimony in IPR2013-00128, you specifically pointed out that
2 3 4	been made or whether it was known that you could. I see that Tsien is suggesting that, and that was known in, and that had been suggested in the literature built before. I don't believe that	2 3 4	 A. You read it correctly. Q. And in your deposition testimony in IPR2013-00128, you specifically pointed out that Tsien does not provide any data for simultaneously
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FIO	/d Romesberg, Ph.D.		University in the City of New York
1	Page 37 data that a nucleotide with a 30 allyl protecting	1	Page 38 Q. I'm not asking you about drawing
2	group could be used successfully for SBS?	2	inferences. I'm asking you for any direct data
3	A. No. I think there was data that	3	just the way you testified that you weren't aware
4	suggested that it could be used.	4	of any data that showed cleavable linker on three
5	Q. What data was that?	5	prime protecting group cleavage. Are you aware of
6	A. Reaction showing that you could	6	any data in the context of actually using it for
7	efficiently deprotect the you could efficiently	7	SBS in an SBS reaction, are you aware of any
8	remove the deprotecting group.	8	data that a 3'-O-allyl protecting group on a
9	Q. That wasn't in the context of SBS, was	9	nucleotide that was used for SBS?
10	it?	10	
11	A. No, but you asked about any data.	11	testimony here referred to simultaneously
12	Q. I asked specifically let me ask the	12	
13	question again, okay?	13	
14	Are you aware of any prior art prior to	14	
15	October 2000 that provided any data that a	15	
16	nucleotide with a 3'-O-allyl protecting group could	16	
17	be used for SBS?	17	two things like that.
18	A. I think that the way you asked the	18	-
19	question, the way I would interpret that is that	19	I'm going to again explain that if one was simply
20	any data would include data that a person of	20	asking if an O-allyl could be deprotected from a
21	ordinary skill in the art would look at a	21	nucleotide, it is common, and any person of
22	particular reaction, let's say, and draw an	22	• •
23	impression about the same or similar reaction that	23	-
24	could be used in SBS. And I believe there was data	24	would have looked for similar reactions, and then
25	for that.	25	
	Page 39		Page 40
1	Page 39 you draw an impression about how optimistic or	1	Page 40 A. No, but he did look at steps that would
1		1 2	
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2	you draw an impression about how optimistic or pessimistic you are about whether or not	2	A. No, but he did look at steps that would be involved in that.
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	ya Romesberg, Ph.D.		
1	Page 41 A. That's right. Not for the complete	1	Page 42 data showing simultaneous cleavage, correct?
2	scheme for all the steps of the sequencing by	2	MR. ZIMMERMAN: Objection. Compound.
3	synthesis reaction.	3	BY MR. SCHWARTZ:
4	Q. And without having any date, the best you	4	Q. That was your testimony?
5	could do is speculate at that point?	5	MR. ZIMMERMAN: Objection. Compound.
6	A. Well, this is the problem I'm having. I	6	MR. SCHWARTZ: Again, you can object.
7	disagree with the statement that there would not	7	MR. ZIMMERMAN: The rules allow me to
8	have been any data, because there are multiple	8	succinctly state my objection. Objection.
9	steps. And you have data that shows whether or not	9	Compound.
10	the O-allyl the 3'-O-allyl variant would have	10	BY MR. SCHWARTZ:
11	satisfied individual steps of the overall reaction.	11	Q. Okay.
12	You did not have any data on the overall reaction,	12	A. What I was referring to in that testimony
13	but you did have data on very relevant steps of	13	was a specific step, and no one had ever shown that
14	that overall reaction. And I think that would have	14	step.
15	been I call that data, and I think a person of	15	Q. Okay. And no one has ever shown all of
16	ordinary skill in the art would have viewed that as	16	the steps involved in SBS for the 3'-O-allyl
17	data relevant to how he felt he or she felt	17	nucleotide, correct?
18	about the O-allyl groups.	18	A. That is correct. No one has shown all of
19	Q. But you didn't refer to that kind of	19	the steps for a 3'-O-allyl nucleotide. That is
20	steps data when you were testifying about the	20	correct.
21	simultaneous cleavage. There was data. You were	21	Q. And without that data, you can only
22	asked about data for the individual steps, and you	22	speculate that it would work?
23	said there was no data for the combined cleavage	23	A. I disagree with that.
24	reaction. That it was not enough to just look at	24	Q. But let's go to an Exhibit 2007. It's an
25	the individual data. You were not aware of any	25	exhibit that is already on the table. It's your
	Page 43		Page 44
1	Page 43 expert declaration in IPR2013-00517. Please turn	1	Page 44 be expected to be successful in Tsien's SBS
1 2		1 2	
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	a Romesberg, Ph.D.		University in the City of New York
1	Page 45 Q. So according to Tsien, if any of these	1	Page 46 solvents.)"
	criteria are not met, Tsien says it wouldn't be	2	That's is what you wrote, correct?
	expected to work for his SBS method, correct?	3	A. That is what I wrote.
4	A. Yes.	4	Q. So it is your opinion that most organic
5	Q. Let's go to Tsien's second requirement,	5	solvents, including methanol, were known to
	which is the availability of mild conditions for	6	denature DNA, correct?
	rapid and quantitative deblocking. We'll talk	7	A. Under conditions where they dehydrate the
	about that for a little bit, okay?	8	DNA, yes.
9	A. Yes.	9	Q. Your testimony is most organic solvents,
10	Q. Please go back to your declaration,	10	including methanol, were known to denature DNA.
11	Exhibit 2007, it was your declaration in	11	You cite to the Lee paper, correct?
12	IPR2013-00517. Do you have that?	12	A. Yes. And I was also trying to be clear
13	A. Yes, I do.	13	that it was due to dehydration.
14	Q. Please go to page 29. Please draw your	14	Q. Let's turn to page 23 of your
15	attention to paragraph 55, and please look down	15	
16	about five lines to a sentence that starts with "in	16	second line on the top of the page that starts with
17		17	
18	Do you see that?	18	Do you see that?
19	A. I do.	19	A. Yes, I do.
20	Q. And in paragraph 55 you write: "In fact,	20	Q. And you write: "This reaction was
21	most organic solvents, including tetrahydrofuran,	20	performed in methanol which was known to denature
22	THF, and methanol were known to denature DNA." And	22	DNA, required two hours, and resulted in only
23	you cite the Lee paper of 1981, and you say "(such	23	75 percent to 91 percent yield." You cite to a
24	unwinding of the DNA duplex caused by dehydration	24	paper by Loubinoux. "Further, stannous chloride
25	is likely to be a common phenomenon in organic	25	
20			
1	Page 47 strands." And you cite to a Dantas paper. Then	1	Page 48 That's what you wrote, correct?
	you write: "Any one of these considerations would	2	A. Could you point me to where I wrote that,
	have led one of ordinary skill to have expected	3	please?
4	stannous chloride to be incompatible which ends in	4	Q. We just looked at that before, remember?
5	Ju's requirements for sequencing by synthesis."	5	A. Right. So due to dehydration.
6	That is what you wrote, correct?	6	5
7	-	0	Q. Most organic solvents?
	A. That is what I wrote, correct, ves.	7	Q. Most organic solvents? A. Yes. Due to dehvdration, veah. Thev're
8	 A. That is what I wrote, correct, yes. Q. And, again, the Tsien reference that you 	7	A. Yes. Due to dehydration, yeah. They're
8	Q. And, again, the Tsien reference that you	7 8	A. Yes. Due to dehydration, yeah. They're not damaging the DNA is what I mean. But they're
8 9 10	Q. And, again, the Tsien reference that you referred to in that statement is the same Tsien	7	A. Yes. Due to dehydration, yeah. They're not damaging the DNA is what I mean. But they're incompatible if they're denaturing the DNA, yes.
9 10	Q. And, again, the Tsien reference that you referred to in that statement is the same Tsien reference as Exhibit 2 1013, correct?	7 8 9 10	 A. Yes. Due to dehydration, yeah. They're not damaging the DNA is what I mean. But they're incompatible if they're denaturing the DNA, yes. Q. And what we looked at, you don't say
9	 Q. And, again, the Tsien reference that you referred to in that statement is the same Tsien reference as Exhibit 2 1013, correct? A. It is. 	7 8 9	A. Yes. Due to dehydration, yeah. They're not damaging the DNA is what I mean. But they're incompatible if they're denaturing the DNA, yes.
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FIO	/d Romesberg, Ph.D.		University in the City of New Yorl
1	Page 49 Q. Let's go back to Tsien, please,	1	Page 50 A. Yes.
2	Exhibit 2013. And please turn to page 23. And	2	Q. Can you please turn to page 174. Please
3	please draw your attention to line 27. There is a	2	draw your attention to line 19, and starting at
4	section entitled "deblocking methods." Do you see	4	line 19 the transcript reads:
-	that?	4 5	"Question: And is it reasonable for
5	A. I do.		
6		6	someone looking to develop an SBS method to consult
7	Q. And starting at line 33, Tsien teaches	7	synthetic chemistry papers?
8	that the deblocking method used in his SBS method	8	"Answer: Yes, it is.
9	must, A, proceed rapidly; B, yield a viable	9	"Question: Does that and does it
10	3'-O-allyl O-H function in high yield; and, C, not	10	depend on what is disclosed in those papers?
11	interfere with future enzyme function or denature	11	"Answer: Yes. So I mean, if I were
12	the DNA strand."	12	looking to develop a sequencing by synthesis
13	Those are Tsien's requirements for	13	method, I would certainly consult synthetic
14	deblocking the 3'-O-allyl protecting group, right?	14	chemistry papers that showed things like efficiency
15	A. He lists those requirements, yes.	15	of protecting group cleavage. I would look at the
16	Q. And Tsien expressly requires that the	16	conditions that were explored to see if they were
17	deblocking conditions used to remove the 3'-O-allyl	17	compatible with sequencing by synthesis, and I
18	protecting group does not denature the DNA,	18	would look at yield to see if they were compatible
19	correct?	19	with the high efficiencies required for sequencing
20	A. Yes.	20	by synthesis.
21	Q. Will you please go back to Exhibit 2035,	21	"Question: And if the conditions
22	that was your transcript from your deposition in	22	disclosed in a particular reference were
23	IPR2013-00517.	23	incompatible with sequencing by synthesis, would
24	A. Okay.	24	there be any expectation of success?
25	Q. You have it?	25	"Answer: That would not provide an
<u> </u>	Page 51		Page 52
1	expectation of success."	1	reads:
2	expectation of success." That was your testimony, correct?	1 2	reads: "Question: Would you expect DNA to be
	expectation of success." That was your testimony, correct? A. Yes.		reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous
2	expectation of success." That was your testimony, correct? A. Yes. Q. So your opinion is that it is reasonable	2	reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous pyrimidine sufficient to solubilize
2 3 4 5	expectation of success." That was your testimony, correct? A. Yes. Q. So your opinion is that it is reasonable for someone looking to develop SBS methods to	2 3	reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous
2 3 4 5 6	expectation of success." That was your testimony, correct? A. Yes. Q. So your opinion is that it is reasonable for someone looking to develop SBS methods to consult synthetic chemistry papers, correct?	2 3 4 5 6	reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous pyrimidine sufficient to solubilize triphenylphosphine?" There was an objection.
2 3 4 5 6 7	 expectation of success." That was your testimony, correct? A. Yes. Q. So your opinion is that it is reasonable for someone looking to develop SBS methods to consult synthetic chemistry papers, correct? A. Yes. 	2 3 4 5	reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous pyrimidine sufficient to solubilize triphenylphosphine?" There was an objection. "Answer: That depends on the DNA. Large
2 3 4 5 6	expectation of success." That was your testimony, correct? A. Yes. Q. So your opinion is that it is reasonable for someone looking to develop SBS methods to consult synthetic chemistry papers, correct? A. Yes. Q. And your opinion is, if the condition	2 3 4 5 6	reads: "Question: Would you expect DNA to be denatured at 20 degrees Celsius in aqueous pyrimidine sufficient to solubilize triphenylphosphine?" There was an objection. "Answer: That depends on the DNA. Large pieces of DNA like chromosomal DNA or maybe even
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	Yu Komesberg, Ph.D.		
1	Page 53	1	Page 54 sequencing by synthesis would be more prone to
1	as an especially critical step, but it is a required step that you do denature that you do	2	denaturation.
2			
3	not dissociate the primer from the template.	3	Q. I'm going to ask you to go back to your
4	Q. And because the primers and the templates	4	declaration in IPR2013-00517 that was marked as
5	are short in length, they are more susceptible to	5	Exhibit 2007. Okay?
6	denaturation?	6	A. I am there.
7	A. Shorter than chromosomal double-stranded	7	Q. Please go to page 34. And please draw
8	DNA, so certainly, yes.	8	your attention to paragraph 60. And please look at
9	Q. They're already not very stable. That	9	the third line down in paragraph 60 that starts
10	was your testimony.	10	with "For example, it was known that."
11	A. Well, they're certainly less stable, so	11	Do you see that?
12	they're more prone to lose structure when you	12	A. I do.
13	perturb them by any method; heating or	13	Q. In paragraph 60 you write: "For example,
14	denaturational, whatever.	14	it was known that stringent requirements for
15	Q. And so that's why my question was, so	15	nucleotide analogs with removable 3'-OH terminators
16	they are that's a particularly important	16	presented a formidable obstacle to developing
17	consideration in SBS because the primers and the	17	stable and efficiently deprotected 3'-OH protecting
18	templates are short and only marginally stable,	18	groups for SBS." And you cite to Metzker at 4259.
19	they are even more susceptible to denaturing	19	That is the Metzker paper that you also use in your
20	conditions, correct?	20	
21	A. Yes. I was just a little I was	21	A. I believe that is correct, yes.
22	commenting on that I did not think that the	22	
23	denaturation was any more criterial than any other	23	5
24	requirement. But relative to a chromosome, the	24	
25	conditions that usually would be used for	25	-
			······································
	Page 55	4	Page 56
1	correct?	1	A. Yes, I think I was quoting a paper there.
2	A. Correct.	2	A. Yes, I think I was quoting a paper there. If you want me to describe what I'm trying to say,
2 3	correct? A. Correct. Q. That's what you wrote?	23	A. Yes, I think I was quoting a paper there.If you want me to describe what I'm trying to say,if I can remember the context, any sequencing by
2 3 4	 correct? A. Correct. Q. That's what you wrote? A. That is what I wrote. 	2 3 4	A. Yes, I think I was quoting a paper there. If you want me to describe what I'm trying to say, if I can remember the context, any sequencing by synthesis method stringently requires you simply
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2 3 4 5 6	 correct? A. Correct. Q. That's what you wrote? A. That is what I wrote. Q. It is your opinion that stringent requirements for nucleotide analogs with removable 	2 3 4 5 6	A. Yes, I think I was quoting a paper there. If you want me to describe what I'm trying to say, if I can remember the context, any sequencing by synthesis method stringently requires you simply of the sequencing by synthesis methods that we're describing, you simply have to remove that
2 3 4 5 6 7	 correct? A. Correct. Q. That's what you wrote? A. That is what I wrote. Q. It is your opinion that stringent requirements for nucleotide analogs with removable 3'-OH terminators presented a formidable obstacle 	2 3 4 5 6 7	A. Yes, I think I was quoting a paper there. If you want me to describe what I'm trying to say, if I can remember the context, any sequencing by synthesis method stringently requires you simply of the sequencing by synthesis methods that we're describing, you simply have to remove that protecting group.
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	a Romesberg, Ph.D.		University in the City of New York
4	Page 57	4	Page 58
1	MR. ZIMMERMAN: If now is a good time.	1	page 25. Please draw your attention to the second
2	THE WITNESS: It will only take a second.	2	line on page 25 that starts with "Furthermore,
3	MR. SCHWARTZ: Fine. And, of course,	3	the."
4	you've been through this, you've been to this rodeo	4	Do you see that?
5	before. If you ever need a break, you are entitled	5	A. Yes.
	to have a break.	6	Q. And you write: "Furthermore, the amount
7	THE WITNESS: Yeah, I can't understand	7	of time required to achieve those yields, three
8	why I have to go to the bathroom all the time.	8	hours to multiple days, would not have been
9	THE VIDEOGRAPHER: We are off the record	9	expected to fulfill the rapid deprotection
10	at 10:07 a.m.	10	requirement of Tsien's DNA sequencing method, or
11	(Recess taken.)	11	provide a fast sequencing system capable of the
12	THE VIDEOGRAPHER: We are on the record	12	ultrahigh-throughput in accordance with Ju's DNA
13	at 10:19 a.m.	13	sequencing method."
14	BY MR. SCHWARTZ:	14	That is what you wrote, correct?
15	Q. Dr. Romesberg, would you please go back	15	A. That is correct.
16	to Tsien. That is Exhibit 1013. And please turn	16	Q. It's your opinion that if it takes three
17	to page 23. We may have looked at this before. At	17	hours to achieve high-yield deprotection of the 3'
18	the bottom of the page, that's where Tsien lists	18	protecting group, it would not have been expected
19	the deblocking requirements. And we discussed this	19	to fulfill Tsien's rapid deprojection requirement?
20	earlier, one of Tsien's deblocking requirements is	20	That's what you wrote?
21	rapid deblocking of the 3'-OH protecting group,	21	A. That is what I wrote.
	correct?	22	Q. Let's go back to Exhibit 2035, your
23	A. Proceed rapidly, yes.	23	deposition transcript in this case, in the 00517
24	Q. Please turn back to Exhibit 2007, your	24	case I should say. Please turn to page 97. Please
	declaration in IPR2013-00517. Please go to	25	draw your attention to line 22, and starting at
20		20	araw your attention to fine 22, and starting at
	Page 59	4	Page 60
1	line 22, the transcript reads:	1	but that would at least maybe be in the realm of
2	line 22, the transcript reads: "Question: Does the rapid removal of a	2	but that would at least maybe be in the realm of doable."
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	line 22, the transcript reads: "Question: Does the rapid removal of a 3' protecting group relate to the eventual application of the SBS method?" And there was an objection. "Answer: So you are asking if the efficiency of removal depends required depends on the application? "Question: Sure. That is a better way of asking it. Thank you. "Answer: So that depends. I mean, so that depends. Are you talking about sequencing by synthesis applications? "Question: Yes. "Answer: Sure, but they all have to be efficient. There is a minimum that you are going to have to be. You are going to have to get 30 nucleotide read lengths minimum in a time that is not in a time that's feasible. "Question: What is a feasible time? "Answer: You know, that's a relative	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 but that would at least maybe be in the realm of doable." That is your testimony, correct? A. Yes. Q. Is it fair to say that your opinion is that SBS requires sequencing somewhere between 25 to 30 bases? A. I think it's my opinion that I'm not sure doable is a word, but, yes, I think depending on the specific application, and, again, getting back to the discussion that we had earlier, if you are doing the if your goal is the application that we are describing here, and that is de novo sequencing of unknown sequences of DNA, then, yeah, you would need read lengths of on that order. Q. And then the testimony we just read you testified that a reasonable time for deprotection of the 3'-O-allyl protecting group in SBS is an hour or less. That is what you testified, right? A. Well, I am hemmed and hawed a little bit
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	line 22, the transcript reads: "Question: Does the rapid removal of a 3' protecting group relate to the eventual application of the SBS method?" And there was an objection. "Answer: So you are asking if the efficiency of removal depends required depends on the application? "Question: Sure. That is a better way of asking it. Thank you. "Answer: So that depends. I mean, so that depends. Are you talking about sequencing by synthesis applications? "Question: Yes. "Answer: Sure, but they all have to be efficient. There is a minimum that you are going to have to be. You are going to have to get 30 nucleotide read lengths minimum in a time that is not in a time that's feasible. "Question: What is a feasible time? "Answer: You know, that's a relative term. I mean, I threw out the idea of five hours being not reasonable. And I think that is probably	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 3	but that would at least maybe be in the realm of doable." That is your testimony, correct? A. Yes. Q. Is it fair to say that your opinion is that SBS requires sequencing somewhere between 25 to 30 bases? A. I think it's my opinion that I'm not sure doable is a word, but, yes, I think depending on the specific application, and, again, getting back to the discussion that we had earlier, if you are doing the if your goal is the application that we are describing here, and that is de novo sequencing of unknown sequences of DNA, then, yeah, you would need read lengths of on that order. Q. And then the testimony we just read you testified that a reasonable time for deprotection of the 3'-O-allyl protecting group in SBS is an hour or less. That is what you testified, right? A. Well, I am hemmed and hawed a little bit because I think that is kind of a tough question. If an hour would have been all that is available,

1	Page 61 possible.	1	Page 62 think, you know, today, looking at what we have, I
2	Q. But your testimony was that an hour or	2	think an hour would not be great.
3	less. That was your testimony?	3	Q. But in terms of this testimony, that
4	MR. ZIMMERMAN: Objection.	4	wasn't today. That testimony was in the past. And
5	Mischaracterizes.	5	you were asked about sequencing by synthesis, and
6	THE WITNESS: Let me look at what I said.	6	your testimony was that an hour might not be great,
7	"I threw out the idea of five hours being not	7	but it would be doable. So my question is, it was
8	reasonable, and I think that is probably right.	8	your testimony that in your opinion SBS required
9	And what time is reasonable? I don't know. An	9	that it occur in an hour or less?
10	hour, that would not be great I don't think, but	10	MR. ZIMMERMAN: Objection. Compound.
11	that would at least maybe be in the realm of	11	BY MR. SCHWARTZ:
12	doable."	12	Q. That was your testimony, correct?
13	Yeah, so there's not I don't have an	13	MR. ZIMMERMAN: Objection. Compound.
14	answer, an exact answer to this question. People's	14	THE WITNESS: I don't believe that was my
15	standards would have adapted to what was required,	15	testimony. If it put it this way, if it was, it
16	but clearly faster is better.	16	shouldn't have been. But I don't believe it was.
17	BY MR. SCHWARTZ:	17	Could you direct me to where I said that it would
18	Q. I am just asking about your testimony.	18	have to be an hour or less.
19	And your testimony was that an hour might be	19	BY MR. SCHWARTZ:
20	doable.	20	Q. "What is a reasonable time? I don't
21	A. I certainly believe that an hour might be	21	know. An hour, that would not be great, I don't
22	doable.	22	think, but at least that would be in the realm of
23	Q. But you also said that an hour would not	23	doable."
24	be great.	24	A. If you ask me does it have to be an hour
25	A. Great is kind of a relative term. I	25	or faster to be a great sequencing by synthesis
1	Page 63 method, I think we could discuss what great means,	1	Page 64 BY MR. SCHWARTZ:
2	but I think the point I was making was that an hour	2	Q. But you did testify what is a reasonable
3	is you certainly could have hoped faster, but I	3	time. I don't know, an hour. That would not be
4	don't think an hour would have precluded using it.	4	great.
5	It wouldn't necessarily have been great, but it	5	That was your testimony?
6	still it certainly would have been doable.	6	A. Yes. That is what I wrote, but, again, I
7	I think probably I mean, in reality,	7	don't believe I was I don't believe there's
8	10 hours would have been doable. It would have not	8	anything in that statement that defines an hour as
9	been great. It would have started to be pretty bad	9	some cutoff. I think it is exactly what it says.
10	because you just would not have been able to	10	I didn't know. An hour. I was speculating that
11	desequence your read lengths wouldn't have been	11	wouldn't be great, but doable.
12	limited by time, but would you still be able to do	12	Q. An hour would not be great?
13	it? Yes. Would you be able to do it in the	13	A. But it would be doable.
14	high-throughput way that Ju describes? No.	14	Q. So Tsien teaches that his SBS method
15	So it really comes down to what does	15	requires rapid and quantitative deblocking. A
16	high-throughput mean. And the problem is that	16	POSA and, again, a POSA in terms of the way you
17	scientists use these kinds of terms loosely. So	17	defined it in October 2000 understood that to
18	when I used when I said "I don't know. An hour,	18	mean that the 3' protecting group should be cleaved
19	that would not be great, I don't think, but that	19	quantitatively in what you consider to be around an
20	would at least be in the realm of doable," what I	20	hour?
21	was saying was that an hour, you know, that's	21	A. I would not be comfortable imposing my
22	doable. It certainly could have been better, but I	22	speculation of an hour being doable but not great.
23	don't believe I ever intended to define an hour as	23	I guess you could interpret that as saying my
24	any sort of cutoff.	24	reading of Tsien in a doable but not great way
25	///	25	would have suggested, I don't know, an hour as a

1deprotection time. But I don't think I meant to 2ever state that that was any sort of an absolute 3efficiently incorporate the DNTPs, i.e., deoxy 12ever state that that was any sort of an absolute 3and triphosphates; and, two, the 33ever state that that was any sort of an absolute 4availability of mild conditions for rapid and 4 quantitative deblocking of the 3-0-allyl 9 protecting group, correct?6A. Would be good, I guess. It would 66A. That is what I wrote. 7 G. And Tsien's SBS method requires quantitative deblocking of the 3-0-allyl 9 protecting group, correct?1A. Iagree that one hour would probably be 12 reasonable time was an hour. 140. Tier's SBS method requires quantitative deblocking of the 3'-0-allyl protecting group, 14 correct?16A. Im sory. Of Tsian's method. 15A. Im requires high and 16 deblocking, ves.17IPAcase go back to your declaration in 17 19 paragraph 52 un vargraph 52 un vargraph 52 un paragraph 52 a sthe disulfide cleavage conditions that would 5 provide an a colum 30, lines 28 to 44, stating 19 paragraph 50 kas stating about is the disulfide cleavage. Do you 21Int experimentation of une supportent that 9 procent that you tick here in paragraph 53 4 as the disulfide cleavage conditions that would 5 provide an expe	1	yd Romesberg, Ph.D.		University in the City of New York
2 eversitate that that was any sort of an absolute 2 nuclocities triphosphates; and, two, the 3 requirement intime, you know, 59 minutes okay, 61 3 availability of mild conditions for rapid and 4 Q. Around an hour? 5 A. Would be good, juess. It would 6 6 A. Would be good, juess. It would 6 A. That is what i worte. 7 7 Q. And Tsien's SBS method requires 8 quantitative deblocking." 8 Q. The question was, you asked yourself what 9 protecting group, correct? 10 A. Image that one hourvould probably be 11 in my – please repeat the question. 11 A. Igne that he hour would probably be 11 in my – please repeat the question. 12 reasonable time. And your maker to would probably be 12 Q. Tsien's SBS method requires quantitative 13 oblocking of the 3-O-allyl protecting group. 12 Q. Tsien's SBS method requires quantitative 14 Q. Wore taiking about Tsien. 14 G. Wore taiking about Tsien. 14 14 D. Wore taiking about Tsien. 14 Q. Wore taiking about the set taiking about th			4	
3 requirement in time, you know, 59 minutes okay, 61 3 availability of mild conditions for rapid and 4 4 minutes not. 4 quantitative deblocking." 6 A. Would be good, Iguess. It would 6 A. That is what Jourote. 7 Certainly be doable. 6 A. That is what Jourote. 8 Q. The question was, you asked yourself what is a reasonable time. And your answer to what is a protecting group, correct? 7 A. And Tslor's SBS method requires quantitative deblocking of the 3'-O-allyl protecting group, correct? 11 A. I agree that one hour would probably be transcript in PR2015017. That is schibit 2007. Please transcript in PR20150017. That is schibit 2007. Please transcript in PR2015017. That is schibit 2007. Please transcript in PR20150017. That is schibit 2007. Please transcript in PR20150017. That is schibit 2007. Please transcript in PR20150018. Please transcript in PR20150018. Please transcript in PR20150018. Please transcript reads: 20 2 example That is what you cle here in paragraph 50. Cleavage of a 3'-OH protecting group, what you cle here in paragraph 51. That is schabt would 20 3 as that disulfide		-		
4 quantitative deblocking." 5 C. Around an hour? 5 6 A. Would be good, Iguess. It would 6 7 Cartainly be doable. 7 9 areasonable time. And your answer to what is a 7 10 reasonable time. And your answer to what is a 7 11 A. Ira great hat one hour would probably be 10 12 reasonable for a good, but not great sequencing by 10 13 synthesis implementation of Ju's method. 11 14 Q. We are talking about Tsien. 12 Carse go back to your declaration in 17 IPR2013-00517. That is Exhibit 2007. Please turn 16 deblocking, yes. 19 paragraph 52. In paragraph 52 you write: "Tsien 16 deblocking, yes. 19 analys DNA sequencing methods require high and 18 quantitative efficiency using nucleotides. See 2 3: O-ality locking groups incluide, one, the 21 2 ability of polymerase enzyme to accurately and 12 ant talking about is the disulfide cleavage. Do you 11 11 a. It requires a minute for ead this to get 23 2 abil				
5 Q. Around an hour? 5 That is what you wrote, correct? 6 A. Would be good, iguess. It would 6 A. That is what i wrote. 7 Q. And Tisin's SBS method requires 8 Q. The question was, you asked yourself what is a reasonable time. And your answer to what is a 9 protecting group, correct? 10 reasonable time. And your answer to what is a 9 protecting group, correct? 11 A. I agree that one hour would probably be 11 in my – please repeat the question. 12 reasonable time was an hour. 10 A. That is what you wrote, correct? 13 argare that one hour would probably be 11 in my – please repeat the question. 12 12 reasonable time was an hour. 13 deblocking of the 3'-O-ally! protecting group, correct? 14 Q. We are taking about Tslein. 14 correct? 13 deblocking, yes. 17 IPR2013-00517. That is Exhibit 2007. Please tum 17 Q. You wrote that it "requires high and 13 quantitative efficiency using nucleotides." right? 19 and Jrs DNA sequencing twy 14 quantitative efficiency using nucleotides." right? 19 and Jrs DNA sequencing twy 22 Q. Please go			-	-
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7 Cartality be doable. 7 Q. And Tsien's SBS method requires 8 Q. The question was, you asked yourself what 9 quantitative doablocking of the 3'-O-allyl 9 reasonable time was an hour. 1 A. I gree that one hour would probably be 11 in my – please repeat the question. 11 A. I gree that one hour would probably be 12 Cartain's best sequencing by 12 reasonable time was an hour. 10 A. I'm sorry. I was distracted by an error 11 in my – please repeat the question. 12 A. I'm sorry. I'was distracted by an error 13 was taking about Tsien. 12 A. I'm sorry. I'was distracted by an error 14 Q. We are taking about Tsien. 12 A. I'm sorry. I'was distracted by an error 15 A. I'm sorry. Of Tsien's by and the second the count of the second the second the count of the second			-	• ·
8 Q. The question was, you asked yourself what 8 quantitative deblocking of the 3'-O-allyl 9 is a reasonable time. And your answer to what is a 9 protecting group, correct? 10 reasonable time was now. 10 A. Ims distracted by an error 11 A. Lagree that one hour would probably be 11 in my - please repeat the question. 12 reasonable for a good, but not great sequencing by 12 Q. We are talking about Tsien. 15 A. I'm sory. Of Tsien's method. 15 A. I'm correct? 16 Q. Please go back to your declaration in 16 deblocking yes. 17 IPR2013-00517. That is Exhibit 2007. Please turn 16 deblocking, yes. 19 paragraph 52. In paragraph 52 you write: "Tsien 19 A. That is what I wrote. 20 ad Ju's DNA sequencing methods require high and 20 21 21 quantitative efficiency using nucleotides. See 21 23 that the criteria for the successful use of 24 Please go back to Exhibit 2080, which is 23 group include, one, the 3 as the disulfide cleavage. Doyou 2 "Questint any ou the na maragraph 52 <			-	
9 is a reasonable time. And your answer to what is a 9 protecting group, correct? 10 reasonable time was an hour. 10 A. I'm sorry. I was distracted by an error 11 in your pelsase repeat the question. 11 in your pelsase repeat the question. 12 reasonable for a good, but not great sequencing by synthesis implementation of Ju's method. 12 Q. Tsion's SBS method requires quantitative disclosing, your and the sequencing and the sequencing by synthesis implementation of Ju's method. 12 Q. Tsion's SBS method requires high and deblocking, yes. 17 IPR2013-00517. That is Exhibit 2007. Please turn to page 27, and please draw your attention to and Ju's DNA sequencing methods require high and 21 13 quantitative efficiency using nucleotides. See 21 12 example Tsien at column 20, lines 28 to 34, stating 24 20 Please go back to Exhibit 2080, which is 23 23 sho transcript in alway out sinclude, one, the 24 21 Please good and please draw your 24 1 line 3, the transcript reads: Page 67 1 am talking about is the disulfide cleavage. Do you 24 2 as the disulfide cleavage of an OH protecting 7 1 am talking about is the disulfide cleavage of a 3-OH 3 the 90 percent may ou cleavage of an OH protecting 3 1 am talking abou		-		-
10 reasonable time was an hour. 10 A. I'm sorry. I was distracted by an error 11 A. I agree that one hour would probably be 11 in my - please repeat the question. 12 reasonable for a good, but not great sequencing by 12 C. Stein's SBS method requires quantitative 13 Q. We are talking about Tsien. 12 C. Stein's SBS method requires quantitative 14 Q. We are talking about Tsien. 14 correct? 17 IPR2013-00517. That is Exhibit 2007. Please turn 16 deblocking, yes. 17 17 Q. You wrote that it "requires high and 10 A. I'm sorry. I'was distracted by an error 14 and Ju's DNA sequencing methods require high and 10 A. I'm sorry. I'was distracted by an error 17 IPR2013-00517. That is Exhibit 2007. Please turn 16 deblocking, yes. 14 quantitative efficiency using nucleotides. See 21 A. That is what I wrote. 20 Please turn to page 200, and please draw your 23 3/O-allyl blocking groups include, one, the 21 line 3, the transcript reads: 1 an talking about is the disulfide cleavage. Do you 21 "Question: Slowing at cleavage of a OH protecting			-	
11 A. lagree that one hour would probably be 11 in my - please (repeat the question. 12 reasonable for a good, but not great sequencing by 3 13 ware talking about Tsien. 11 in my - please repeat the question. 14 Q. We are talking about Tsien. 13 deblocking of the 3'-O-allyl protecting group. 14 Q. We are talking about Tsien. 14 correct? 15 A. I'm sorn. Of Tsien's method. 15 A. It requires very high efficient 16 De please go back to your declaration in 16 deblocking, yes. 17 IPR2013-00517. That is Exhibit 2007. Please turn 16 deblocking, yes. 11 in my - please go back to Exhibit 2080, which is guantitative efficiency using nucleotides. 20 and Ju's DNA sequencing methods require high and 20 21 a. That is what i wrote. 20 22 example Tsien at column 20, lines 28 to 34, stating 20 23 that the criteria for the successful use of 21 24 the so percent that you cite here in paragraph 58 context? 4 as the disulfide cleavage of an OH protecting 2 2		-		
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15A. I'm sorry. Of Tsien's method.15A. It requires very high efficient16Q. Please go back to your declaration in16deblocking, yes.17IPR2013-00617. That is Exhibit 2007. Please turn16deblocking, yes.18to page 27, and please draw your attention to18quantitative efficiency using nucleotides, "right?19paragraph 52. In paragraph 52 you write: "Tsien20A. That is what I wrote.20and Ju's DNA sequencing methods require high and18quantitative efficiency using nucleotides. See21quantitative efficiency using nucleotides. See2122A. Jub blocking groups include, one, the2323that the criteria for the successful use of24243-0-allyl blocking groups include, one, the232"Question: Okay. And it's greater than33the 90 percent that you cite here in paragraph 584as the disulfide cleavage conditions. that would5provide an expectation that the disulfide cleavage6conditions, looking at cleavage of an OH protecting7group with greater than 90 percent the sufficient. It's certainly19percent was sufficient. It's certainly19percent was sufficient. It's certainly11on be sufficient. Sufficiency is really something13sufficient. Sufficiency is really something14that would not be sufficient. It's certainly15be sufficient. Sufficiency is really something16approaching 100."1				
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	va Kolliesberg, Fli.D.		
1	Page 69 A. Can I answer that in a second?	1	greater than 90 percent efficiency."
2	Q. Sure.	2	That was your question. And your answer
3	A. It seems to me I'm talking about	3	was: "95 percent would not be sufficient.
4	disulfide cleavage.	4	Sufficiency is really something approaching 100."
5	Q. In relation to cleavage of the 3'-OH	5	A. The problem that I'm having here is that
6	protecting group, correct?	6	at the time, we were talking about and I don't
7	A. I think I'm talking about the efficiency	7	know if that's what we are talking about here, but
8	of disulfide cleavage.	8	one of the issues was simultaneous deprotection of
9	Q. Of the 3'-OH protecting group.	9	a nucleobase linker and the 3' blocking group. So
10	A. No.	10	I'm not sure I'm talking about conditions where you
11	Q. You're talking about cleavage conditions	11	get simultaneous cleavage.
12	could get cleavage of a 3'-OH protecting group.	12	Let me answer the question that I think
13	A. I'm sorry. I don't follow that. I know	13	you're getting at is that I think that we would
14	one of the issues that we were talking about in	14	need you would need for Tsien's sequencing by
15	this case was disulfide cleavage of a nucleobase.	15	synthesis method, I think the answer is applicable
16	Q. Okay, but let's look at the transcript	16	to that, that you would need something greater than
17	again. Starting at page 200, line 3.	17	95 percent for cleavage of the 3' hydrox group, if
18	"Question."	18	that's the question that you're asking.
19	Are you with me?	19	Q. Yes.
20	A. Yes.	20	A. Then I would agree with it. I want to
21	Q. "Okay. And it's greater than 90 percent	21	caveat that I'm not absolutely sure that that is
22	that the cite here in paragraph 58. It's the	22	specifically precisely what I am talking about
23	disulfide cleavage conditions that would provide an	23	here, but I am happy to answer that question.
24	expectation that the disulfide cleavage conditions	24	Q. Okay. And, in fact, you said that
25	could get cleavage of a 3'-OH protecting group with	25	sufficiency is really something approaching 100?
	D		
1	Page 71 A. The closer you get to 100, the better the	1	Page 72 BY MR. SCHWARTZ:
2			
	method's doing to be.	2	Q. Let's go back to Exhibit 2035, your
	method's going to be. Q. Sufficiency is really something closer to	2	Q. Let's go back to Exhibit 2035, your deposition transcript in the IPR2013-00517 case.
3	Q. Sufficiency is really something closer to	2 3 4	deposition transcript in the IPR2013-00517 case.
3 4	Q. Sufficiency is really something closer to 100, correct?	3	deposition transcript in the IPR2013-00517 case. Please turn to page 156. Please draw your
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	ya Romesberg, Ph.D.		University in the City of New York
1	Page 73 A. Yes.	1	Page 74 chain terminator?
2	Q. I'm putting in front of you another	2	"Answer: The sentence on page 3?"
3	document that has not yet been marked. I am	3	And there was an objection.
4	putting you in front of you, Dr. Romesberg, a	4	"Question: Yeah, we yes.
5	document entitled "Video Deposition of Floyd	5	"Answer: It says that they are not very
6	Romesberg Ph.D. in IPR2013-00266.	6	efficient chain terminators.
7	MR. SCHWARTZ: Court reporter, could you	7	"Question: And what does that mean to be
8	please mark this as Exhibit 2082?	8	a not very efficient chain terminator?
9	THE WITNESS: I'm sorry, so this is going	9	"Answer: That it is not a good
10	to be	10	substrate.
11	MR. SCHWARTZ: 2082.	11	"Question: And what do you mean by not a
12	THE WITNESS: In the future, this is	12	good substrate?
13	going to be today, this will be referred to as?	13	"Answer: It is not recognized well and
14	(Exhibit 2082 was marked for	14	
15	identification.)	15	That was your testimony, correct?
16	BY MR. SCHWARTZ:	16	A. That is what is written, yes.
17	Q. 2082. Please turn to page 66. And	17	Q. And when you are referring to a chain
18	please draw your attention to line 17. And	18	terminator, you are referring to the 3' protecting
19	starting at line 17 the transcript reads:	19	group, correct?
20	"Question: Okay. So what is the	20	A. I would have to check. A 3' protected
21	concentration needed to reach high incorporation?	21	nucleotide is certainly a chain terminator.
22	"Answer: Of these substrates would be	22	Q. So a 3' protecting group that is not
23	one millimolar.	23	efficiently incorporated by a polymerase means that
24	"Question: Does that same sentence	24	it is not a good substrate for the polymerase,
25	mention whether or not it is it's an efficient	25	-
1	A. Yes.	1	Page 76 of ordinary skill"?
1 2	A. Yes.	1	of ordinary skill"?
	A. Yes.Q. And the 3' protecting group that is not a	1 2 3	of ordinary skill"? A. I do.
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	a Komesberg, Fil.D.		
1	Page 77 the 3' protecting group, that would result in an	1	Page 78 for SBS, correct?
2	SBS system becoming out of phase, correct?	2	A. That was the answer to the question that
3	A. The sequencing read would lose phase,	3	I was answering.
4	yes.	4	Q. That's your opinion, right? That if more
5	Q. And when the sequencing I'm trying to	5	than half of the strands are out of phase after a
6	use your words. When the sequencing system loses	6	given number of cycles, that would not be
7	phase, it's no longer capable of generating	7	sufficient for SBS?
8	reliable sequencing information, correct?	8	A. I think there are ways that you can
9	A. Certainly once it's lost phase, that is	9	recover data, and where exactly it fails to be
10	correct.	10	recoverable, I'm not exactly sure. But this sounds
11	Q. Please go back to Exhibit 2080. That's	11	about right to me.
12	your deposition transcript in IPR2013-00128. And	12	Q. So you stand by your testimony?
13	please turn to page 252. Please draw your	13	A. More or less. Certainly losing phase is
14	attention to line 6. And starting at line 6 the	14	a problem for sequencing.
15	transcript reads:	15	Q. And if more than half the strands were
16	"Question: If more than half of the	16	out of phase, you testified that would not be
17	strands were out of phase after a given number of	17	sufficient for SBS?
18	cycles, would that be sufficient for SBS?	18	MR. ZIMMERMAN: Objection. Asked and
19	"Answer. No. No."	19	answered.
20	That was your testimony, correct?	20	THE WITNESS: I think there are methods
21	A. Yes.	21	that allow you to still sequence as you are losing
22	Q. It's okay. You can smile.	22	phase. I don't know how far currently people can
23	It is your opinion that if more than half	23	do that. I don't know that the specific software
24	the strands are out of phase after a given number	24	algorithms that allow you to do that as to how far
25	of SBS cycles, then that would not be sufficient	25	you can read into a lost phase sequence, a losing
	Page 79		Page 80
1	phase sequence. So I don't know exactly at what	1	MR. ZIMMERMAN: The answer is the answer.
2	point. I think that intuitively I would more or	2	You're going to make him sit here and punch in .93
3	less agree with what I said here, although I don't	3	ten times on a calculator?
4	know the exact value.	4	MR. SCHWARTZ: Yes, I am going to ask him
5	BY MR. SCHWARTZ:	5	to confirm that is 48.4 percent.
6	Q. I am going to take you up on your	6	MR. ZIMMERMAN: Take as much time as you
7	suggestion that we do some calculations.	7	need, Dr. Romesberg.
8	Does anybody have a piece of paper?	8	THE WITNESS: The answer I get is
9	A. You're going to make me do calculations?	9	48 percent.
10	Q. You bet.	10	
11	Let's assume that we have 93 percent	11	Q. And that means that 93 percent cleavage
12	efficiency of cleavage of the 3' protecting group	12	
13	at each cycle. After 10 cycles there would be .93	13	•
14	to the 10th of the sequence polynucleotide	14	
15	containing the correct sequence, correct?	15	
16	A. Um-hmm. Sorry, yes.	16	
17	Q. And so I just want you to do that	17	declaration. Take your time.
18	calculation, .93	18	
19 20	A. Can I use my calculator?	19	
	Q. Absolutely.	20	Q. Okay, so we're at Exhibit 1012?A. Yes.
21	A. Actually, I am not sure my calculator	21	
21 22	has	22	Q. That's your declaration in IPR2018-00291,
21 22 23	has Q. Mine doesn't either. You'll have to do	22 23	Q. That's your declaration in IPR2018-00291, correct?
21 22	has	22	Q. That's your declaration in IPR2018-00291, correct?A. Yes.

1 10	yd Romesberg, Ph.D.		University in the City of New Yorl
1	Page 81 paragraph 40 you have a header which reads: "It was	1	Page 82 Q. And you write: "Thus, a person of
2	known that the 3' capping group should be small."	2	ordinary skill in the art would not have had an
3	Do you see that?	3	expectation that a small and removal protecting
4	A. I do.	4	group would also possess all of the listed
5	Q. Please keep that open, but I'd like you	5	properties of Ju and Tsien's protecting group."
6	to go back to Exhibit 2007. That is your	6	That is what you wrote, correct?
7	declaration in IPR2013-00517. And please go to	7	A. Yes.
8	page 32. Please draw your attention to paragraph	8	Q. And, again, the Tsien reference that
9	58, and please draw your attention to the second		you're talking about is the same Tsien reference
10	sentence. And you write: "One skilled in the art	9 10	that we have been talking about in this case?
11	would have understood that the physical properties	11	A. It is.
12	underlying, one, nucleotide stability; two,	12	
13	polymerase recognition; and, three, reinitiation of	13	
14	DNA synthesis following deprotection are not	14	expectation that a small and removable protecting
15	necessarily related to; four, size; and, five, the	15	group would also possess all of the listed
16	chemical reversibility. A protecting group can	16	properties of the Tsien's protecting group,
17	have any subset of the aforementioned properties	17	
18	independent of the other properties."	18	A. Sure, yes. I assume that you're going to
19	And to save time, I'm not going to read	19	have me clarify this, but
20	the rest of paragraph 58 into the record, but I	20	Q. No, I'm just asking you that you heard
21	would ask that you turn to page 34 where you have a	21	the question. I could repeat it again. It is your
22	concluding sentence. It starts with "Thus, a	22	opinion that a person of ordinary skill in the art
23	person of ordinary skill in the art."	23	would not have had an expectation that a small and
24	Do you see that?	23	-
25	A. Yes.	25	
20	7. 103.	20	of the listed properties of fisien's protecting
L			
1	Page 83 group?	1	Page 84 deblocking stage, correct?
1	Page 83 group? A. Yes.	1	deblocking stage, correct?
2	group? A. Yes.	1 2 3	deblocking stage, correct? A. That is listed as No. 3, yes.
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2 3	group? A. Yes. Q. Go back to Tsien, please. That is Exhibit 1013, back to pages 20 and 21. And, again,	2 3	 deblocking stage, correct? A. That is listed as No. 3, yes. Q. It is your opinion that "a person of ordinary skill in the art" and, again, I'm using
2 3 4	group? A. Yes. Q. Go back to Tsien, please. That is Exhibit 1013, back to pages 20 and 21. And, again, Tsien's required properties for successfully	2 3 4	 deblocking stage, correct? A. That is listed as No. 3, yes. Q. It is your opinion that "a person of ordinary skill in the art" and, again, I'm using it in the way you do in October 2000 "would not
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	Page 85 1 (Exhibit 1035 was previously marked for	1	Page 86 protecting groups, correct?
	2 identification.)	2	A. Including conditions that under which
	BY MR. SCHWARTZ:	3	you could get deprotection.
	4 Q. I am going to put in front of you a	4	Q. But you relied on this paper to form your
	5 different exhibit.	5	opinions about deprotection efficiencies. You say
	6 I am putting in front of you Exhibit 1035	6	so in your declaration.
	7 entitled "Cleavage of Allyl Ethers with Palladium	7	A. So I included two specific examples of
	8 Carbon" by Boss and Scheffold. You relied on this	8	deprotection. This is a paper that I relied
	9 paper to form your opinions about deprotection	9	particularly strongly on some aspects that
1	0 efficiency of allyl groups, correct?	10	weren't there certainly is data in here that
1	1 A. Actually, no.	11	lead that I included in to form my opinion
1	2 Q. Okay. Why don't we take you back to your	12	
1	3 declaration that we looked at in IPR2018-00291.	13	
1	4 MR. ZIMMERMAN: 10 of 12. Go to page 64.	14	of the reaction as well.
1	5 THE WITNESS: On what page?	15	Q. As an additional question, none of the
1	6 BY MR. SCHWARTZ:	16	
1	7 Q. Page 32. And do you see where it says	17	correct?
1	8 "It was also known it's paragraph 64 "it was	18	A. They are not.
1	9 also known that allyl groups were generally	19	Q. And when you opined that Boss talked that
2	0 efficiently removed using palladium including under	20	allyl groups were efficiently removed, you were
2	1 aqueous conditions," and you cite to Exhibit 1035,	21	referring to the first four entries in table one
2	2 which is Boss?	22	that reported yields of greater than 95 percent,
2	3 A. Yes.	23	correct?
2	4 Q. So you relied on this paper to form your	24	A. I was.
2	5 opinions about deprotection efficiencies of allyl	25	Q. And that data comes from structures four,
\vdash	Page 87		Page 88
	1 five, six and seven, correct?	1	represented the efficient deprotection so, yes.
	 five, six and seven, correct? A. That is correct. 	1 2	
			represented the efficient deprotection so, yes.
	A. That is correct.	2	represented the efficient deprotection so, yes. Q. Let me ask you a few questions about
	A. That is correct.Q. Is it fair to say that yields of	2 3	 represented the efficient deprotection so, yes. Q. Let me ask you a few questions about those compounds in Table 1. As an initial
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	a Romesberg, Ph.D.		University in the City of New York
1	Page 89 Q. And it's a primary alcohol?	1	Page 90 And if it to be clear, if it is not
			rate limiting that means that in a series of steps,
2	A. It is a primary.	2	
3	Q. So a person of ordinary skill in the art	3	if you have a step that is slower than everyone
4	understood that the structure that is most relevant	4	else, that step is called rate limiting, and that
5	in terms of the type of alcohol that is on the	5	is the step that is going to determine the rate of
6	protecting group is compound 10 to a nucleotide	6	the reaction. And if you are competing with other
7	3'-OH because they're both secondary alcohols,	7	things, that determines the efficiency of the
8	correct?	8	deprotection in this case.
9	A. I think that is hard to say. You have to	9	These Palladium-catalyzed reactions are
10	understand the mechanism of these reactions. They	10	multi-step. The palladium has to bind. A
11	are very complex, and even to understand any one	11	nucleophile has to attack. You then have to do a
12	even today, it's a mechanistic study. What you're	12	rearrangement that involves displacement of that
13	getting at is whether or not the departure of the	13	leaving group. That step may or may not be rate
14	hydroxyl, that deprotection step, whether that	14	limiting, so the nature of the actual leaving group
15	weight limits the reaction. Sometimes it does;	15	may or may not identify commonalities between them.
16	sometimes it doesn't. And so sometimes the	16	Q. Is it true that in terms of the compounds
17	efficiency of the stability, what you're getting at	17	that we are looking at in the allyl on an aromatic
18	is whether or not the stability of that leaving	18	alcohol is generally easier to cleave than an allyl
19	group would impinge will affect the reaction.	19	on a secondary alcohol?
20	And that is not known. It could, but it might not,	20	A. It might be, but it is not an absolute.
20	-	20 21	C
	and so I think it is a little bit risky to say that		Q. And is it true that an allyl on a primary
22	that compound in particular represents the sort of	22	alcohol is generally easier to cleave than a allyl
23	yield, the sort of reaction that you would expect	23	on a secondary alcohol?
24	with a nucleotide, because that step is often not	24	A. If it's the rate limiting step, then,
25	rate limiting.	25	yes; if it is not, then, no. Let me rephrase that.
	Page 91		Page 92
1	Page 91 The actual step itself, what you said is correct.	1	Page 92 aromatic alcohols that we're talking about here,
1 2		1 2	aromatic alcohols that we're talking about here, generally speaking, the aromatic is easier to
2	The actual step itself, what you said is correct.		aromatic alcohols that we're talking about here,
2	The actual step itself, what you said is correct. But whether you actually can even see that because	2	aromatic alcohols that we're talking about here, generally speaking, the aromatic is easier to
23	The actual step itself, what you said is correct. But whether you actually can even see that because whether that those differences contribute to the	2 3	aromatic alcohols that we're talking about here, generally speaking, the aromatic is easier to cleave than a secondary?
2 3 4	The actual step itself, what you said is correct. But whether you actually can even see that because whether that those differences contribute to the rate of the reaction and the efficiency for the	2 3 4	aromatic alcohols that we're talking about here, generally speaking, the aromatic is easier to cleave than a secondary? A. It depends if that step is what limits
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	yu Komesberg, Ph.D.		
1	think any of the functionality in DNA is Page 93	1	Page 94 there is a sentence that starts "Furthermore, it
		2	was known that a nucleotide."
2	particularly poisoning to a palladium-catalyzed	-	
3	reaction.	3	Do you see that?
4	I don't see any I'm not envisioning	4	A. I do.
5	any functionality in what I imagine is the standard	5	Q. And you write: "Furthermore, it was known
6	implementation of a linker, fluora-4, in a	6	that a nucleotide having a 3'-O azido ester was not
7	sequencing by synthesis methodology that would be	7	a substrate for polymerases examined by Metzker,"
8	incompatible with palladium, I'm not imagining	8	and you cite the Metzker '94 paper. And that is
9	that. I'm not seeing that, if that is what you are	9	the same Metzker paper that you used in your
10	asking.	10	declarations in these cases, correct?
11	MR. ZIMMERMAN: I think we need to change	11	A. That is correct.
12	the tape.	12	Q. And then you write: "Thus, 3'-O azido
13	THE VIDEOGRAPHER: This marks the end of	13	esters were known to interfere with polymerase
14	media No. 1 of the deposition of Floyd Romesberg	14	recognition and incorporation of the modified
15	Ph.D. We are off the record at 11:12 a.m.	15	nucleotide despite the small size and removability
16	(Recess taken.)	16	of this protecting group."
17	THE VIDEOGRAPHER: We are on the record	17	That is what you wrote, correct?
18	at 11:18 a.m., and this marks the beginning of	18	A. That is what is written.
19	media No. 2 of the deposition of Floyd Romesberg	19	Q. So it is your opinion that Metzker 1994
20		20	showed that 3'-O azido esters were known to
21	BY MR. SCHWARTZ:	21	interfere with polymerase recognition and
22	Q. Dr. Romesberg, please go back to	22	incorporation of the modified nucleotide, despite
23	Exhibit 2007. It's your declaration in	23	the small size and removability of this protecting
	IPR2013-00517. Please turn to page 33, and please	23 24	· · · ·
		24	
25	draw your attention to the middle of the page where	25	A. That is what is written.
	Deve AF		D
	Page 95		Page 96
1	Q. Please turn to page 34 of the	1	THE VIDEOGRAPHER: We are on the record
2	Q. Please turn to page 34 of the declaration, and please look at the second sentence	2	THE VIDEOGRAPHER: We are on the record at 11:25 a.m.
2 3	Q. Please turn to page 34 of the declaration, and please look at the second sentence and you write: "Thus, a person of ordinary skill in	2 3	THE VIDEOGRAPHER: We are on the record at 11:25 a.m. BY MR. SCHWARTZ:
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-	ya Romesberg, Ph.D.		
1	Page 97 Q. So a person with ordinary skill would	1	Page 98 redacted where 3'-O-allyl protecting group was
2	have had an expectation that 3'-O acetyl esters	2	removed from nucleotides under aqueous conditions."
3	would be expected to interfere with polymerase	3	Please look at paragraph 75 on page 40.
4	recognition and incorporation of the 3'-O acetyl	4	In paragraph 75 you write: "In my opinion, these
5	ester nucleotide, correct?	5	notebook pages report that even under optimized
6	A. With the polymerases examined by Metzker,	6	conditions long time periods are required to
	yes.	7	achieve 3'-0 allyl protecting group cleavage
8	Q. Sitting here today are you aware of any	8	efficiencies that are sufficient for SBS. For
9	other polymerases that had been tested prior to	9	example, 3'-O-allyl protecting group was at best
10	October 2000 with the acetyl ester?	10	only redacted this is insufficient for SBS."
11	A. I did not research that, so off the top	11	The passages I read into the record is
12	of my head I could not come up with an example. I	12	what you wrote, correct?
13	am not positive that people did not try, but I	13	A. That is what is written, yes.
14	don't know of one.	14	Q. And in your opinion, the Illumina
15	Q. Let's go back to your declaration and	15	laboratory notebooks you reviewed show that even
16	IPR2013-00517, Exhibit 2007, and please look at	16	under optimized conditions in aqueous media, long
17	paragraph 67 on page 37. In paragraph 67 you	17	time periods are required to achieve 3'-O-allyl
18	write: "I have reviewed several notebook pages	18	protecting group cleavage efficiencies that are
19	provided by Illumina dated" blank. And the date	19	sufficient for SBS, correct?
20	is and that was redacted.	20	A. Under the optimized conditions that they
21	Do you see that?	21	reported, that is what I'm saying.
22	A. Yes, I do.	22	Q. And you conclude from looking at those
23	Q. Please look at paragraph 68. In	23	notebooks that the 3'-O-allyl protecting group
24	paragraph 68 you write: "Notebook SLB090 describes	23	
25		25	A. I don't believe I concluded that. I
20		20	
	Dama 00		Page 100
1	Page 99	1	
1	believe I concluded that the conditions the	1	from the bottom of page 15, there is a sentence
2	believe I concluded that the conditions the precise result they reported under the optimized	2	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in
2 3	believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient	2 3	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art."
2 3 4	believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes.	2 3 4	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that?
2 3 4 5	believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in	2 3 4 5	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. do.
2 3 4 5 6	 believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in 2014, you were aware of Metzker 1994 and the data 	2 3 4 5 6	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. I do. Q. You write: "A person of ordinary skill in
2 3 4 5 6 7	believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in 2014, you were aware of Metzker 1994 and the data in Metzker 1994 for the 3'-O-allyl nucleotide,	2 3 4 5 6 7	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. I do. Q. You write: "A person of ordinary skill in the art seeking to develop sequencing by synthesis
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in 2014, you were aware of Metzker 1994 and the data in Metzker 1994 for the 3'-O-allyl nucleotide, correct? A. I was, yes. Q. Please go back to Exhibit 2081, that is your second declaration in IPR2013-00266. MR. ZIMMERMAN: I don't think we have 2081. THE WITNESS: Yeah, I thought we skipped 2081. 2082? MR. SCHWARTZ: You don't have this before. I have skipped it before. (Exhibit 2081 was marked for identification.) BY MR. SCHWARTZ: Q. And so are you ready? A. Um-hmm. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. I do. Q. You write: "A person of ordinary skill in the art seeking to develop sequencing by synthesis methods, would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations." That is what you wrote, right? A. That is what you wrote. Q. And it is your opinion that a person of ordinary skill in the art seeking to develop an SBS method would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations, correct? A. Again, I believe that is what is written. Q. Please go back to Tsien. To Tsien's paper, Exhibit 1013.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in 2014, you were aware of Metzker 1994 and the data in Metzker 1994 for the 3'-O-allyl nucleotide, correct? A. I was, yes. Q. Please go back to Exhibit 2081, that is your second declaration in IPR2013-00266. MR. ZIMMERMAN: I don't think we have 2081. THE WITNESS: Yeah, I thought we skipped 2081. 2082? MR. SCHWARTZ: You don't have this before. I have skipped it before. (Exhibit 2081 was marked for identification.) BY MR. SCHWARTZ: Q. And so are you ready? A. Um-hmm. Q. I put in front of you Exhibit 2081, which 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. I do. Q. You write: "A person of ordinary skill in the art seeking to develop sequencing by synthesis methods, would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations." That is what you wrote, right? A. That is what l wrote. Q. And it is your opinion that a person of ordinary skill in the art seeking to develop an SBS method would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations, correct? A. Again, I believe that is what is written. Q. Please go back to Tsien. To Tsien's paper, Exhibit 1013. A. I have it.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 believe I concluded that the conditions the precise result they reported under the optimized conditions that they reported would be insufficient for SBS, yes. Q. And when you came to that conclusion in 2014, you were aware of Metzker 1994 and the data in Metzker 1994 for the 3'-O-allyl nucleotide, correct? A. I was, yes. Q. Please go back to Exhibit 2081, that is your second declaration in IPR2013-00266. MR. ZIMMERMAN: I don't think we have 2081. THE WITNESS: Yeah, I thought we skipped 2081. 2082? MR. SCHWARTZ: You don't have this before. I have skipped it before. (Exhibit 2081 was marked for identification.) BY MR. SCHWARTZ: Q. And so are you ready? A. Um-hmm. Q. I put in front of you Exhibit 2081, which is your second declaration in IPR2013-00266. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	from the bottom of page 15, there is a sentence that starts with "The person of ordinary skill in the art." Do you see that? A. 1 do. Q. You write: "A person of ordinary skill in the art seeking to develop sequencing by synthesis methods, would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations." That is what you wrote, right? A. That is what I wrote. Q. And it is your opinion that a person of ordinary skill in the art seeking to develop an SBS method would recognize the need to use a nucleotide with a 3' protecting group that could be incorporated by a polymerase with high fidelity at moderate to low concentrations, correct? A. Again, I believe that is what is written. Q. Please go back to Tsien. To Tsien's paper, Exhibit 1013. A. I have it. Q. Please turn to page 38. On page 38 Tsien

	yu Komesberg, Fil.D.		
1	Page 101 Incorporation of Labeled Nucleotide Analogs Into	1	that is what I wrote, so, yes. Page 102
2	DNA."	2	Q. A POSA understood that a nucleotide
3	Do you see that?	3	concentration of 200 micromolars is higher than
4	A. I do.	4	what typically would be expected for SBS, correct?
5	Q. You can read through this entire	5	A. Would a person of ordinary skill in the
6	paragraph if you need to in order to answer my	6	art thought 200 micromolar was higher than what?
7	question, but for now I'll just ask you to look at	7	Q. Well, when you were talking about
8	lines 14 through 19. And Tsien teaches the use of	8	moderate to low concentrations for SBS, would it be
9	three unlabeled 3' blocked deoxy nucleotide	9	higher than the moderate to low concentrations?
10	triphosphate dNTP analogs at a concentration of 1.5	10	A. 200 micromolar, no. That's probably in
11	micromolar each and one 3' blocked fluorescently	11	the moderate range. That is certainly not in the
12	labeled dNTP analog at a concentration of 30	12	low range.
13	micromolar for his SBS method, correct?	13	Q. But Tsien himself uses them at much lower
14	A. That would seem to be correct.	14	
15	Q. And Tsien's teaching of 1.5 to 30	15	A. That would seem to be the case here, yes.
16	micromolar concentrations excuse me of	16	(Exhibit 1016 was previously marked for
17	nucleotides is consistent with your opinion that a	17	identification.)
18	person of ordinary skill in the art seeking to	18	
19	develop sequencing by synthesis methods would	19	Q. I am putting in front of you
20	recognize the need to use a nucleotide with a 3'	20	Exhibit 1016, a paper entitled "Termination of DNA
21	protecting group that could be incorporated by a	21	Synthesis by Novel 3' Modified Deoxyribonucleoside
22	polymerase with high fidelity of moderate to low	22	5'-triphosphates" by Metzker that was published in
23	concentrations, correct?	23	
24	A. I think that's just the question that you	24	This is the same Metzker 1994 paper you
25	asked earlier, and if that's the case, then, yes,	25	
20			
1	Page 103	1	Page 104
1	correct?	1	which there's a lot of data, and then table two is
2	A. That is correct.	2	which there's a lot of data, and then table two is sort of a summary of that. And so it is not
2 3	correct?A. That is correct.Q. Please look at the abstract. In the	2 3	which there's a lot of data, and then table two is sort of a summary of that. And so it is not exactly those assays, it is compiled from the
2 3 4	correct? A. That is correct. Q. Please look at the abstract. In the abstract the authors write "Eight modified-dNTPs	2 3 4	which there's a lot of data, and then table two is sort of a summary of that. And so it is not exactly those assays, it is compiled from the assays.
2 3 4 5	correct? A. That is correct. Q. Please look at the abstract. In the abstract the authors write "Eight modified-dNTPs were synthesized and tested in two different	2 3 4 5	which there's a lot of data, and then table two is sort of a summary of that. And so it is not exactly those assays, it is compiled from the assays.Q. But if you look on the right side of
2 3 4 5 6	correct? A. That is correct. Q. Please look at the abstract. In the abstract the authors write "Eight modified-dNTPs were synthesized and tested in two different template assays for incorporation activity. From	2 3 4 5 6	 which there's a lot of data, and then table two is sort of a summary of that. And so it is not exactly those assays, it is compiled from the assays. Q. But if you look on the right side of page 4263 of Tsien, it talks about a
2 3 4 5 6 7	correct? A. That is correct. Q. Please look at the abstract. In the abstract the authors write "Eight modified-dNTPs were synthesized and tested in two different template assays for incorporation activity. From this"	2 3 4 5 6 7	 which there's a lot of data, and then table two is sort of a summary of that. And so it is not exactly those assays, it is compiled from the assays. Q. But if you look on the right side of page 4263 of Tsien, it talks about a A. I'm sorry
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yd Romesberg, Ph.D.		University in the City of New York
Page 105 discovery projects to mean a preliminary test that	1	Page 106 mean, certainly he compares to the dideoxy
		termination pattern, which sort of is Sanger
		sequencing. So I didn't ask myself were each of
		his assays precisely Sanger sequencing, but I think
-		essentially that is correct.
	6	Q. Included in that group of seven 3'-O
-		protecting groups is a 3'-O-allyl protecting group,
-	8	correct?
	9	A. It is compound 3, yes.
-	10	Q. And going back to the abstract we just
o 11	11	looked at that from those screening tests that are
•	12	reported in table two, Metzker identified two 3'-O
	13	protected nucleotides that Metzker says may be used
-	14	as alternative terminators in Sanger sequencing,
-	15	correct?
A. They can be used in that way.	16	A. Yes.
5	17	Q. And Metzker states in the abstract that
-	18	"the 3'-O methyl dNTPs were shown to terminate DNA
	19	synthesis and may be used as alternative
	20	terminators in Sanger sequencing."
Q. Going back to table two of Metzker.	21	That is what he states, correct?
Metzker tested modified nucleotides with seven	22	A. That is what is written.
different 3'-O protecting groups to see how they	23	Q. Metzker does not say that the 3'-O-allyl
behaved in Sanger sequencing, correct?	24	dNTP was shown to terminate DNA synthesis and may
A. Was it exactly Sanger sequencing? I	25	be used as an alternative terminator in Sanger
Page 107		Page 108
	1	(Exhibit 2053 was marked for
A. He does not say that.	2	identification.)
Q. I would like you to keep Metzker '94 in	3	BY MR. SCHWARTZ:
front of you. I'm going to mark another paper.	4	Q. Dr. Romesberg, I'm putting in front of
MR. SCHWARTZ: This will require a new	5	you a document entitled "Sequencing technologies,
stamp, Court Reporter, 2053.	6	the next generation," by Michael Metzker. It was
(Reporter clarification.)	7	published in 2010. The author of this article is
MR. SCHWARTZ: Some of these are going to	8	the same Michael Metzker who is an author of the
be a little bit out of order. I'll be coming back	9	Metzker 1994 paper we were just looking at,
and forth, so instead of changing my whole index,	10	correct?
I'm just going to ask you to mark this as 2053.	11	A. That is correct.
This is actually something already marked in these	12	Q. Please turn to page 34 of that paper, and
,	13	please draw your attention to the left column which
5	14	contains a section header entitled "Cyclic
	15	reversible termination."
	16	Do you see that?
MR. SCHWARTZ: It's so far only of record	17	A. I do.
	18	Q. And please look at the second paragraph
in the method case.		
MR. ZIMMERMAN: Because we offered the	19	in that section that starts with "The key to the
MR. ZIMMERMAN: Because we offered the witness today on the method. And you're not taking	19 20	CRT Method."
MR. ZIMMERMAN: Because we offered the witness today on the method. And you're not taking him on the method, but you're using exhibits from	19 20 21	CRT Method." Do you see that?
MR. ZIMMERMAN: Because we offered the witness today on the method. And you're not taking him on the method, but you're using exhibits from the method case. Just want that to be clear from	19 20 21 22	CRT Method." Do you see that? A. I do.
MR. ZIMMERMAN: Because we offered the witness today on the method. And you're not taking him on the method, but you're using exhibits from the method case. Just want that to be clear from the record.	19 20 21 22 23	CRT Method." Do you see that? A. I do. Q. CRT means Cyclic Reversible Termination,
MR. ZIMMERMAN: Because we offered the witness today on the method. And you're not taking him on the method, but you're using exhibits from the method case. Just want that to be clear from	19 20 21 22	CRT Method." Do you see that? A. I do. Q. CRT means Cyclic Reversible Termination,
	discovery projects to mean a preliminary test that is used to identify compounds that have promising properties? A. It can be used in that way. That is not the only way it is used. Q. But that is a way that it is used? A. It is used in that way. It is also used as a preliminary analysis that is not superficial. A screen itself does not necessarily have to be rough or approximate. It is somehow limited and is expected to then lead to information that would help you move forward. Q. And is it fair to say that screens are also used to decide which compounds not to move forward with? A. They can be used in that way. Q. And that would be because they don't have necessarily the properties you are looking for in your discovery project, right? A. Yes. Q. Going back to table two of Metzker. Metzker tested modified nucleotides with seven different 3'-O protecting groups to see how they behaved in Sanger sequencing, correct? A. Was it exactly Sanger sequencing? I Page 107 sequencing, right? A. He does not say that. Q. I would like you to keep Metzker '94 in front of you. I'm going to mark another paper. MR. SCHWARTZ: This will require a new stamp, Court Reporter, 2053. (Reporter clarification.) MR. SCHWARTZ: Some of these are going to be a little bit out of order. I'll be coming back and forth, so instead of changing my whole index, I'm just going to ask you to mark this as 2053. This is actually something already marked in these IPRs, but it wasn't in all four of the early four. MR. ZIMMERMAN: Is it a record only in the method case or is it a record only in the method case or is it a record only in the method case or is it a record only in the method case or is it a record only in the method case or is it a record only in	discovery projects to mean a preliminary test that1is used to identify compounds that have promising2properties?3A. It can be used in that way. That is not4the only way it is used.5Q. But that is a way that it is used?6A. It is used in that way. It is also used7as a preliminary analysis that is not superficial.8A screen itself does not necessarily have to be9rough or approximate. It is somehow limited and is10expected to then lead to information that would11help you move forward.12Q. And is it fair to say that screens are13also used to decide which compounds not to move14forward with?15A. They can be used in that way.16g. And that would be because they don't have17necessarily the properties you are looking for in18your discovery project, right?19A. Yes.20Q. Going back to table two of Metzker.21behaved in Sanger sequencing, correct?24A. Was it exactly Sanger sequencing? I25sequencing, right?1A. He does not say that.2Q. I would like you to keep Metzker '94 in3front of you. I'm going to mark another paper.4MR. SCHWARTZ: This will require a new5stamp, Court Reporter, 2053.6(Reporter clarification.)7MR. SCHWARTZ: Some of these are going tobe a little bit out of order. I'll be coming back<

· · · ·			
1	Page 109 Q. And the CRT method is another name for	1	Page 110 A. Yes.
2	SBS, correct?	2	Q. So in Metzker's 2010 paper, he states
3	A. I would have to read that. It is not a	3	that the work he reported in his 1994 paper on
4	term that I use. In fact, I don't think it is a	4	terminators for Sanger sequencing provided the
5	particularly commonly used term. I'd have to see	5	basis for development of 3'-O reversible blocking
6	whether that refers to a specific step of SBS or if	6	groups for SBS, correct?
7	it refers to SBS overall, certainly is a step.	7	A. That is what it states.
8	Q. Please draw your attention to the section	8	Q. Is it fair to say that a reader of
9	that starts with "The use of Dideoxy Nucleotide."	9	Metzker's 1994 paper understood that the work that
10	Do you see that?	10	it was reported in the 1994 paper on terminators
11	A. Yes.	11	for Sanger sequencing provided the basis for his
12	Q. Metzker writes "The use of a Dideoxy	12	development of 3'-O reversible blocking groups for
13	nucleotide which acts as a chain terminator in	13	
14	Sanger sequencing provided the basis for the	14	A. If you're asking me if a person of
15	initial development of reversible blocking groups	15	ordinary skill in the art would have read the 1994
16	attached to the 3' end nucleotides."	16	paper and understood that Metzker was looking to
17	That is what Metzker wrote, correct?	17	develop reversible terminators for what he
18	A. That is what is written, yes.	18	called he didn't use the term sequencing by
19	Q. And Metzker refers to reference 19 and 20	19	synthesis, but for that, then, yes, I think a
20	support for that statement, correct?	20	reader I think he states that in the 1994 paper.
21	A. Yes.	21	Q. And is it fair to say the person of
22	Q. And would you please turn to the	22	ordinary skill in the art and, again, whenever I
23	references section and look at references 19 and	23	use POSA, I'm referring to POSA that you defined,
24	20. And reference 19 is the same Metzker 1994	24	goes back to October 2000 is it fair to say that
25	paper that we just looked at, correct?	25	a person of ordinary skill in the art understand
	B 444		B 110
1	Page 111 that a cleavable 3'-O protecting group that works	1	Page 112 an inefficient terminator in SBS?
1	that a cleavable 3'-O protecting group that works		an inefficient terminator in SBS?
	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for	2	an inefficient terminator in SBS? A. I want to be careful about how I answer
2	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS?	2 3	an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a
2 3	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it	2	an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient
2 3 4	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also	2 3 4 5	an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what?
2 3 4 5 6	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it	2 3 4 5 6	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all
2 3 4 5 6	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary	2 3 4 5 6 7	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you
2 3 4 5 6 7	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it	2 3 4 5 6	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very
2 3 4 5 6 7 8	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion	2 3 4 5 6 7 8	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define
2 3 4 5 6 7 8 9	that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes.	2 3 4 5 6 7 8 9	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19	 that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes. Q. And by the same token, is it fair to say that a POSA understood that 3' protecting group that inhibits polymerase activity in Sanger sequencing would also likely inhibit polymerase activity in SBS? A. I don't know what you mean by the word "inhibit." Q. The way that Metzker uses the word inhibit in his paper. A. No, I do not believe that that would be a 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look at any one of them if it's not clear by context, which I hope it would be. But in this case it's really not useful to simply categorize something as interference means inefficient. And I'll tell you why. You can imagine a modification of A to C. The natural triphosphate that polymerases use as natural substrates. C is actually recognized less efficiently than A. So would I call that
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes. Q. And by the same token, is it fair to say that a POSA understood that 3' protecting group that inhibits polymerase activity in Sanger sequencing would also likely inhibit polymerase activity in SBS? A. I don't know what you mean by the word "inhibit." Q. The way that Metzker uses the word inhibit in his paper. A. No, I do not believe that that would be a necessary fatal flaw to moving forward with an 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look at any one of them if it's not clear by context, which I hope it would be. But in this case it's really not useful to simply categorize something as interference means inefficient. And I'll tell you why. You can imagine a modification of A to C. The natural triphosphate that polymerases use as natural substrates. C is actually recognized less efficiently than A. So would I call that interference? No, I mean, it's a natural substrate. So I think the question that you're
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes. Q. And by the same token, is it fair to say that a POSA understood that 3' protecting group that inhibits polymerase activity in Sanger sequencing would also likely inhibit polymerase activity in SBS? A. I don't know what you mean by the word "inhibit." Q. The way that Metzker uses the word inhibit in his paper. A. No, I do not believe that that would be a necessary fatal flaw to moving forward with an analog. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look at any one of them if it's not clear by context, which I hope it would be. But in this case it's really not useful to simply categorize something as interference means inefficient. And I'll tell you why. You can imagine a modification of A to C. The natural triphosphate that polymerases use as natural substrates. C is actually recognized less efficiently than A. So would I call that interference? No, I mean, it's a natural substrate. So I think the question that you're asking is at what point does interference preclude
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes. Q. And by the same token, is it fair to say that a POSA understood that 3' protecting group that inhibits polymerase activity in Sanger sequencing would also likely inhibit polymerase activity in SBS? A. I don't know what you mean by the word "inhibit." Q. The way that Metzker uses the word inhibit in his paper. A. No, I do not believe that that would be a necessary fatal flaw to moving forward with an analog. Q. Is it fair to say that a person of 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look at any one of them if it's not clear by context, which I hope it would be. But in this case it's really not useful to simply categorize something as interference means inefficient. And I'll tell you why. You can imagine a modification of A to C. The natural triphosphate that polymerases use as natural substrates. C is actually recognized less efficiently than A. So would I call that interference? No, I mean, it's a natural substrate. So I think the question that you're asking is at what point does interference preclude its use in sequencing by synthesis. And that's a
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 that a cleavable 3'-O protecting group that works for Sanger sequencing would be a good candidate for development of a blocking group for SBS? A. If it worked for Sanger sequencing, it would satisfy certain requirements that would also be required for sequencing by synthesis, so it would be a piece of data that a person of ordinary skill in the art would look to to form an opinion about sequencing by synthesis, yes. Q. And by the same token, is it fair to say that a POSA understood that 3' protecting group that inhibits polymerase activity in Sanger sequencing would also likely inhibit polymerase activity in SBS? A. I don't know what you mean by the word "inhibit." Q. The way that Metzker uses the word inhibit in his paper. A. No, I do not believe that that would be a necessary fatal flaw to moving forward with an analog. Q. Is it fair to say that a person of ordinary skill in the art understood that the 3' 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 an inefficient terminator in SBS? A. I want to be careful about how I answer this question, because inefficient is not a particularly useful term here. Inefficient relative to what? Q. Well, you used the word inefficient all the time in what you A. And I believe I used it very qualitatively, and if you wish for me to define more specifically what I meant, I'm happy to look at any one of them if it's not clear by context, which I hope it would be. But in this case it's really not useful to simply categorize something as interference means inefficient. And I'll tell you why. You can imagine a modification of A to C. The natural triphosphate that polymerases use as natural substrates. C is actually recognized less efficiently than A. So would I call that interference? No, I mean, it's a natural substrate. So I think the question that you're asking is at what point does interference preclude its use in sequencing by synthesis. And that's a slightly hard question to answer without

1	to say that if it's not accepted as a substrate,	1	Page 114 were inhibitors in 1994?
2	then that is not going to be a good candidate for	2	A. I think if you looked at the 2010 paper
3	sequencing by synthesis.	3	and those compounds that he found inhibited in 1994
4	If it is recognized as well as a natural	4	weren't there, I think that you would have
5	substrate, that's awesome. People would be really	5	logically assumed that he got rid of them, he
6	excited, and they would want to move forward in the	6	didn't pursue them because they were not promising.
7	sequence by synthesis application. But there's a	7	Q. That is the basis for my question to you,
8	whole range of gray between the two. And at what	8	the same for inefficient. Those that he found to
9	point it becomes so inefficiently recognized that	9	be inefficient, he didn't move forward with those.
10	it could be not be used as a substrate, it's	10	He made that clear in the 2010 paper, correct?
11	certainly there's an area in there where that is	11	A. Are you asking me if this is the only
12	going to happen.	12	metric that he used to decide to move forward?
13	Q. I think maybe we're getting a little bit	13	Q. No, I am not asking you if it is the only
14	too complicated here. Basically, Metzker in 2010	14	
15	pointed to the '94 paper and said that the '94	15	A. Certainly the efficiency of which it
16	paper provided the basis for his moving forward	16	caused termination as he described it in the '94
17	with protecting groups for SBS, correct?	17	paper would have been a component of his thought
18	A. Correct. That would be his moving	18	process, of his decision process to decide to what
19	forward with his opinions of what efficiency was	19	to with which in his process to decide what
20	required.	20	he was moving forward with.
21	Q. Okay. And in the 1994 paper he lists	21	Q. And those are reasonable decisions that
22	some nucleotides as being, for example, inhibitors	22	Metzker made. You said his thought process, but
23	of the polymerase. And I'm just saying would a	23	
24	person understand that he didn't choose those to	24	
25	move forward with in his SBS studies because they	25	A. Everyone is going to weigh things
	Page 115		Dava 446
1	differently. I don't know exactly what he was	1	Page 116 SUCCESS, CORRECT.
2	thinking. I don't know exactly what motivations he	2	Q. For DNA sequencing?
3	had to choose one or the other, but it was	3	A. That is correct.
4	certainly a requirement that you have some	4	Q. If I refer to Vent (exo-) DNA polymerase,
5	recognition. He would have never moved forward	5	if I refer to that only as "Vent," will you
6	with something that has no recognition if he had	6	understand that I am referring to the Vent (exo-)
7	multiple things that had recognition, he may have	7	DNA polymerase in Metzker '94?
8	chosen one over the other for other reasons.	8	A. I will now.
9	Q. Let's go back to the allyl compound in	9	Q. Thank you.
10	Metzker, that's compound 3 as you said, right?	10	Metzker reports that the incorporation of
11	When I say "Metzker," I should have clarified that	11	the 3'-O-allyl dATP by the Vent polymerase was
12	Metzker 1994 since there may be several Metzkers on	12	incomplete at the final concentration of 250
13	the table.	13	micromolar, correct?
14	The 3'-O-allyl dATP had no incorporation	14	A. That is what he writes, yes.
15	with seven of the eight polymerases Metzker used in	15	Q. Please turn to page 4265. Please draw
16	the screening assay, correct?	16	your attention to the second paragraph on the right
17	A. That is what it is listed.	17	side of the page that starts with "Some notable
18	Q. Based on those results, a POSA understood	18	correlations."
19	that the 3'-O-allyl dATP would most likely not work	19	Do you see that?
20	for DNA sequencing with any of those seven	20	A. I do.
21	polymerases, correct?	21	Q. And please look at the second sentence in
22	A. With those polymerases under the	22	that paragraph that starts with "First, the
0			
23	condition employed, a POSA would have not been	23	protecting groups."
23	condition employed, a POSA would have not been encouraged would not have had an expect would	23 24	protecting groups." Do you see that?
	encouraged would not have had an expect would		

	a Romesberg, Ph.D.	University in the City of New York			
4	Page 117		Page 118		
1	Q. And Metzker writes "First, the protecting	1	Do you remember that?		
2	groups containing ether linkages at the 3'	2	A. I do.		
3	position, compounds 1, 3, 4, 7 and 8, were	3	Q. So is it fair to say that a person of		
4	incorporated by some of the polymerases."	4	ordinary skill would expect that the 3'-O-allyl		
5	That is what Metzker wrote, correct?	5	dATP could show even less incorporation with Vent		
6	A. That is what he wrote.	6	if it was used at a 10-fold lower concentration		
7	Q. So when Metzker writes that "Compound 3	7	that Tsien suggests?		
8	was incorporated by some of the polymerases," the	8	A. That is right. But Tsien is suggesting a		
9	reader of Metzker understands that the	9	different reaction than what Metzker is looking at,		
10	incorporation Metzker refers to is incomplete	10	and so it's very difficult to compare that.		
11	incorporation with the Vent polymerase at 250	11	Q. I am not asking you to compare it, I'm		
12	micromolar, correct?	12	just saying		
13	A. Under the conditions that he employed,	13	A. You just did.		
14	yes. I'm sorry. Could you repeat that question?	14	-		
15	Q. So when Metzker writes that "Compound 3	15	,		
16	was incorporated by some of the polymerases," the	16			
17	reader of Metzker understands that the	17			
18	incorporation Metzker refers to is incomplete	18	, ,		
19		19			
	incorporation with the Vent polymerase at 250 micromolar, correct?		Q. And my question was, at those lower		
20		20	concentrations of nucleotide, would a person of		
21	A. Under the conditions that he explored	21	ordinary skill reasonably expect that the		
22	that he is reporting, yes.	22			
23	Q. And earlier we looked at the Tsien	23	be even less than he reported at 250 micromolars?		
24	publication, and Tsien suggested using nucleotides	24			
25	in the low micromolar range, 1.5 to 30 micromolar.	25	Q. Please draw your attention to the last		
	Page 119		Page 120		
1	sentence page withdraw the question.	1	contains approximately percent of the unprotected		
1 2	sentence page withdraw the question. Please turn to page 4263 and please draw	1 2			
	sentence page withdraw the question.		contains approximately percent of the unprotected		
2	sentence page withdraw the question. Please turn to page 4263 and please draw	2	contains approximately percent of the unprotected nucleotide, correct?		
2 3	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where	2 3	contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes.		
2 3 4	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where there is a header "3-O Methyl dATP compound 1	2 3 4	contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes. Q. The unprotected dATP?		
2 3 4 5	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where there is a header "3-O Methyl dATP compound 1 incorporation."	2 3 4 5	 contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes. Q. The unprotected dATP? A. Yes. 		
2 3 4 5 6	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where there is a header "3-O Methyl dATP compound 1 incorporation." Do you see that?	2 3 4 5 6	 contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes. Q. The unprotected dATP? A. Yes. Q. When Metzker reports that compound 3, the 		
2 3 4 5 6 7	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where there is a header "3-O Methyl dATP compound 1 incorporation." Do you see that? A. I do.	2 3 4 5 6 7	 contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes. Q. The unprotected dATP? A. Yes. Q. When Metzker reports that compound 3, the 3'-O-allyl dATP, had incomplete termination with Vent at 250 micromolar, it is possible that any 		
2 3 4 5 6 7 8	sentence page withdraw the question. Please turn to page 4263 and please draw your attention to the right side of the page where there is a header "3-O Methyl dATP compound 1 incorporation." Do you see that? A. I do. Q. And please draw your attention to the last sentence in that section that starts with "All	2 3 4 5 6 7 8	 contains approximately percent of the unprotected nucleotide, correct? A. Triphosphate, yes. Q. The unprotected dATP? A. Yes. Q. When Metzker reports that compound 3, the 3'-O-allyl dATP, had incomplete termination with Vent at 250 micromolar, it is possible that any incorporation Metzker observed for 3'-O-allyl dATP 		
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2 3 4 5 6	Page 121 seeing. Some level of the reduced termination, the asterisk that he puts there is exactly due to that, because there is natural dATP incorporating, and it is allowing synthesis to continue.	1 2 3	Page 122 1992 that I've marked as Exhibit 2084. Please draw your attention to the top far-left side of the
2 3 4 5 6 7 8 9	asterisk that he puts there is exactly due to that, because there is natural dATP incorporating, and it		•
3 4 5 6 7 8 9	because there is natural dATP incorporating, and it	2	-
4 5 6 7 8 9		3	page, and under the date of September 1992 it reads
6 7 8 9	o y	4	"A scientific newsletter from New England Biolabs."
7 8 9	Q. I think maybe we're talking past one	5	Do you see that?
7 8 9	another. When he said the termination is	6	THE WITNESS: I do.
8 9	incomplete, he's suggesting that there is some?	7	MR. ZIMMERMAN: Objection. Lacks
9	A. Termination.	8	foundation.
	Q. There is some termination, but that it is	9	BY MR. SCHWARTZ:
	incomplete, meaning that it continues to go on.	10	Q. New England Biolabs is a supplier of
11	And I'm saying that is it possible that that is	11	biochemical products for the scientific community,
12	because of the contamination of the unprotected	12	correct?
	dATP?	13	MR. ZIMMERMAN: Same objection.
14	A. The incomplete termination is because of	14	BY MR. SCHWARTZ:
	the presence of the contaminating dATP. Yes, I	15	Q. You can answer the question.
16	think that is certainly correct.	16	A. NEB is a company that sells such
17	Q. I am going to put in front of you another	17	reagents, yes.
18	document.	18	Q. And including products for DNA
19	MR. SCHWARTZ: This will require a	19	sequencing, correct?
20	marking. 2084.	20	A. I think so.
21	(Exhibit 2084 was marked for	21	Q. Do you recall reading NEB transcripts in
22	identification.)	22	the course of your own work?
23	BY MR. SCHWARTZ:	23	A. No, I did not.
24	Q. I am putting in front of you a document	24	Q. On the far right side of the page is an
1	entitled "The NEB Transcript," dated September 19,	25	
			······································
1	Page 123 Do you see that?	1	Page 124 A. I do.
2	A. I do.	2	Q. And under that header it reads
			-
3	MR. ZIMMERMAN: Objection. Foundation.	3	"CircumVent thermal cycle DNA sequencing is a
	BY MR. SCHWARTZ	4	simple and rapid method which generates sequence
5	Q. And within the index there is an entry	5	information from less template than any other
	entitled "CircumVent Thermal Cycle DNA Sequencing	6	protocol. The kit uses the highly thermostable
	Kit."	7	Vent (exo-) DNA polymerase."
8	Do you see that?	8	That is what it says, right?
9	MR. ZIMMERMAN: Can I have a standing	9	A. That is what it says.
10	objection to this exhibit on foundation, or do you	10	Q. Now, please go back to Metzker 1994,
11	want me to make the objection every time?	11	which is Exhibit 10,016 [sic], and go back to table
12	MR. SCHWARTZ: You can have a standing	12	two. The Vent polymerase used in Metzker 1994 is
13	objection, and the foundation will become clear in	13	the Vent (exo-) DNA polymerase, correct?
14	a minute.	14	A. That is correct.
15	BY MR. SCHWARTZ:	15	Q. Now, please turn to Page 4261 of Metzker.
16	Q. Okay. Do you see where there's an entry	16	And on the right side towards the bottom of page
17	"CircumVent Thermal Cycle DNA Sequencing Kit," in	17	4161 is a header entitled "Polymerases."
	the index, page 12?	18	Do you see that?
19	A. Yes.	19	A. I do.
20	Q. Please turn to the next page, which is		
		20	Q. And please draw your attention to the
21	page 12. And I will represent that Exhibit 284	21	last sentence in that section. Metzker states that
22	2084 is not the complete transcript. It's only the	22	he got the Vent (exo-) DNA polymerase from New
23	portion that refers to the CircumVent Thermal Cycle	23	England Biolabs, correct?
1	DNA Sequencing Kit.	24	A. That is written, yes.
25	Do you see that?	25	Q. So the Vent polymerase in Metzker 1994 is

2 the 3 4 4 1 ha 5 a th 6 dor 7 8 9 say 10 11 12 tra 13 wh 14 15 16 Do 17 tha 18 mi 19 rea 20 se 21 flue 22 0 3 ine 4 sec 5 6 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 ine 10 cal 11 BY 12 res 13 res 14 15 15 I du 16 bas 17 18	i Kolliesberg, Fli.d.					
4 I hat 5 a th 6 dor 7 8 8 poll 9 say 10 11 12 tra 13 wh 14 15 16 Dot 17 the 18 mi 19 rea 20 see 21 flu 22 0 3 inee 4 sec 5 6 6 Mis 7 8 8 inee 10 cal 11 BY 12 0 3 inee 10 cal 11 BY 12 0 13 res 14 15 15 I de 17 18 18 ba 19 tha 10 in	Page 125 he same Vent polymerase that is being discussed in he NEB September 1992 transcript, correct?	2	Page 126 your attention to the second sentence there which starts with "However, when fluorescent tagged."			
5 a th 6 dor 7 8 pol 9 say 10 11 12 tra 13 wh 14 15 16 Dot 17 the 18 mi 19 rea 20 se 21 flu 22 3 23 24 23 24 25 wi 1 exp 3 ine 4 sec 5 Mis 7 8 8 inef 9 inte 10 cal 11 BY 12 0 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc <td>A. I would have to look at this. As I said,</td> <td>3</td> <td>Do you see that?</td>	A. I would have to look at this. As I said,	3	Do you see that?			
6 dor 7 8 pol 9 say 10 11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 see 21 flue 23 24 23 24 23 24 23 3 1 exp 3 ine 4 sec 5 6 6 Mis 7 8 10 cal 11 BY 12 9 13 res 14 15 15 Ide 16 bas 17 18 18 ba 19 tha 20 inc	haven't looked at this before. This seems to be	4	A. Ido.			
7 8 pol 9 say 10 11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 see 21 flue 22 0 23 24 23 24 23 24 23 a 1 exp 2 0 3 ine 4 seco 5 6 6 Mis 7 8 10 cal 11 BY 12 main 13 res 14 15 15 1d 16 bas 17 18 18 ba 19 tha 20 inc	thermal cycle kit. It seems to use Vent. I	5	Q. And the transcript reads, "However, when			
8 pol 9 say 10 11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 22 3 23 24 25 wi 1 exp 3 ine 4 sec 5 6 6 Mis 7 8 8 inef 9 inte 10 cal 11 BY 12 m 13 res 14 15 15 I du 16 bas 17 18 18 ba 19 tha 20 inc	on't know what else is in the kit.	6	fluorescent-tagged dideoxy dNTPs are used in the			
9 say 10 11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flue 23 24 23 24 23 ine 4 sec 5 6 6 Mis 7 8 8 inet 10 cal 11 BY 12 10 13 res 14 15 15 I de 17 18 ba 19 tha 20 inc	Q. I'm just asking you specifically the	7	sequencing method, the results can be mixed.			
10 11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flue 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 ine! 9 inte 10 cal 11 BY 12 main 13 res 14 15 15 Ide 17 18 18 ba 19 tha 20 inc	olymerase. It's the same polymerase that Metzker	8	Inefficient incorporation leads to different			
11 12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 23 24 23 24 25 wi 1 exp 3 ine 4 sec 5 6 6 Mis 7 8 8 inel 9 inte 10 cal 11 BY 12 0 13 res 14 15 15 I de 17 18 18 ba 19 tha 20 inc	ays that he used?	9	intensities and peak heights on the resulting			
12 tra 13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 22 23 24 22 23 24 25 wir 1 exp 2 0 3 ine 4 sec 5 Mis 7 8 8 inet 10 cal 11 BY 12 res 13 res 14 15 15 I du 16 bas 17 18 18 ba 19 tha 20 inc	A. It seems to be the same polymerase.	10	fluorograms, thus, the software has more difficulty			
13 wh 14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 23 24 23 24 25 wi 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 9 inte 10 cal 11 BY 12 10 13 res 14 15 15 I de 17 18 19 tha 20 inc	Q. Please go back to that exhibit, the	11	calling correct bases."			
14 15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 inet 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	ranscript, Exhibit 2084 and turn to the next page,	12	That is what the transcript states,			
15 16 Do 17 the 18 mi 19 rea 20 se 21 flu 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 inef 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 17 18 ba 19 tha 20 inc	which is page 13.	13	right?			
16 Do 17 the 18 mi 19 rea 20 se 21 flu 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 17 18 18 ba 19 tha 20 inc	A. I'm sorry, which? 2084?	14	A. That is what is written, yes.			
17 the 18 mi 19 rea 20 se 21 flu 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	Q. Yes, page 13 is the next page in that.	15	Q. So the NEB transcript teaches that adding			
17 the 18 mi 19 rea 20 se 21 flu 22 23 24 25 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	Do you see that? And please draw your attention to	16				
19 rea 20 se 21 flu 22 23 24 25 25 win 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 17 18 18 ba 19 tha 20 inc	the middle column. And towards the bottom of the	17	inefficient incorporation by Vent polymerase,			
19 rea 20 se 21 flu 22 23 24 25 25 win 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 17 18 18 ba 19 tha 20 inc	niddle column on page 13, there is a header which	18				
21 flu 22 23 24 25 wir 1 exp 2 0 3 iner 4 sec 5 6 6 Mis 7 8 9 inter 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 19 tha 20 inc	eads "Can the CircumVent kit be used for	19				
21 flu 22 23 24 25 wir 1 exp 2 0 3 iner 4 sec 5 6 6 Mis 7 8 9 inter 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 19 tha 20 inc	sequencing templates to be used on automated	20	describe what the modifications are, but they say			
22 23 24 25 wir 1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	luorescent sequencers."	21	that inefficient incorporation can lead to			
23 24 25 wi 1 exp 2 0 3 ine 4 sec 5 6 Mis 7 8 ine 9 inte 10 cal 11 BY 12 13 res 14 15 I do 15 I do 16 bas 17 18 ba 19 tha 20 inc	Do you see that?	22	•			
24 25 wir 1 exp 2 0 3 iner 4 sec 5 6 6 Mis 7 8 9 inter 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	A. I do.	23	-			
25 wir 1 exp 2 0 3 iner 4 sec 5 6 6 Mis 7 8 9 inter 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 19 that 20 incertail	Q. The NEB transcript answers that question	24	,			
1 exp 2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 inel 9 intel 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	with a yes, but with a caveat. Now, please draw	25				
2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 ine 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc			<u> </u>			
2 0 3 ine 4 sec 5 6 6 Mis 7 8 8 ine 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	Page 127 xploring, yes.	1	Page 128 the September 1992 transcript, and it teaches that			
3 ine 4 sec 5 6 6 Mis 7 8 8 inel 9 inte 10 cal 11 BY 12 13 13 res 14 15 15 I de 16 bas 17 18 19 that 20 inc	Q. And the transcript states that, that	2	fluorescently labeled nucleotides are inefficiently			
4 sec 5 6 6 Mis 7 8 8 inel 9 intel 10 cal 11 BY 12 13 14 15 15 I de 16 bas 17 18 19 that 20 inc	nefficient incorporation leads to unreliable	3	incorporated by the same Vent polymerases Metzker			
5 6 Mis 7 8 inet 9 inte 10 cal 11 BY 12 13 res 14 15 I do 16 bas 17 18 ba 19 tha 20 inc	equencing. That's what it says, right?	4	used in those experiments, correct?			
6 Mis 7 8 inef 9 inte 10 cal 11 BY 12 13 res 14 15 I du 16 bas 17 18 ba 19 tha 20 inc	MR. ZIMMERMAN: Objection.	5	A. The polymerase is the same, so given the			
7 8 inef 9 inte 10 cal 11 BY 12 13 res 14 15 I do 16 bas 17 18 ba 19 tha 20 inc	lischaracterizes.	6	caveats that we just described, there can be some			
8 inef 9 inte 10 cal 11 BY 12 13 res 14 15 I do 16 bas 17 18 ba 19 tha 20 inc	THE WITNESS: The document says	-	fluorophore link or modifications that reduce			
 9 inte 10 cal 11 BY 12 13 res 14 15 I de 16 bas 17 18 ba 19 tha 20 inc 	efficient incorporation leads to different	8	efficiency and complicate base calling, according			
10 cal 11 BY 12 13 res 14 15 1 da 15 1 da 16 bas 17 18 ba 19 tha 20 inc	Itensities and peak heights and more difficulty in	9	to NEB.			
11 BY 12 13 res 14 15 I do 16 bas 17 18 ba 19 tha 20 inc	calling correct bases.	10	Q. Given that Metzker reports that the			
12 13 res 14 15 15 I do 16 bas 17 18 18 ba 19 tha 20 income	3Y MR. SCHWARTZ	11	3'-O-allyl dATP without a label is capable of			
13 res 14 14 15 I de 16 bas 17 18 18 ba 19 tha 20 inc	Q. So would you say that that is what	12	incomplete termination even at high nucleotide			
14 15 I du 16 bas 17 18 ba 19 tha 20 inc	results in unreliable sequencing?	13	concentrations, a person of ordinary skill in the			
15 I do 16 bas 17 18 ba 19 tha 20 inc	A. You have as much information about it as	14				
16 bas 17 18 ba 19 tha 20 inc	do. It says that difficulty in calling correct	15	that nucleotide could result in even less			
17 18 ba 19 tha 20 inc	Dases.	16				
18 ba 19 tha 20 inc	Q. So please go back to Metzker 1994, and go	17	A. No, I disagree with that. He has not			
19 tha 20 inc	back to table two. Metzker reports in table two	18	reported that the 3'-O-allyl is inefficiently			
20 inc	hat the unlabeled 3'-O-allyl nucleotide	19	incorporated.			
	ncompletely terminated the sequencing reaction	20	-			
21		21	Q. No, I said incomplete termination.			
	using Vent (exo-) DNA polymerase at 250 micromolars		A. That's inefficient incorporation. So he			
	of the nucleotide, correct?	22	has not reported that it, on its own, is			
	A. Under the conditions that he is using	23	inefficient at termination. As we just discussed,			
	nere, that is what the document says.	24	•			
25	Q. And we just looked at the NEB transcript,	25	Q. But if you add a fluorescent label to			

FIO	d Romesberg, Ph.D.		University in the City of New York
1	Page 129 that nucleotide, it could result in even less	1	Page 130 that the O-allyl modified dAPT at a 100 fold more
	termination than Metzker reported at the 250	2	concentration than unmodified dAPT was competing
	micromolar, correct?	3	with the incorporation of that unmodified dAPT,
4	A. That is if if the added linker, as was	4	because you are getting some termination.
	the case apparently with the NEB report, if the	5	Q. Please go back to page 63 of Metzker '94.
	added linker reduced the efficiency, then, yes,	6	And please look at the header "Terminator Screen"
	that would presumably, in an additive way, reduce	7	on the right side of the page, and please look at
	the termination. But, again, that is termination	8	the last sentence in that section that starts with
	in the presence of 1 percent dATP.	9	"Compounds 1, 8, 7."
10	Q. Yeah, I'm just referring to Metzker's	10	Do you see that?
11	studies.	11	A. I do.
12	A. As am I, and his Metzker study has	12	Q. And Metzker writes "Compounds 1, 8, and 7
13	1 percent dATP in the mixture. No one would	13	showed specific termination and were further
14	sequence that way. No one would include dATP that	14	evaluated with respect to their
15	you're allowing to compete with your modified dATP	15	concentration-dependent effects."
16	where you're relying on the modified dAPT	16	That is what Metzker writes?
17	terminate.	17	A. Correct.
18	Q. Right. So the fact that there is that	18	Q. A person of ordinary skill reading
19	there is unlabeled, unmodified dAPT there would be	19	Metzker understood that Metzker was advancing
20	understood to change the way that one would expect	20	compounds 1, 8, and 7 for further evaluation based
21	that nucleotide to behave if that unmodified dATP	21	on the screening tests that were reported at
	wasn't there, correct?	22	least based on the screening tests that were
23	A. A POSA would understand that having that	23	reported in table two, correct?
24	contaminating dAPT would dramatically change this,	24	A. At least, but that data was available to
25	yes. I guess just to be clear, a POSA would know	25	him when he made the choice to conduct further
	Page 131		Page 132
	experiments concentration-dependent effects with 1,	1	Q. But he didn't move 3 forward?
2	8, and 7.	2	A. He did not move 3 forward.
3	Q. And a person reading a person of	3	Q. Metzker 1994 did not describe any
	ordinary skill reading Metzker 1994 also understood	4	condition to remove the 3'-O-allyl group, correct?
5	that, based on those screening tests, Metzker did	5	A. That is correct.
6	not advance compound 3 for further evaluation,	6	Q. He certainly didn't describe any
7	correct?	7	condition to remove the 3'-O-allyl group that would
8	A. No. If you say based on at least those	8	be compatible with DNA synthesis, right or
9	screening effects, then I might agree with it.	9	DNA let me strike that.
10	Q. Let me change the question then.	10	Metzker 1994 did not describe any
11	A person of ordinary skill reading	11	condition to remove the 3'-O-allyl group that is
12	Metzker 1994 understood that at least based on the	12	compatible with DNA, correct?
13	screening test, Metzker did not advance compound 3	13	A. That is correct. And in fact, I think
14	for further evaluation?	14	that's probably related to why he did not move 3
15	A. So just to be clear, because I'm not sure	15	forward. Because he had the conditions to easily
16	I understand that turn of the phrase "at least	16	deprotect 7. He wasn't the chemist. So he
17	based on," he had that information available. He	17	probably didn't think about reactions to
18	certainly had other information available too. And	18	deprotect 3.
19	he chose to move forward with 1, 8, and 7, and what	19	Q. But he didn't describe any conditions?
20	you are getting at, obviously, is that he did not	20	A. That is right.
21	move 3 forward, and he had the data from his	21	Q. Six lines down in the abstract, which is
22	kinetic data that's reported in table two, and he	22	on page we read the first part of the abstract,
23	had a world of other data as well. And I don't	23	and I'm just going to read the next part of the
24	know what it was that fed into his choice to not	24	abstract there is a sentence that starts with
25	move 3 forward.	25	"3'-O 2-nitrobenzyl dAPT."
1			-

	a Romesberg, Ph.D.	University in the City of New York	
1	Page 133 Do you see that?	1	Page 134 MR. SCHWARTZ: Do you want to take a
2	A. I do.	2	break for lunch at any particular time? I don't
3	Q. And Metzker writes "3'-O- 2-nitrobenzyl	3	know. What are your plans?
4	dAPT is a UV sensitive nucleotide, and was shown to	4	MR. ZIMMERMAN: When are you hungry?
5	be incorporated by several thermostable DNA	5	THE WITNESS: I generally don't eat big
6	polymerases. Base-specific termination and	6	lunches. So whenever anyone else is hungry.
7	efficient photolytic removal of the 3' protecting	7	MR. SCHWARTZ: Maybe we should break now.
	group was demonstrated. Following deprotection,	8	MR. ZIMMERMAN: If there is a natural
		-	
9	DNA synthesis was reinitiated by the incorporation	9	stopping point or a place, we are not wedded to a
10	of natural nucleotides into DNA. The	10	particular time.
11	identification of this labile terminator and the	11	THE VIDEOGRAPHER: We are off the record
12	demonstration of one-cycle stop/start DNA synthesis	12	at 12:16 p.m.
13	are initial steps in the development of a novel	13	(Recess taken.)
14	sequencing strategy."	14	THE VIDEOGRAPHER: We are on the record
15	That is what Metzker writes, correct?	15	at 1:19 p.m.
16	A. That is correct.	16	BY MR. SCHWARTZ:
17	Q. So Metzker advanced a nitrobenzyl	17	Q. Good afternoon, Dr. Romesberg. I am
18	nucleotide for further testing and demonstrated it	18	going to put in front of you another exhibit. That
19	was capable of one complete cycle of SBS, correct?	19	doesn't require marking. I'm putting in front of
20	A. I think that is precisely why he advanced	20	you Exhibit 107. It is entitled "Termination of
21	it, because it was easy to show it would be capable	21	the DNA Synthesis by N6-Alkylated, Not
22	of that complete step.	22	3'-O-Alkylated Photocleavable 2'-Deoxyadenosine
23	Q. And Metzker didn't advance the 3'-O-allyl	23	Triphosphates," by Wu, et al.
24	cap nucleotide for SBS testing?	24	You reviewed this paper in forming your
25	A. He did not.	25	opinion in the current IPRs, correct?
	Page 135		Page 136
1	A. That is correct.	1	BY MR. SCHWARTZ:
2	Q. The corresponding author of this paper is	2	Q. DNA polymerase?
3	Michael Metzker, correct?	3	MR. ZIMMERMAN: Yes.
4	A. Correct.	4	MR. SCHWARTZ: I thought I did say DNA.
5	Q. And that's the same Michael Metzker who	5	THE WITNESS: That is what is written.
6	is the first author of Metzker 1994, correct?	6	BY MR. SCHWARTZ:
7	A. That is correct.	7	Q. And the authors cite to reference 8 as
8	Q. Please turn to page 6348 of the paper,	8	support for those statements, correct?
9	and please look at the discussion section on the	9	A. They do.
10	left side. Please draw your attention to seven	10	Q. And can you take a look at the reference
11	lines down in the discussion section where the	11	list. And reference 8 is the Metzker '94 paper
12	sentence starts with "This is unlike the situation	12	we've been looking at, correct?
13	for."	13	A. That is correct.
14	Do you see that?	14	Q. So the Wu paper which includes Metzker as
15	A. I do.	15	an author refers back to the screening test
16	Q. And the author writes "This is unlike the	16	reported in Metzker 1994, correct?
17	situation for 3' modified nucleotides which	17	A. That is correct.
18	typically act as poor substrates for DNA	18	Q. And the Wu authors which include Metzker
19	polymerases. For example, screening 3'-O-allyl	19	characterize the screening test in Metzker '94 as
	dATP with eight different polymerases revealed	20	showing that the allyl capped nucleotide had
20			
20 21	limited activity at high micromolar concentrations,	21	limited activity at high micromolar concentrations
		21 22	limited activity at high micromolar concentrations with only one out of eight polymerases tested,
21	limited activity at high micromolar concentrations,		
21 22	limited activity at high micromolar concentrations, with only Vent (exo-) DNA polymerase."	22	with only one out of eight polymerases tested,
21 22 23 24	limited activity at high micromolar concentrations, with only Vent (exo-) DNA polymerase." That is what the authors wrote, correct?	22 23	with only one out of eight polymerases tested, correct?
1	Page 137 of the fact that there were contaminants in the	1	Page 138 BY MR. SCHWARTZ:
---	---	--	--
2	1994 studies, correct?	2	Q. That is what they said, correct?
3	A. That is correct.	3	A. They say that, yes.
4	Q. So despite the fact that they were aware	4	Q. And a person reading Metzker 1994 in
5	of contaminants in the 1994 studies, even years	5	October of 2000 understood that the 3-O-allyl ATP
6	later Metzker concluded that the 3'-O-allyl dAPT	6	had only limited activity at high micromolar
7	had limited activity with Vent at high micromolar	7	concentrations with only one of the eight
8	concentrations.	8	polymerases?
9	That is what he says, right?	9	A. I would disagree with that.
10	A. He says that. I just, by way of	10	Q. So you disagree with Metzker's
11	clarification, I think I don't know that I would	11	interpretation of his own data?
12	call that a conclusion. He is using sort of	12	A. Lam.
13	justification for what he is doing. He is	13	Q. But you do agree that Metzker himself
14		14	stated in 2007 that the O-allyl had limited
15	he is then saying that the that all the old ones	15	activity high micromolar concentrations?
16	were not incorporated efficiently. Which,	16	A. I agreed that he stated what you read.
17	incidentally, I disagreed with, but that is not	17	Q. And the Wu authors also stated that "the
18	your question.	18	3-O-allyl dAPT is a poor substrate for DNA
19	Q. And the Wu authors, which include	19	polymerase." That's what they said?
20	Metzker, characterize the screening test in Metzker	20	A. I agree that what you read is out of
21	'94 as showing that the allyl cap nucleotide had	21	their paper.
22	limited activity at high micromolar concentration	22	Q. And they said that knowing that in their
23	with only one of the eight polymerases?	23	1994 paper they had some contaminate in each of
24	MR. ZIMMERMAN: Objection. Asked and	24	their nucleotides, correct?
	answered.	25	A. I guess I don't know that the first
			-
			Page 140
1	Page 139 author knew that, because I don't think he was in	1	Page 140 Page 140
1	author knew that, because I don't think he was in	1	papers?
2	author knew that, because I don't think he was in the group at the time that that paper was studied,	1 2 3	A. I have.
2 3	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was.	2	papers?A. I have.Q. Is it your practice to read the papers
2 3 4	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have	2 3	papers?A. I have.Q. Is it your practice to read the papersthat you are corresponding author on?
2 3	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he	2 3 4	papers?A. I have.Q. Is it your practice to read the papers
2 3 4 5	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but	2 3 4 5	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on.
2 3 4 5 6	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those	2 3 4 5 6 7	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if
2 3 4 5 6 7 8	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct?	2 3 4 5 6 7	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a
2 3 4 5 6 7	 author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct? A. I don't know who wrote the paper. Wu is 	2 3 4 5 6 7 8	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a corresponding author, that you would permit
2 3 4 5 6 7 8 9	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct?	2 3 4 5 6 7 8 9	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a
2 3 4 5 6 7 8 9 10	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct? A. I don't know who wrote the paper. Wu is first author on the paper, Metzker is corresponding	2 3 4 5 6 7 8 9 10	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a corresponding author, that you would permit information that you did not believe to be accurate
2 3 4 5 6 7 8 9 10 11	author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct? A. I don't know who wrote the paper. Wu is first author on the paper, Metzker is corresponding author.	2 3 4 5 6 7 8 9 10 11	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a corresponding author, that you would permit information that you did not believe to be accurate to be in that paper?
2 3 4 5 6 7 8 9 10 11 11	 author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct? A. I don't know who wrote the paper. Wu is first author on the paper, Metzker is corresponding author. Q. Let's just make it simple. That is what 	2 3 4 5 6 7 8 9 10 11 12	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a corresponding author, that you would permit information that you did not believe to be accurate to be in that paper? A. No.
2 3 4 5 6 7 8 9 10 11 12 13	 author knew that, because I don't think he was in the group at the time that that paper was studied, or that paper was written, Metzker certainly was. I certainly think that that first author would have had access to read the paper. Do I know that he thought about that, no, but Q. But the Wu authors do characterize those studies that way, correct? A. I don't know who wrote the paper. Wu is first author on the paper, Metzker is corresponding author. Q. Let's just make it simple. That is what the paper says, right? 	2 3 4 5 6 7 8 9 10 11 12 13	 papers? A. I have. Q. Is it your practice to read the papers that you are corresponding author on? A. I write the papers that I'm corresponding author on. Q. And is it your practice that, even if there are other authors on the paper where you're a corresponding author, that you would permit information that you did not believe to be accurate to be in that paper? A. No. Q. So would you agree that corresponding
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2 3 4 5	Page 141		
2 3 4 5	that they compete with natural dAPT for	1	Page 142 choice based on the photodeprotection. So I don't
3 4 5	incorporation. Metzker seems to characterize that	2	think that he's mischaracterizing things. I just
4 5	as "not great." I would characterize that as	3	wouldn't characterize I just wouldn't use the
5	surprisingly good. I guess it's a matter of	4	perspective that he I would not take the
n	perspective or taste, I guess. I don't think he	5	perspective he is taking. BY MR. SCHWARTZ:
	was lying or falsifying anything. There is no	6	
	quantitative data there that's you know, it's he	7	Q. But you would agree that he did interpret
	simply says he uses a qualitative term that's	8	his 1994 data as saying that the 3-O-allyl ATP is a
	not wrong. You can buy again, I don't	9	poor substrate for DNA polymerase.
10	particularly like the qualitative way he describes	10	A. I would not agree with that. Those are
11	it, but what I'll also say is this is life.	11	the words he uses. It is attempting to take words
12	People do this. People use qualitative	12	so literally.
13	statements to justify where they're at in the	13	Q. I am only asking you if Metzker is saying
14	program right then. He had chosen, I think,	14	
15	probably because he was seduced into this idea that	15	A. He wrote those words. And that is not
16	you could photodeprotect very easily a base	16	how a person of ordinary skill in the art would
17	protected analog sorry, a 3' protected analog.	17	have written them.
18	That turned out to be incorrect, as I know you	18	Q. Those are the words in his paper?
19	know.	19	MR. ZIMMERMAN: Objection. Asked and
20	And I think he was enticed into that	20	answered.
21	approach because he thought it was so easy to	21	THE WITNESS: Those are the words in his
22	deprotect with light. And so that is why, in fact,	22	
23	I am almost certain moved compound 8 on in 1994,	23	(Exhibit 2025 was previously marked for
24	not because it was particularly better as a	24	identification.)
25	substrate than O-allyl. I think he made that	25	///
4	Page 143 BY MR. SCHWARTZ:	4	Page 144
1 2		1 2	 Q. It's about halfway down. A. Yes, I found it.
	Q. I am going to put in front of you another document. I'm putting in front of you	2	Q. You are with me?
	Exhibit 2025, a document entitled "Improved	4	A. Lam.
	Nucleotide Selectivity and Termination of 3'-OH	5	Q. And the Litosh authors write "We note
	Unblocked Reversible Terminators by Molecular	6	that 3'-O-allyl dAPT is not efficiently
	Tuning of 2-nitrobenzyl Alkylated HOMedU	7	incorporated by Vent (exo-) polymerase."
	Triphosphate." And, again, the corresponding	8	That is what the authors wrote, correct?
	author of this paper is Michael Metzker, correct?	9	A. That is what they wrote.
10	A. That is correct.	10	Q. And they refer to reference 46 to support
11	Q. And this is the same Michael Metzker who	11	that statement, right?
12	is the first author of Metzker '94, correct?	12	A. They do.
13	A. It is.	13	Q. Reference 46 is the Metzker 1994 paper,
14	Q. Please turn to Page 10 of the paper.	14	isn't it?
15	Please draw your attention to the first full	15	A. It is.
16	paragraph on the left side of the page that starts	16	Q. So the authors of the Litosh paper, which
10	with "The Use of HOMedU."	17	includes Michael Metzker as the corresponding
17	Do you see that?	18	author, states that the 3'-O-allyl dAPT is not
	A. Yes, I do.	19	efficiently incorporated by Vent (exo-) polymerase,
17	Q. Will you please look a little over	20	correct?
17 18			
17 18 19	halfway down that paragraph where there is a	21	A. Those are the words your reading of
 17 18 19 20 21 22 	halfway down that paragraph where there is a sentence that starts "We note 3-O-allyl dAPT is."	22	those words is correct.
 17 18 19 20 21 22 23 	sentence that starts "We note 3-O-allyl dAPT is." Do you see that?		those words is correct. Q. And they refer to the Metzker 1994 paper
 17 18 19 20 21 22 23 24 	sentence that starts "We note 3-O-allyl dAPT is."	22	those words is correct.

· · · · ,	a Romesberg, Ph.D.		
1	Page 145 Q. And again, they made that statement in	1	Page 146 correct?
2	this paper, which was published in 2011, aware of	2	A. I don't know that he said that. I would
3	the fact that they had some contamination of the	3	have to look to see that he said that, but the
4	nucleotides that were used in the Metzker 1994	4	Q. He says that it is not efficiently
5	studies, correct?	5	incorporated in 2011. And he says in 2007 that
6	A. I assume they were still aware of that.	6	it's a poor substrate for the polymerase, correct?
7	I don't know, maybe he wanted to forget it. It's	7	A. So he doesn't refer to sequencing by
8	frankly, it's a mistake on his part, and he may	8	synthesis as a whole, but, you know, this is why we
9	have not been proud of that.	9	qualify people's opinions as having to be a person
10	Q. So you're putting yourself into Metzker's	10	of ordinary skill in the art. A person of ordinary
11	mind now?	11	skill in the art, I imagine, would have been
12	A. No, I'm thinking of how I would feel if I	12	confused by these statements.
13	had done that, and I'm just feeling how most people	13	
14	would have. He is drawing conclusions that are	14	A. A person of the ordinary skill in the art
15	really not, in my opinion, founded. And I think	15	would have seen 1 percent contamination. A person
16	they wouldn't have been to the person of ordinary	16	of ordinary skill in the art would have understood
17	skill in the art as well.	17	that dAPT would be very efficiently incorporated at
18	Q. We looked at two papers, post 1994, where	18	its natural rate, and would therefore be
19	Michael Metzker is the corresponding author, and in	19	contributing to the lack of termination mediated by
20	both of those papers he is interpreting the	20	the protected analog. It is sophomore organic
21	experiment in the 1994 paper, correct?	21	chemistry. I think that anyone practicing in that
22	A. That is correct.	22	field would have been interested in that, and then
23	Q. And he is interpreting those experiments	23	maybe if they then read these papers been confused
24	to conclude that the 3'-O-allyl dATP was not	24	by Metzker stepping away from it.
25	suitable for his studies as a DNA terminator,	25	And again, I think what they would have
	Dave 447		B 446
	Page 147		Page 148
1	concluded is he is interested in these other guys,	1	BY MR. SCHWARTZ:
1 2	concluded is he is interested in these other guys, and he is passing this aside because this is now	1 2	BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA?
	concluded is he is interested in these other guys, and he is passing this aside because this is now his direction.		BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA? A. Yes.
2	concluded is he is interested in these other guys, and he is passing this aside because this is now his direction.Q. This is all speculation on your part,	2	 BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA? A. Yes. Q. You consider Metzker a POSA, and Metzker
2 3 4 5	 concluded is he is interested in these other guys, and he is passing this aside because this is now his direction. Q. This is all speculation on your part, isn't it? 	2 3 4 5	BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA? A. Yes. Q. You consider Metzker a POSA, and Metzker as a POSA wrote that the 3'-O-allyl dAPT is not
2 3 4 5 6	 concluded is he is interested in these other guys, and he is passing this aside because this is now his direction. Q. This is all speculation on your part, isn't it? A. It is. It is. 	2 3 4 5 6	 BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA? A. Yes. Q. You consider Metzker a POSA, and Metzker as a POSA wrote that the 3'-O-allyl dAPT is not efficiently incorporated by a Vent polymerase,
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 3 24	 concluded is he is interested in these other guys, and he is passing this aside because this is now his direction. Q. This is all speculation on your part, isn't it? A. It is. It is. Q. Do you have any evidence that persons skilled in the art actually believed in what you just said? A. Yes, because what I just said is sophomore chemistry. It's basic chemistry. And that is why we put a certain burden my understanding is that's why we put a certain limit on what we define as a person of ordinary skill in the art. And I think a person of ordinary skill in the art would have fully appreciated the natural substrates for polymerase are very efficiently incorporated. Q. But Metzker himself who had even extraordinary skill in the art would you agree that he had more than a person of ordinary skill in the art? How would you characterize that? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	BY MR. SCHWARTZ: Q. Do you consider Metzker a POSA? A. Yes. Q. You consider Metzker a POSA, and Metzker as a POSA wrote that the 3'-O-allyl dAPT is not efficiently incorporated by a Vent polymerase, right? A. So if you really want to talk about why he said this, I think it's clear. First of all, efficiently is a relative term, and so he's not saying anything wrong, and he is justifying the direction of research that he's going. He's got NIH grants that describe how great his approach is. Unfortunately, everyone in science has to do this. We have to justify what we are doing relative to other things. You're right. This is all speculation. I don't know why he wrote those words, but it is not speculation that there was 1 percent dAPT in those reactions and that that 1 percent dAPT led to two and a half micromolar of the natural substrate. That's a concentration that is typical of a standard PCR reaction. So it's

	va Romesberg, i n.b.		Oniversity in the only of New Tork
1	Page 149 something that would not have been obvious to a	1	Page 150 efficiently incorporated by the polymerase?
2	POSA.	2	A. Just because I would like to understand
3	Q. And when Metzker wrote in the other	3	the context in which I said that, where are you
4	paper, the 2007 paper, that the 3'-O-allyl dAPT is	4	reading that?
5	a poor substrate for DNA polymerase, that was also	5	Q. We can go back. We already have
	Metzker as a POSA writing that, correct?		
6 7	A. Yeah. And if you want me to speculate	6	testimony on this, but we can go back to your
		7	deposition transcript in IPR2013-00517. I forget
8	why I think he might have written that, speculate,	8	what exhibit that is.
9	I'm happy to do so.	9	A. I feel like I can answer the question now
10	Q. I'm not asking you to speculate. I'm	10	without looking further. That would be in the
11	only asking you to answer my question.	11	context of a sequencing by synthesis method.
12	A. He wrote that. And I would consider him	12	Q. Let's go back to table two of Metzker
13	a POSA.	13	1 0
14	Q. And earlier we looked at Tsien, and you	14	A. I am there.
15	agreed that one of Tsien's specific requirements is	15	
16	efficient incorporation of the 3' block nucleotide	16	
17	by the polymerase, correct?	17	A. That is correct.
18	A. Correct.	18	Q. And compound 8 in table two is a
19	Q. You also agree that to be successful for	19	3'-O-Methyl dTTP, correct?
20	SBS, the nucleotide has to be efficiently	20	A. That is correct.
21	incorporated by the polymerase?	21	Q. The only difference between compound 1
22	A. Are you asking my opinion?	22	and compound 8 is the base, correct?
23	Q. No, you actually said it, I'm just asking	23	A. That is correct.
24	if you remember that it was your opinion that to be	24	Q. The base in compound 1 is an adenine.
25	successful for SBS, the nucleotide has to be	25	-
	Page 151		Page 152
1	Page 151 A. That is correct.	1	Page 152
1 2		1 2	
	A. That is correct.		right?
2	A. That is correct.Q. I want to focus on table two's results	2	right? A. That is correct.
2 3	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? 	2 3	right? A. That is correct. Q. But none of those four polymerases was
2 3 4	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. 	2 3 4	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine
2 3 4 5 6	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to 	2 3 4 5	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right?
2 3 4 5 6	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, 	2 3 4 5 6	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is
2 3 4 5 6 7	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? 	2 3 4 5 6 7	 right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying.
2 3 4 5 6 7 8	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. 	2 3 4 5 6 7 8	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in
2 3 4 5 6 7 8 9	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to 	2 3 4 5 6 7 8 9	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the
2 3 4 5 6 7 8 9 10	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to incorporate the methyl capped thymine nucleotide, 	2 3 4 5 6 7 8 9 10	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the methyl capped thymine analog and the methyl capped
2 3 4 5 6 7 8 9 10 11	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to incorporate the methyl capped thymine nucleotide, right? 	2 3 4 5 6 7 8 9 10 11	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the methyl capped thymine analog and the methyl capped adenine analog, correct?
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2 3 4 5 6 7 8 9 10 11 12 13 14	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to incorporate the methyl capped thymine nucleotide, right? A. That is what is reported. Q. Again, the only difference between those two Methyl capped nucleotides is that one had an 	2 3 4 5 6 7 8 9 10 11 12 13 14	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the methyl capped thymine analog and the methyl capped adenine analog, correct? A. That would seem to be the case. Q. Please turn to page 4266. Look at the right column. The second full paragraph that
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to incorporate the methyl capped thymine nucleotide, right? A. That is what is reported. Q. Again, the only difference between those two Methyl capped nucleotides is that one had an adenine base and the other had a thymine base, 	2 3 4 5 6 7 8 9 10 11 12 13 14 15	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the methyl capped thymine analog and the methyl capped adenine analog, correct? A. That would seem to be the case. Q. Please turn to page 4266. Look at the right column. The second full paragraph that starts with "Several unexpected results."
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	 A. That is correct. Q. I want to focus on table two's results for the AMV RT polymerase, okay? A. Okay. Q. The AMV RT polymerase was able to incorporate the methyl capped adenine nucleotide, correct? A. That is correct. Q. But the same polymerase was unable to incorporate the methyl capped thymine nucleotide, right? A. That is what is reported. Q. Again, the only difference between those two Methyl capped nucleotides is that one had an adenine base and the other had a thymine base, correct? A. That is correct. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	right? A. That is correct. Q. But none of those four polymerases was capable of incorporating the methyl capped adenine nucleotide, right? A. That would seem to be what the table is saying. Q. You agree none of the polymerases in table two were capable of incorporating both the methyl capped thymine analog and the methyl capped adenine analog, correct? A. That would seem to be the case. Q. Please turn to page 4266. Look at the right column. The second full paragraph that starts with "Several unexpected results." Do you see that? A. Yes.
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	d Romesberg, Ph.D.	University in the City of New York			
1	Page 153 nucleotide analog with one particular base, it may	1	Page 154 analogs. I don't think the people would have known		
2	or may not incorporate the similar analog	2	that is the case. I think they probably would have		
3	containing a different base, correct?	3	expected it, and I think that is probably why he is		
4	A. That is what is stated there. That is	4	calling it unexpected. He opens that paragraph by		
5					
	the conclusion that is obvious in the conclusion	5	saying "several unexpected results." And so I		
6	that is stated there.	6	think that is a bit of a surprise.		
7	Q. So is it fair to say that a polymerase's	7	I would also point out, again, that all		
8	ability to incorporate a nucleotide analog with one	8	of those O-methyl ATP reactions have dATP in there.		
9	base doesn't provide an expectation that that	9	So some of those dashes may have been because they		
10	polymerase will also incorporate a similar	10	were being outcompeted by the natural dAPT. So I		
11	nucleotide with a different base?	11	think a POSA would have viewed that conclusion		
12	A. I would probably have to disagree with	12	5		
13	that. It certainly doesn't prove that it is the	13	Q. That is your speculation?		
14	case, but I think a POSA would have expected it to	14	A. That is my speculation. It is not my		
15	be, because you are making a modification that is	15	speculation that those reactions had natural dAPT		
16	distant from you are talking about the	16	in there and those would have officially		
17	modification that change distant from the position	17	incorporated. Again, each of those reactions had		
18	you are modifying and most people think that a	18	1 percent of the natural dAPT in there, and it is		
19	Watson-Crick base pair is formed. It doesn't	19	not my speculation that a POSA would have		
20	particularly matter if it's a GC, AT, TA, CG, so I	20	recognized that as an efficient substrate. That is		
21	think most people would have dismissed that and	21	my definition of a POSA.		
22	assumed that it was sort of general.	22	Q. But you will agree that Metzker's own		
23	I mean it was very common for people to	23	studies demonstrate that just substituting a base		
23			with the same polymerase can result in completely		
	simply characterize one analog and sort of expect	24			
25	that to transfer, to be transferable to other	25	different results in terms of incorporation by the		
	Page 155	1	Page 156		
1	polymerase, correct?	1	and the methyl capped thymine nucleotide, a POSA		
2	A. No, I don't agree with that. I agree	2	and the methyl capped thymine nucleotide, a POSA understood it was possible that for the allyl		
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1	Page 157 Tertiary Alcohol of 3-C-Methyl-N-Acetyllactosamine	1	Page 158 Q. Now the scheme that is used to deprotect
2	Produces a Novel Analogue of the LeX-Trisaccharide"	2	the allyl in Qian is not performed in aqueous
3	by Qian that was published in 1998.	3	media, correct?
4	This is the same Qian paper that you	4	A. It is performed in methanol.
5	referred to in these IPRs, correct?	5	Q. And you agree that protecting groups for
6	A. That is correct, Qian.	6	use is SBS must be removable under aqueous
7	Q. Please turn to Page 2185. Please draw	7	conditions, correct?
8	your attention to scheme two. You refer to scheme	8	A. That is correct.
9	two to form your opinion that the allyl protecting	9	Q. And you just said this one, instead of
10	group is removed quantitatively, correct?	10	aqueous conditions, uses methanol?
11	A. Yes, that is correct.	11	A. That is what's reported.
12	Q. And the compounds in Qian are not	12	Q. And you testified earlier that methanol
13	nucleotides or nucleosides, correct?	13	-
14	A. That is correct. They are sugars, which	14	A. Neat methanol is a DNA denaturance. As
15	is closely related to a the sugar of the	15	long as to the extent that it dehydrates DNA, it
16	nucleotide of which the 3'-O-hydroxyl group is part	16	is a denaturer. There is no reason these reactions
17	of, meaning that that unit is a sugar. And these	17	could haven't been performed with added water. The
18	are sugars.	18	reaction scheme they're showing is absolutely
19	Q. These are disaccharides, correct?	19	compatible with water.
20	A. They're two sugars attached.	20	Q. So you are suggesting that somebody could
21	Q. Just so that we are understand each	21	modify this reaction scheme, is that what you are
22	other, a nucleotide used for sequencing does not	22	
23	have a disaccharide, does it?	23	A. I am suggesting it would have been
24	A. No, but a nucleotide used for sequencing	24	
25	has a sugar unit. These have sugar units.	25	
	· · ·		
1	Page 159 that this would have been equivalent in water or in	1	Page 160
	that this would have been equivalent in water or in	1	he skipped several sentences.
2	that this would have been equivalent in water or in water/methanol mixtures, if not pure water.	2	he skipped several sentences. MR. SCHWARTZ: I did.
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	vu Kolliesberg, Fli.d.		University in the City of New Tork
1	Page 161 was not did not have compatible 3' protecting	1	Page 162 that was compatible with DNA. A palladium chloride
2	group deblocking conditions.	2	mediated deprotection of an allyl group of an
3	A. If the reaction was done in neat	3	O-allyl would have been known that it could be done
4	pyridine, then I would have said then and I would	4	in water, and so when you see it done in methanol,
5	say now that that is not compatible with any DNA	5	you would have had a POSA would have had a high
6	based method.	6	expectation that that would be doable in water.
7	Q. So let's go back to the Qian paper. The	7	MR. SCHWARTZ: Can I just take a minute
8	Qian paper is using conditions that are not	8	to ask can I go off the record for a second,
9	compatible with SBS, correct?	9	please?
10	A. The difference is that in one case, the	10	THE VIDEOGRAPHER: This marks the end of
11	transferability to conditions that would be are	11	media No. 2 of the deposition of Floyd Romesberg
12	trivial and well known and commonly practiced. In	12	Ph.D. We are off the record at 1:57 p.m.
13	the other case, they were not.	13	(Recess taken.)
14	Q. But you agree that the conditions in the	14	THE VIDEOGRAPHER: On the record at
15	paper that you are referring to in your declaration	15	2:00 p.m. And this marks the beginning media No. 3
16	in these cases those conditions are not compatible	16	of the deposition of Floyd Romesberg Ph.D.
17	with SBS?	17	BY MR. SCHWARTZ:
18	MR. ZIMMERMAN: Objection. Asked and	18	Q. Dr. Romesberg, going back to the
19	answered.	19	reaction, the palladium reaction in Qian.
20	THE WITNESS: A POSA reading the	20	Palladium chloride is only soluble in methanol or
21	Loubinoux paper would have thought that is	21	an organic solvent, correct? It has to have an
22	unlike those conditions are unlike there	22	organic solvent?
23	would not have been there would not have been	23	A. It is not highly soluble in water, so to
24	high expectation of success of performing a	24	solve that problem, people had worked out additives
25	reaction of that particular reaction in a way	25	to get it into water. The most common one is a PEG
20		20	to get it into water. The most common one is a r EO
	Page 163		Page 164
1		1	
1	additive. Trucol is a PEG. Palladium chloride	1	not themselves compatible with SBS, correct?
2	additive. Trucol is a PEG. Palladium chloride forms polymers, and that precludes it from acting	2	not themselves compatible with SBS, correct? A. That is correct.
23	additive. Trucol is a PEG. Palladium chloride forms polymers, and that precludes it from acting as a catalyst because they precipitate, they become	2 3	not themselves compatible with SBS, correct?A. That is correct.Q. Let's go back to the Boss reference that
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	Yu Komesberg, Fil.D.		
1	Page 165 A. I still have the former one you gave me	1	Page 166 And you can have a calculator.
2	with very little written on it if that saves you	2	A. Look, as easy as this is, it is
3	the trouble.	3	surprisingly tricky to do on your feet in front of
-			
4	Q. Did you write anything on that paper?	4	when 10 people are watching you. If you tell me
5	We'll keep that paper, because I will	5	that is correct, I'll believe it. It sounds about
6	mark it at the end. And so we're going to walk	6	right. If you really want me to sit here and do
7	through this calculation for the concentration of	7	that freshman chemistry, I'll do it.
8	methanol in that reaction, okay? And I will	8	Q. Yeah, let's do it. 7.9
9	provide you with some information which I will	9	A. All right. I will do it.
10	represent to you is accurate, and your counsel, if	10	Q. 7.9 grams, when the molecular weight of
11	they think it is inaccurate, has a chance to take	11	32 grams per mole is .25 moles, correct?
12	that up with you on your redirect.	12	A. Is 7.9 divided by 32.
13	So why don't you write down that the	13	Q. And it's .25 grams.
14	molecular weight of methanol is 32 grams per mil.	14	A. You want me to check that?
15	I'm sorry, 32 grams per mole.	15	Q. Ido.
16	And the density of methanol is .79 grams	16	-
17	per mil. So 10 mils of methanol weighs 7.9 grams,	17	,
18	correct?	18	
19	MR. ZIMMERMAN: Do you need a calculator,	19	
20	Dr. Romesberg?	20	Q. The so the molarity of methanol in the
			-
21	THE WITNESS: No. 10 mils would be	21	mixture of 10 mils of methanol and two mils of
22	7.9 grams, that's correct.	22	water, that is to say you've got a total of .012
23	BY MR. SCHWARTZ:	23	liters of solution, would be .25 moles over .012
24	Q. The 7.9 grams, again, the molecular	24	,
25	weight of 32 grams per mole is .25 moles, correct?	25	A. That sounds right.
	Page 167		Dana 400
	i ago i ei		Page 168
1	Q. Would you do that calculation, .25 over	1	in freshman chemistry in 1984, so I guess I trust
1 2		1 2	
	Q. Would you do that calculation, .25 over		in freshman chemistry in 1984, so I guess I trust
2	Q. Would you do that calculation, .25 over 0.12?	2	in freshman chemistry in 1984, so I guess I trust my calculation. You got the same result, so I
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	yu Komesberg, Fil.b.		
1	Page 169 Q. It's reasonable though, could he he		Page 170 that seems to be what it is saying.
2 3	could be using that, correct? A. Could have.	2	Q. You actually relied on this paper in some
4	MR. SCHWARTZ: I'm going to mark another	4	of your IBS IPRs. A. That's why I said I have read it, but not
5	paper. This is going to require a marking. Court	5	in years. That is, I believe, 2014.
6	Reporter, this will be Exhibit 2085.	6	Q. You testified earlier that DNA primer
7	(Exhibit 2085 was marked for	7	templates that were used in SBS are more
8	identification.)	8	susceptible to denaturation than large pieces of
9	BY MR. SCHWARTZ	9	DNA.
10	Q. I'm putting in front of you a document	10	Do you remember that?
11	entitled "The Relationship of Structure to the	11	A. Yes.
12	Effectiveness of Denaturing Agents for	12	Q. Would you agree that the susceptibility
13	Deoxyribonucleic Acid" by Levine, et al. published	12	of DNA primary templates in an SBS reaction would
14	in 1973.	14	
15	Please turn to page 171 of the paper. On	15	
16	the top left side you see there is a table two?	16	A. To the extent that they are smaller, that
17	A. I see it.	17	-
18	Q. Table two is a list of reagents that were	18	Q. Now, first reagent listed in table two is
19	tested for their ability to denature DNA, correct?	19	methyl alcohol.
20	A. I see it.	20	Do you see that?
21	Q. And the DNA being tested here is	21	A. I do.
22	bacteria 5 DNA, correct?	22	Q. And that is the same as methanol,
23	A. That seems to be the case. I have not	23	
24	read this paper. Actually, I have read this paper,	24	A. It is.
25	surprisingly, but I haven't read it in years, so	25	Q. And table two indicates that 3.5 molar
1	Page 171 methanol denatures 50 percent of bacteriophage DNA	1	Page 172 page 169, about nine lines up from the bottom,
2	under the conditions of the assay, correct?	2	
3	A. That is correct.		•
-		3	73 degrees there is essentially."
4	Q. And the conditions of the assay are given	3 4	73 degrees there is essentially." Do you see that?
4	Q. And the conditions of the assay are given on page 168.	-	73 degrees there is essentially." Do you see that? A. do.
	on page 168.	4	Do you see that? A. I do.
5	on page 168. MR. ZIMMERMAN: To the extent you need	4 5 6	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees
5 6	on page 168.	4 5 6	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the
5 6 7	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do.	4 5 6 7	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent."
5 6 7 8	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ:	4 5 6 7 8	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the
5 6 7 8 9	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data	4 5 6 7 8 9	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct?
5 6 7 8 9 10	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as	4 5 6 7 8 9 10	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct.
5 6 7 8 9 10 11	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as	4 5 6 7 8 9 10 11	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two
5 6 7 8 9 10 11 12	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows?	4 5 6 7 8 9 10 11 12	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two
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5 6 7 8 9 10 11 12 13 14	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered	4 5 6 7 8 9 10 11 12 13 14	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They
5 6 7 8 9 10 11 12 13 14 15	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered aqueous conditions at close to neutral pH at	4 5 6 7 8 9 10 11 12 13 14 15	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They basically took the system to the brink of
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5 6 7 8 9 10 11 12 13 14 15 16 77	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered aqueous conditions at close to neutral pH at 73 degrees C. Do you see that?	4 5 6 7 8 9 10 11 12 13 14 15 16 17	Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They basically took the system to the brink of denaturation by heating it up. 73 degrees. That is the point.
5 6 7 8 9 10 11 12 13 14 15 16 17 18	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered aqueous conditions at close to neutral pH at 73 degrees C. Do you see that? A. I do.	4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	 Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They basically took the system to the brink of denaturation by heating it up. 73 degrees. That is the point. Q. But it says "at 73 degrees there is no
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5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered aqueous conditions at close to neutral pH at 73 degrees C. Do you see that? A. I do. Q. And if you go back to table two it indicates that the assay conditions were also done	4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They basically took the system to the brink of denaturation by heating it up. 73 degrees. That is the point. Q. But it says "at 73 degrees there is no denaturation of DNA"? A. They took it to the brink of
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5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24	on page 168. MR. ZIMMERMAN: To the extent you need time to verify that, please do. BY MR. SCHWARTZ: Q. The assay conditions for the data reported in table two of Levine. Do you see where it said denaturation of DNA was determined as follows? A. I do. Q. And it was 30 minutes under buffered aqueous conditions at close to neutral pH at 73 degrees C. Do you see that? A. I do. Q. And if you go back to table two it indicates that the assay conditions were also done at an ionic strength of 0.043. Do you see that? A. I do. Q. And if you go back to page 169, if you	 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 	 Do you see that? A. I do. Q. And the sentence reads: "At 73 degrees there is essentially no denaturation of DNA in the absence of a denaturing agent." That is what Levine wrote, correct? A. That is correct. Q. So the condition the reaction condition used to test the compounds in table two of Levine were not themselves denaturing, correct? A. No, that is not true at all. They basically took the system to the brink of denaturation by heating it up. 73 degrees. That is the point. Q. But it says "at 73 degrees there is no denaturation. Q. But they say that "at 73 degree there is essentially no denaturation in the absence of a denaturation in the absence of a denaturing agent."
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1	Page 173 A. None of the strands are dissociating,	1	Page 174 Mischaracterizes the document.
2	detectably. But they have the whole point of	2	You can answer.
3	elevating the temperature was to give them a regime	3	MR. SCHWARTZ: No, I'm asking about his
4	where they could see changes.	4	testimony.
5	Q. What temperature do you think is in	5	MR. ZIMMERMAN: I know, but the guestion
6	your opinion, what temperature does denature DNA?	6	mischaracterizes the document, so I'm going to make
7	A. It depends on the sequence, the sequence	7	the objection.
8	length, the sequence context. It's variable. It	8	You can answer.
9	depends on the conditions. 73 is up there.	9	BY MR. SCHWARTZ:
10	Q. Would 80 degrees denature DNA?	10	Q. The sentence reads: "At 73 degrees there
11	A. PCR reactions, which they have much	11	is essentially no denaturation of DNA in the
12	longer strands of DNA, routinely heat up to	12	absence of a denaturing agent." That is what it
13	90 degrees, 88 degrees, 89 degrees, and assume	13	says, right? And then you had a position about
14	complete denaturation in no methanol, in just	14	that that you said I think that that is close to
15	aqueous standard conditions that favor duplex	15	where DNA denatures. You were making a point,
16	formation. So 73 is a lot closer to 89 or 88 than	16	right?
17	it is to room temperature of 37. These strands are	17	A. Yes. The point that I was trying to
18	close to dissociating.	18	make and it wasn't for this specific strand,
19	Q. I mean it is your point that 73 degrees,	19	because I don't know the temperature at which it
20	even though Levine reports that there is no	20	would dissociate. 73 degrees what DNA duplex
21	denaturation of the DNA in the absence of a	21	is susceptible to heating. Heating is a
22	denaturing agent, your position is that that is	22	denaturation is a denature you have a free
23	close to the level of where DNA would be denatured,	23	energy that stabilizes a duplex. When you increase
	if I understood you correctly?	24	the temperature, the free energy stabilizing the
25	MR. ZIMMERMAN: Objection.	25	duplex is decreasing, and you can get that to an
			· · · · · · · ·
1	Page 175 arbitrarily small value, and it won't dissociate	1	is not speculation. Page 176
1	arbitrarily small value, and it won't dissociate	1	is not speculation.
2	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether	2	is not speculation. Q. But do you remember relying on this very
23	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature		is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS?
2 3 4	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature or addition of methanol or whatever sort of	2 3	 is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS? A. I do.
23	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature or addition of methanol or whatever sort of denature you are interested in, then makes that	2 3 4 5	 is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS? A. I do. Q. And you relied on this paper and this
2 3 4	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature or addition of methanol or whatever sort of denature you are interested in, then makes that free energy increasingly less favorable. And then	2 3 4 5	 is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS? A. I do. Q. And you relied on this paper and this very table to make conclusions about what agents
2 3 4 5 6 7	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature or addition of methanol or whatever sort of denature you are interested in, then makes that free energy increasingly less favorable. And then you will start to see the dissociation of that.	2 3 4 5	 is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS? A. I do. Q. And you relied on this paper and this very table to make conclusions about what agents would be DNA denaturing agents.
2 3 4 5 6 7 8	arbitrarily small value, and it won't dissociate yet. And then a little more denaturation, whether that be provided by further increase in temperature or addition of methanol or whatever sort of denature you are interested in, then makes that free energy increasingly less favorable. And then you will start to see the dissociation of that. So what they are doing is they are taking	2 3 4 5 6 7 8	 is not speculation. Q. But do you remember relying on this very paper and this very table in your IPRs against IBS? A. I do. Q. And you relied on this paper and this very table to make conclusions about what agents would be DNA denaturing agents. Do you remember that?
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	a Romesberg, Ph.D.		University in the City of New York
1	Page 177 So it is not disingenuous, and I would	1	Page 178 was that you needed, at any given temperature, less
2	still believe this, that if I look at that table,	2	pyridine to cause denature than most other things.
3	you see pyridine is 0.09 molar. That's much less	3	And that is true.
4	than methanol. And I believe my point was, and I	4	Q. And going back to the entry for methanol
5	would have to go look to rethink this if you want	5	that you just brought that up again, 3.5
6			methanol 3.5 molar methanol is reported in this
	me to be sure, but I believe my point was that	6	•
7	pyridine is a relatively strong denaturant. And I	7	paper to cause 50 percent denaturation of the
8	believe that conclusion is correct.	8	bacteriophage DNA in 30 minutes, correct?
9	Q. But you did rely on this paper and on	9	A. At 73 degrees.
10	that table to form that conclusion?	10	Q. Which the paper reports there was
11	A. In the way I just described it, yes.	11	essentially no denaturation of DNA at that
12	Which is not the same as the way we are looking at	12	temperature, the paper reports that, right?
13	it now. Just to be clear.	13	A. I'm not sure if you don't understand what
14	Q. And in what way is it different?	14	I'm saying or if you just want me to say it again.
15	A. I was comparing pyridine to other things.	15	They took this by heating it up to a temperature
16	Q. I haven't even asked you any questions	16	where the free energy stabilizing the duplex was
17	about this table yet, so all I've asked you is	17	almost taken to zero. That allowed them to add
18	what the reaction conditions were.	18	different denaturants and get dissociation so
19	A. No, you are asking me about methanol	19	everything they added, they would be able to see
20	being a denaturant, and that is shown there in the	20	denaturation. They wanted to do that so they can
21	concentration.	21	see how they all compared. That is one conclusion
22	Q. Yes.	22	you can draw from this table.
23	A. And that is in this table.	23	That was not the conclusion that I drew
24	Q. Yes.	24	from the table four years ago in that IPR that you
25	A. The only point that I was making before	25	evoked. The conclusion I drew at that time was
25	A. The only point that I was making before	25	evoked. The conclusion rulew at that time was
4	Page 179	4	Page 180
1	that pyridine under these given conditions is a	1	A. 3.5. Let me get my calculator. 20.8
2	stronger denaturant than methanol. Not that		
		2	divided by 3.5. I get 5.94.
3	pyridine would or wouldn't have denatured it at	3	Q. So that's almost a six fold difference,
4	these exact conditions.	3 4	Q. So that's almost a six fold difference, correct?
4 5	these exact conditions. Q. Does this paper report that 3.5 molar	3 4 5	Q. So that's almost a six fold difference, correct?A. Yeah, that's almost a six fold.
4	these exact conditions. Q. Does this paper report that 3.5 molar methanol caused 50 percent denaturation of the	3 4	 Q. So that's almost a six fold difference, correct? A. Yeah, that's almost a six fold. Q. So a POSA reading this paper, Levine,
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1109	a Romesberg, Ph.D.		University in the City of New York
1 2	Page 181 A. 20 molar methanol at room temperature? Q. 20.8 molar methanol.	1 2	Page 182 A. At 73 degrees. Q. Which the paper reports there was no
3	MR. ZIMMERMAN: Objection. Incomplete	2	denaturation without the denaturant, right?
	hypothetical.	4	A. I know you're making
5	THE WITNESS: Are you asking if it would	5	MR. ZIMMERMAN: Objection.
	denature DNA or if the entire process would be	6	Mischaracterizes the document again.
	compatible with that much methanol?	7	MR. SCHWARTZ: I disagree with your
	BY MR. SCHWARTZ:	8	saying that it mischaracterizes.
9	Q. Let's start with the later. Would the	9	MR. ZIMMERMAN: You're leaving out the
10	process be compatible with that much methanol?	10	word "essentially." Every time you characterize
11	A. Probably not. It would probably affect the	11	you leave out "essentially," so that is a
12	polymerase.	12	mischaracterization of the document.
13	Q. And at the bottom of table two there is	13	MR. SCHWARTZ: The document says that at
14	an entry for acetonitrile.	14	73 degrees there is essentially no denaturation in
15	Do you see that?	15	the absence of denaturing agent. That is what the
16	A. Bottom of table two.	16	document says, correct?
17	Q. Of that Levine paper.	17	THE WITNESS: Yes.
18	A. Oh, yes.	18	MR. SCHWARTZ: Can we take a break?
19	Q. And table two reports	19	Let's take a break here.
20	A. I'm sorry, acetonitrile? At the bottom?	20	MR. ZIMMERMAN: Sure
21	Q. At the bottom right.	21	THE VIDEOGRAPHER: We are off the record
22	A. Oh, yeah, sorry. Yes, I found it.	22	at 2:26 p.m.
23	Q. And table two reports that 1.2 molar	23	(Recess taken.)
24	acetonitrile causes 50 percent denaturation of	24	THE VIDEOGRAPHER: We are on the record
25	bacteriophage DNA, correct?	25	at 2:46 p.m.
	Page 183		Page 184
1	BY MR. SCHWARTZ:	1	BY MR. SCHWARTZ
2	Q. Dr. Romesberg, just to step back for a	2	Q. I am putting in front you, Dr. Romesberg,
	moment. We were referring earlier today about	3	a document entitled "Deoxyribonucleic Acids" by Kit
	allyl deprotections with palladium chloride in	4	that was published in 1963. And this is a paper
	water. Can you point me to any reference prior to	5	that you relied on in your declarations in the
	October of 2000 showing an allyl deprotection with		IPR2013-00517 case to form opinions about reagents
	palladium chloride in water?		that are DNA denaturants, correct?
8	A. No. I cannot. What I can show you is	8	A. I believe that is correct, yes.
	a I'm sure I could find papers that talk about	9	Q. And please turn to page 57 of Kit. And
10	using palladium chloride in water.	10	please draw your attention to the last paragraph on
11	Q. But can you point me to any paper that	11	the page, and five lines up from the bottom of the
12	reports deprotection of an allyl using palladium	12	page there is a sentence that starts with
13	chloride in water?	13	"Thiocyanate."
14	A. Not that I know of. In general,	14 15	Do you see that? A. I do.
15	palladium chloride deprotections of allyl were most used in sugar chemistry, and those substrates were	15 16	A. 100. Q. And Kit writes "Thiocyanate,
17	insoluble in water. People would not have desired	17	trichloroacetate, and perchlorate are more powerful
18	to run the reactions in water.	17	DNA denaturing agents than those prototypical
10	Q. I think I'm ready to go to another	10	hydrogen bond breakers, urea and clonidinian
20	exhibit.	20	chloride, and are matched only by certain organic
20	MR. SCHWARTZ: Court Reporter, this will	20	solvents such as formalyde, dimethylformalyde, and
22	be marked as Exhibit 2086.	22	dimethylsulfoxide."
22	(Exhibit 2086 was marked for	23	That is what Kit writes, correct?
23	identification.)	24	A. That is what is written.
25	,	25	Q. And Kit teaches that perchloric acid is a
			•

	yu Komesberg, Fil.D.		
1	Page 185 very powerful DNA denaturant, correct?	1	Page 186 really care if it is needed or not. Because
2	A. That is what is written there, yes.	2	they're sort of generic conditions.
3	Q. Please go back to the Boss reference that	3	And so I imagine that is what they were
4	was Exhibit 1035. And please go back to Table 1.	4	doing here.
5	And if you look at the legend of Table 1, the	5	Q. But clearly it indicates that perchloric
6	legend indicates that deprotection conditions for	6	acid is present in these deprotection reactions.
7	all of the compounds except compound 8 contained	7	MR. ZIMMERMAN: Objection. Asked and
8	perchloric acid, correct?	8	answered.
9	A. That is correct.	9	BY MR. SCHWARTZ
10	Q. And the acid is there because it is	10	Q. And, again, we looked at this before. I
11	required for deprotecting an allyl using palladium	11	know that you don't like doing calculations, so I
12	two, correct?	12	won't ask you to do it if you don't want to, but I
13	A. That is not correct.	13	would the concentration of perchloric acid
14	Q. There is acid there, correct?	14	that's listed there is either 1 millimolar to
15	A. There is acid, yes.	15	5 millimolar, correct?
16	Q. Okay.	16	A. I think that I remember reading that.
17	A. So let me be clear about that, in	17	Q. It's actually in the paper.
18	principle, it might have been important. The way	18	A. Okay.
19	people run these reactions if you look at the	19	Q. Do you see it? Turn it over and look at
20	mechanism, there is a spot where if you do not have	20	note a under the table.
21	a powerful nucleophile, your electrophile has to be	21	A. Yes.
22	made a better leaving group. And in those cases it	22	Q. And I want to calculate what the
23	might be beneficial to add a proton, very common	23	concentration of perchloric acid is when it's
24	for people if they were not conducting mechanistic	24	1 millimolar one millimole per 12 mils. That's
25	studies to simply add some acid because they don't	25	a simple calculation. I got 83 millimolars. Would
20		20	a simple calculation. I got oo minimolars. Would
	Page 187		Page 188
1		1	
1	you like to confirm that it is 83 millimolars?	1	1.1.
2	you like to confirm that it is 83 millimolars? A. No. I'll take your word for it.	2	1.1. A. Sounds about right.
2 3	you like to confirm that it is 83 millimolars?A. No. I'll take your word for it.Q. And 5 millimoles of perchloric acid in 12	2 3	1.1.A. Sounds about right.Q. And your counsel will have an opportunity
2 3 4	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. 	2 3 4	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417
2 3 4 5	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 	2 3 4 5	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4.
2 3 4 5 6	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? 	2 3 4 5 6	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay.
2 3 4 5 6 7	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the 	2 3 4 5 6 7	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the
2 3 4 5 6 7 8	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. 	2 3 4 5 6 7 8	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me?
2 3 4 5 6 7 8 9	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a 	2 3 4 5 6 7 8 9	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation.
2 3 4 5 6 7 8 9 10	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with 	2 3 4 5 6 7 8 9 10	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the
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2 3 4 5 6 7 8 9 10 11 12	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information 	2 3 4 5 6 7 8 9 10 11 12	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4.
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2 3 4 5 6 7 8 9 10 11 12 13 14	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton 	2 3 4 5 6 7 8 9 10 11 12 13 14	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1.
2 3 4 5 6 7 8 9 10 11 12 13 14 15	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration. So when the hydrogen when the 	2 3 4 5 6 7 8 9 10 11 12 13 14 15	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration?
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight?
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration. So when the hydrogen when the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that
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2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. Q. And likewise, when the hydrogen the proton concentration is .417 molar, the pH will be 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that down already. There were two concentrations of perchloric acid given in that figure.
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. Q. And likewise, when the hydrogen the proton concentration is .417 molar, the pH will be the minus log of .417, correct? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that down already. There were two concentrations of perchloric acid given in that figure. A. So it was 0.417 and 0.083 pH of 1.1 and
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. Q. And likewise, when the hydrogen the proton concentration is .417 molar, the pH will be the minus log of .417, correct? A. Yes. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that down already. There were two concentrations of perchloric acid given in that figure. A. So it was 0.417 and 0.083 pH of 1.1 and 0.84?
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. Q. And likewise, when the hydrogen the proton concentration is .417 molar, the pH will be the minus log of .417, correct? A. Yes. Q. We have a calculator that does logs? 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that down already. There were two concentrations of perchloric acid given in that figure. A. So it was 0.417 and 0.083 pH of 1.1 and 0.84? Q. Yes. So please go back to the Kit
2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24	 you like to confirm that it is 83 millimolars? A. No. I'll take your word for it. Q. And 5 millimoles of perchloric acid in 12 mils. I calculated it was 417 millimolars. Do you want to confirm that it is 417 millimolar? A. If I bought the first, I bought the second. Q. I am going to ask you now to do a calculation of pHs. And I will provide you with some information so use that table or use that piece of paper. The first piece of information I'll provide to you is that the formula for determining pH is pH equals minus log of the proton concentration is .083 molar, pH would be the minus log of .083, correct? A. That is correct. Q. And likewise, when the hydrogen the proton concentration is .417 molar, the pH will be the minus log of .417, correct? A. Yes. 	2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22	 1.1. A. Sounds about right. Q. And your counsel will have an opportunity to correct me if I am wrong. And the pH for .417 molar is about .4. A. Okay. Q. I mean, would you like to do the calculation to confirm me? A. No. It's a straightforward calculation. Q. So maybe you can write that down that the pH of .083 molar is about 1.1, and the pH of .417 molar is about pH of .4. A. What was the first number? Q. I'm sorry, 1.1. A. What was the hydrogen ion concentration? Q083. A. Zero eight? Q083. I think you may have written that down already. There were two concentrations of perchloric acid given in that figure. A. So it was 0.417 and 0.083 pH of 1.1 and 0.84?

	ya Komesberg, Fili.b.	1	
1	Page 189 there is a sentence that starts with "DNA is stable	1	Page 190 DNA is stable, correct?
2	over."	2	A. That is correct.
3	Do you see that?	3	Q. And pH of 0.4, the pH of perchloric acid
4	A. Yes.	4	at .417 molar is also outside the range of 2.7 to
5	Q. And Kit writes "DNA is stable over the	5	12 where DNA is stable, correct?
6	approximate pH range of 2.7 to 12	6	A. That is correct.
7	MR. ZIMMERMAN: That's not the same. You	7	Q. So the concentration of perchloric acid
8	said that is the start of the sentence?	8	that was used to deprotect the allyl in the Boss
9	MR. SCHWARTZ: I said about eight lines	9	deprotection reaction would denature DNA, correct?
10	down there is a sentence that starts with "DNA is	10	A. The Kit reference suggests that it would.
11	stable over."	11	Q. And that deprotection reaction in Boss
12	MR. ZIMMERMAN: There is not a sentence	12	would not be considered a mild condition that would
13	that starts with that.	13	not denature DNA, correct?
14	MR. SCHWARTZ: It's in the middle of the	14	A. That is correct.
15	sentence. I apologize.	15	Q. Let's switch topics for a minute for a
16		16	few minutes, and let's talk about depurination. If
17	Q. But do you see where I am?	17	it's been known before 2000 that glycosidic
18	-	18	linkages and purines are susceptible to acid
19	Q. And it says "DNA is stable over the	19	hydrolysis, correct?
20	approximate pH range of 2.7 to 12, but denatures at	20	A. That is correct.
21	higher or lower pHs."	21	Q. I am going to mark the paper.
22	That is what Kit writes, right?	22	(Exhibit 2057 was marked for
23	A. That is what's written.	23	identification.)
24	Q. So pH 1.1, the pH of perchloric acid at	24	MR. SCHWARTZ: This is, again, this is
25	0.83 molars is outside the range of 2.7 to 12 where	25	one of these papers that was present in only one
			Dave 400
1	Page 191 IPR, but not all the IPRs, so we're going to mark	1	Page 192 this one?
2	it to make sure it is in all the IPRs. And it is	2	MR. SCHWARTZ: This is marked as 2057.
3	2057.	3	BY MR. SCHWARTZ
4	MR. ZIMMERMAN: So in the case where	4	Q. So I'm putting in front of you a paper
5	we're not doing the deposition in that case today,	5	entitled "Instability and Decay of the Primary
6	and the case that you refused to do the deposition	6	Structure of DNA" by Lindahl, which was published
7	on that case today, you are marking the exhibits	7	in 1993. Now, please draw your attention to the
8	from that case.	8	first page of Lindahl. It's Page 709. And on the
9	MR. SCHWARTZ: Not the exhibits from that	9	left side of page 709 is a section entitled "DNA
10		10	Hydrolysis."
11		11	Do you see that?
12		12	A. Ido.
13		13	Q. And please draw your attention to the
14		14	sentence that begins five lines down. It starts
15		15	with "The instability of DNA."
16		16	Do you see that?
17		17	A. I do.
18		18	Q. Lindahl writes "The instability of DNA
19	5	19	glyclosal bonds under physiological solvent
20		20	conditions was investigated 20 years ago. It was
21	0	21	measured by using DNA with carbon 14-labeled purine
22		22	or pyrimidine residues and determining the rates of
23	, , , ,	23	release of free bases as a function of temperature,
24	•	24	pH, ionic strength, and nucleic acid secondary
25		25	
	- · · · · · · · · · · · · · · · · · · ·		

FIO	yd Romesberg, Ph.D.		University in the City of New Yor
1	Page 193 That is what Lindahl writes, correct?	1	Page 194 abstract. The abstract reads "The rate of
2	A. That is what is written.	2	depurination of double stranded bacillus subtilis
3	Q. And Lindahl cites to references five	3	DNA radio actively labeled in the purine residues
4	through seven in support for that statement,	4	has been followed as a function of temperature, pH,
5	correct?	_	and ionic strength. In the magnesium containing
	A. That is correct.	5	
6		6	buffer of physiological ionic strength, the rate
7	Q. And please turn to the references list.	7	constant for depurination of DNA is 4 times 10 to
8	And reference five is a paper by Lindahl and Nyberg	8	the minus ninth per second at 70 degrees and pH
9	published in 1972, correct?	9	7.4."
10	A. That is correct.	10	That's what the authors write, correct?
11	MR. SCHWARTZ: I am going to mark that	11	A. That is what is written.
12	paper. That is going to be 2087, Court Reporter.	12	Q. Please turn to page 3616 of the paper,
13	(Exhibit 2087 was marked for	13	and please draw your attention to figure nine at
14	identification.)	14	the top of the page. Figure nine shows the pH
15	MR. ZIMMERMAN: Is this already of record	15	dependence of the rate of acid catalyzed
16	in any of the IPRs?	16	depurination of DNA, correct?
17	MR. SCHWARTZ: No.	17	A. It does.
18	BY MR. SCHWARTZ:	18	Q. And if you compare figure nine to the
19	Q. So I am putting in front of you a	19	statement we just read in the abstract, the rate of
20	document entitled "Rate of Depurination of Native	20	depurination reported in the abstract of four times
21	Deoxyribonucleic Acid" by Lindahl and Nyberg,	21	10 to the minus nine per second at pH 7.4 is
22	published in 1972. This is reference 5 in Lindahl	22	consistent with where the dotted line in figure
23	1993, correct?	23	nine intersects the X axis, correct?
24	A. It appears to be.	24	A. Yeah, I guess it is. It's a little hard
25	Q. Please direct your attention to the	25	to read, but I guess, yes.
	Page 195		Page 196
1	Q. And figure nine shows that as the pH	1	Q. We were just talking about the logs. We
2	becomes more the acidic, the rate of acid catalyzed	2	can do the calculation. I know you don't like to
3	depurination greatly increases, correct?	3	do calculations, but we're talking about four times
4	A. It increases, yes.	4	10 to the minus nine per second at pH 7.4?
5	Q. And if you drop from pH 7.4 to about	5	A. So four times 10 to the minus it's
6	pH 4, the rate of acid catalyzed depurination of	6	just that rapidly, again, is a relative term. That
7	DNA increases about three logs, correct?	7	is what it is.
8	A. That is correct.	8	Q. And go back to Boss, please,
9	Q. That is an increase of about a thousand	9	Exhibit 1035. Let's look at Table 1, and we
10	fold, correct?	10	calculated that the pH of the allyl deprotection
11	A. Three logs is a thousand fold.	11	reactions that use perchloric acid is between 1.1
12	Q. And if we were to extrapolate figure nine	12	and 0.4, correct? We put that on that piece of
13	to pH 1, the rate of acid catalyzed depurination of	13	paper?
14	DNA could increase by about another three logs?	14	A. The pHs, yes.
15	A. If that same linear relationship holds,	15	Q. So let me ask the question again so we
16	then that's correct.	16	5
17	Q. And that would be an increase of about a	17	We calculated that the pH of the allyl
18	million fold from pH 7.4, right?	18	deprotection reactions that use perchloric acid in
19	A. That would be correct.	19	Boss is between 1.1 and 0.4, correct?
20	Q. So Lindahl shows that DNA is rapidly	20	A. Yes.
21	depurinated in strong acid, correct?	21	Q. And Lindahl shows that those conditions
22	A. Well, it shows it down to pH 4.5.	22	will lead to rapid DNA depurination all the way
23	Q. And he shows that it's rapidly	23	down to pHs below 1, correct?
24	depurinated?	24	MR. ZIMMERMAN: Objection.
25	A. Rapidly?	25	Mischaracterizes the reference. Assumes facts not
1		1	

1	in evidence. Page 197	1	Page 198 A. So he doesn't show anything about pH 1,
2	THE WITNESS: Lindahl shows rates, I	2	
3	believe, down to pH 4.5, and the pH and it shows	3	Q. He does show that it would be depurinated
4	pH rates for that that vary from four times 10 to	4	at pH 4.5?
5	the minus nine, and up by about three orders of	5	A. Yes.
6	magnitude as we just discussed.	6	Q. And if you dropped it down to pH 1.1,
7	BY MR. SCHWARTZ:	7	would you also expect depurination at pH 1.1?
8	Q. And if we extrapolate from that graph,	8	A. I would expect that. My only point was
9	assuming you can extrapolate from the graph.	9	that Lindahl does not actually show that. But I
10	A. Well, Lindahl does not show that.	10	
11	Q. I know.	11	
12	A. If you extrapolate it and it remains	12	0
13	linear, then you can multiply that by another	13	
14	thousand.	14	
15	Q. So it would be a million fold if you did	15	
16	that extrapolation?	16	-
17	A. If it's linear over that range, then	17	-
18	would that be correct.	18	-
19	Q. And Lindahl shows that in pHs between 1.4	19	-
20	and 0.4 would be expected to lead to DNA	20	
21	depurination, correct?	21	
22	A. Lindahl does not I would have to look	22	
23	at this carefully, but I do not think Lindahl	23	
24	reports stabilities experiments under pH of 4.5.	24	
25	Q. Right.	25	
			, ,
1	Page 199	1	Page 200 MR. ZIMMERMAN: Objection. Asked and
1	suggesting, nor would anyone suggest that pH conditions of 1 pH 1 would be stable would be		answered.
3	useful for a DNA would be considered mild for a	3	THE WITNESS: I hope that I clarified
4	DNA reaction. As I tried to explain, this Boss		what I was relying on it for.
5	reference added perchloric acid, and I was simply	5	BY MR. SCHWARTZ:
6	including this reference to illustrate that water	6	Q. You used it in your declaration to say
-	did not poison the plate and catalyze reaction.	7	that allyl deprotection was a mild condition,
8	The reaction that's much closer to	8	
9			right?
9	demonstrating mild conditions for DNA is the Oian	a	right?
10	demonstrating mild conditions for DNA is the Qian	9	A. Allyl deprotections can be run under a
10	or Qian, Q-I-A-N, paper which is very which is	10	A. Allyl deprotections can be run under a wide variety of conditions.
11	or Qian, Q-I-A-N, paper which is very which is mild. So if you really want to talk about whether	10 11	 A. Allyl deprotections can be run under a wide variety of conditions. Q. But you were relying on that paper?
11 12	or Qian, Q-I-A-N, paper which is very which is mild. So if you really want to talk about whether or not palladium chloride mediated deprotections of	10 11 12	 A. Allyl deprotections can be run under a wide variety of conditions. Q. But you were relying on that paper? A. To make that statement, yes. Under a
11 12 13	or Qian, Q-I-A-N, paper which is very which is mild. So if you really want to talk about whether or not palladium chloride mediated deprotections of O-allyls are compatible with DNA, we should really	10 11 12 13	 A. Allyl deprotections can be run under a wide variety of conditions. Q. But you were relying on that paper? A. To make that statement, yes. Under a wide variety of conditions, some of which are
11 12 13 14	or Qian, Q-I-A-N, paper which is very which is mild. So if you really want to talk about whether or not palladium chloride mediated deprotections of O-allyls are compatible with DNA, we should really be talking we shouldn't be calculating pHs for,	10 11 12 13 14	 A. Allyl deprotections can be run under a wide variety of conditions. Q. But you were relying on that paper? A. To make that statement, yes. Under a wide variety of conditions, some of which are compatible with would be considered mild for
11 12 13 14 15	or Qian, Q-I-A-N, paper which is very which is mild. So if you really want to talk about whether or not palladium chloride mediated deprotections of O-allyls are compatible with DNA, we should really be talking we shouldn't be calculating pHs for, you know, this Boss paper. If I didn't make this	10 11 12 13 14 15	 A. Allyl deprotections can be run under a wide variety of conditions. Q. But you were relying on that paper? A. To make that statement, yes. Under a wide variety of conditions, some of which are compatible with would be considered mild for DNA, and some of which would not be.
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	yu Romesberg, Fli.D.		
1	Again, please look at Table 1. All of the allyl	1	Page 202 Q. Of a protecting group?
2	deprotection conditions in Boss are between three	2	A. You're not sequencing the protecting
3	hours and six days, correct?	3	group here.
4	A. That is correct.	4	Q. Do you want to go back to your testimony?
5	Q. And three hours to six days is not	5	We can go back to it.
6	consistent with SBS, is it?	6	A. Sure. Let's go back to it.
7	A. Three hours, as we went before, depending	7	Q. Okay. It was in IPR2013-00517. That was
8	on what requirements you insisted on, what	8	Exhibit 2007, and page 25.
9	efficiency meant to you, what applications you were	9	A. If I could take a minute to read this.
10	interested in. Maybe six days, very unlikely.	10	Q. Do you know where I'm directing you to?
11	Q. You don't remember your testimony when	11	A. Well, 25 is a pretty short very little
12	you said that three hours was not consistent with	12	text on 25, so I assume it's the text.
	SBS?	13	Q. It is where it says "Furthermore, the
14	A. In the context of what I was describing	14	amount of time required."
15	then, that is what I said, and that is what I	15	Do you see that?
16	meant. If you relax one's standards, three hours	16	A. Yes, I do.
17	would not be great. It would take a very long	17	Q. And you write: "Furthermore, the amount
18	time, but it is not per se inconsistent with	18	of time required to achieve those yields, three
19	running a sequencing by synthesis reaction.	19	hours to multiple days, would not have been
20	Q. But your position in those other IPRs was	20	expected to fulfill the rapid deprotection
21	that three hours was not compatible with SBS. That	21	requirement of Tsien's DNA sequencing method."
22	was your position there, correct?	22	You wrote that, right?
23	A. I believe I was stating something about	23	A. Yeah, and so just to be very clear with
24	whether or not it satisfied the requirements of	24	what I just said, and I was correct. I was talking
25	efficient high-throughput sequencing.	25	about fulfilling the rapid requirements or the
	encient high throughput sequenoing.	25	about running the rapid requirements of the
	Page 203		Page 204
1		1	
1	high-throughput, the fast sequence requirements of	1	of paper where you have your handwritten
2	high-throughput, the fast sequence requirements of Tsien.	2	of paper where you have your handwritten calculations and notes on.
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1	Page 205 Q. Why don't we turn to the claims.	1	Page 206 allyl as small.
2	A. Do you want to point me to a column.	2	Q. What is your understanding of the
3	Q. I'm just going to start with this. It is	3	chemically cleavable chemical group?
4	column 34 of the claims.	4	A. That it can be removed by a reaction that
5	Do you see after the structure it says	5	allows it to by a chemical reaction to produce a
6	wherein RA represents a small chemically cleavable	6	free three-prong hydroxyl group that is compatible
7	chemical group capping the oxygen at the 3'	7	with continued DNA synthesis.
8	position of the deoxyribose of the	8	Q. Do you see in the structure that is drawn
9	deoxyribonucleotide analog."	9	for this particular adenine nucleotide that the tag
10	Do you see that?	10	is attached to the base via a structure called Y?
11	A. Ido.	11	A. A box called Y, yes, I see that.
12	Q. What is your understanding of what is a	12	-
13	small group?	13	
14	A. Small. A small group. I am okay. I	14	-
15	accept Dr. Ju's definition, or his use of "small."	15	cleavable label a link or part of a label that
16	He doesn't define it. He represents it with	16	would attach what he calls the tag.
17	examples, which he uses MOM and allyl. Those are	17	5
18	both three carbons, so I am okay with that	18	
19	definition. I am okay with accepting his	19	•
20	definition of small.	20	talking about why the tag and why start and stop,
21	Do you want me to talk about small	21	because they're all going to be chemical bonds, but
22	outside of this? I'm happy to do so.	22	clearly the spirit here is that there is a part
23	Q. So you understood from reading this what	23	
24		24	·
25	A. I understand that Dr. Ju evokes MOM and	25	
		-	······································
1	Page 207 small in the other patents which I can put in	1	Page 208 Allyl ethers are cleaved by treatment
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2	small in the other patents which I can put in front of you if you want, but I'll represent it uses the same terminology would be the same as	2 3	Allyl ethers are cleaved by treatment with mercury II in acetone water. And he points to Gigg and Warren in 1968.
23	small in the other patents which I can put in front of you if you want, but I'll represent it uses the same terminology would be the same as you just described, your understanding of small? I	2 3 4	Allyl ethers are cleaved by treatment with mercury II in acetone water. And he points to
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110	yd Romesberg, Ph.D.	University in the City of New York			
	Page 209	4	Page 210		
1	both unsubstituted allyls as well as substitute	1	his reaction. Just like people would describe		
2	allyls, correct? A. No.	2	templates. That's a very common descriptor. It's		
3		3 not that they were all different, it's just you're			
4	Q. It's not a genus that includes those?	4	sequencing templates. Because there are many of		
5	A. You can stretch it to that if you want,	5	them. In a reaction, there is many molecules. So		
6	but it's certainly not common language.	6	if that caused confusion, I apologize. I never		
7	Q. But it is a collection of groups?	7	meant to implicate that there were different allyl		
8	A. You could stretch it to include that.	8	groups. If people wanted, like I said, to stretch		
9	It's kind of a slang, it's kind of a vernacular. A	9	the definition and to sort of use it as a slang,		
10	lot of chemical names have very long complex names,	10	that's certainly not unheard of. But an allyl		
11	and a lot of times people don't want to write them	11	group is something, and it's a very specific thing.		
12	out. A lot of times people don't even know how to	12 13	And it would be what anyone would think of.		
13	actually name things. So it's very common for		Q. But you didn't use the term "allyl		
14	people to use sort of a slang. So that's an allyl	14	group," you used the term "allyl groups." That is		
15	group. That's an allyl-type group. That's got an	15	what you wrote, correct?		
16	allyl in it, so let's call it an allyl protecting	16	A. Yeah, just methyl groups. That's a very		
17 18	group. It's kind of like that's a red car.Q. You yourself use the term 3'-O-allyl	17 18	common expression. If you want to modify methyl groups, you can use this sort of thing. And		
19	groups when you refer to Tsien, right?	19	there's no derivatives of a methyl group. It's		
20	A. That's right. He wasn't doing single	20	sort of like it the plural is referring to		
21 22	molecule experiments.	21 22	there are many of them. You're deprotecting		
	Q. And you also refer to Tsien saying allyl	22	you're not doing a single molecule reaction.		
23	ethers, correct?		You're not modifying one. You're removing a lot of		
24	A. Again, not single molecule experiments.	24	allyl groups, one for every molecule in there.		
25	What I meant was that there are many molecules in	25	Q. You know how to use the word group if you		
4	Page 211	1	Page 212		
1	wanted to; you used the word groups. You know how	1	molecule, but still plural. That's all I meant by		
2	wanted to; you used the word groups. You know how to use the word ether if you wanted to; you used	2	molecule, but still plural. That's all I meant by this.		
2 3	wanted to; you used the word groups. You know how to use the word ether if you wanted to; you used the word in plural, correct? You chose those	2 3	molecule, but still plural. That's all I meant by this.Q. You had meant that Tsien cites to Gigg		
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	yu Romesberg, Ph.D.		Oniversity in the City of New Fork
1	Page 213 Warren.	1	Page 214 BY MR. SCHWARTZ:
2	MR. SCHWARTZ: You look like you don't	2	Q. So none of those are three carbon allyl
3	agree	3	groups, correct?
4	MR. ZIMMERMAN: Not at all. Did you say	4	A. That is correct.
5	four hours and 10 minutes. We are almost at five	5	Q. And throughout Gigg and Warren there are
6	hours.	6	disclosures of allyl ethers that are both
7	THE VIDEOGRAPHER: I have to do the math.	7	unsubstituted and substituted, correct?
8	MR. ZIMMERMAN: We are 4:49 right now.	8	A. Apparently, yes.
9	BY MR. SCHWARTZ:	9	Q. And throughout Gigg and Warren there are
10	Q. All right. So we that is the same	10	also disclosures of allyl ethers other than a three
11	Gigg and Warren that is cited in Tsien, correct?	11	carbon allyl ether, correct?
12	A. I believe so, yes.	12	A. That is correct. Obviously, from what
13	Q. And please turn to page 1906 of Gigg and	13	you just pointed out. At least.
14	Warren, and on the left side towards the bottom of	14	
15	the page are shown compounds XL, IX, L, and L2.	15	5
16	Do you see that?	16	support of an allyl the allyl groups
17	A. I do.	17	understood that Tsien's reference to allyl ether is
18	Q. And none of those are three carbon allyl	18	
19	ethers, correct?	19	
20	A. I think whatever that is, LI is.	20	A. No. People as I said, people
21	Q. I didn't ask about LI. I said XL, IX, L,	21	sometimes use the word allyl, and describing this
22	and L2.	22	as an allyl ether, while technically still a little
23	A. Right. That is correct.	23	bit weird, is a little less weird than their
24	MR. ZIMMERMAN: He came up with 4:42.	24	generalizing a group of things. But still, the
	///	25	only thing in here that in here strictly properly
1	Page 215 referred to as an allyl group is the single allyl,	1	Page 216 better than that?
	the three carbon allyl. Other substituted ones	2	A. As I already said, yes, that is
3	technically should have different names, like there		
	···· , ···, ····	3	correct
4	should be like, for example, in you're going to	3 4	correct Q. Let's go back to Exhibit 1001, which was
5	should be like, for example, in you're going to test my nomenclature. The compound, the first one,	4	Q. Let's go back to Exhibit 1001, which was
_	test my nomenclature. The compound, the first one,	-	Q. Let's go back to Exhibit 1001, which was the patent the '852 patent.
5 6	test my nomenclature. The compound, the first one, which really have be called II methyl allyl, but	4 5 6	Q. Let's go back to Exhibit 1001, which was the patent the '852 patent.A. I'm sorry, where?
5 6 7	test my nomenclature. The compound, the first one, which really have be called II methyl allyl, but like I said, people are a little bit loose with	4 5	 Q. Let's go back to Exhibit 1001, which was the patent the '852 patent. A. I'm sorry, where? Q. The patent. The '852 patent that is
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5 6 7 8	test my nomenclature. The compound, the first one, which really have be called II methyl allyl, but like I said, people are a little bit loose with those because people don't like to write out things	4 5 6 7 8	 Q. Let's go back to Exhibit 1001, which was the patent the '852 patent. A. I'm sorry, where? Q. The patent. The '852 patent that is Exhibit 1001. Please go to column six, line 23. And in column 23 the patent reads "The MOM."
5 6 7 8 9	test my nomenclature. The compound, the first one, which really have be called II methyl allyl, but like I said, people are a little bit loose with those because people don't like to write out things like that.	4 5 6 7 8 9	 Q. Let's go back to Exhibit 1001, which was the patent the '852 patent. A. I'm sorry, where? Q. The patent. The '852 patent that is Exhibit 1001. Please go to column six, line 23.
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	yd Romesberg, Ph.D.	University in the City of New York			
1	Page 217 to figure 13.	1	Page 218 3:48 p.m.		
2	So the patent specification, Dr. Ju's	2	(The videotaped deposition of		
3	patent specification, specifically calls the CH2CH	3	FLOYD ROMESBERG, PH.D., VOLUME I		
4	double bond CH2 group the allyl group, correct?	4	concluded at 3:48 p.m.)		
5	A. As most people would, yeah. It's the	5	concluded at 5.40 p.m.)		
6	allyl group, and he parenthetically draws it out.	6			
7	Q. And that structure is not disclosed in	7			
8	Tsien, correct?	8			
9	A. It effectively is. Its name is disclosed	9			
10	in Tsien. It's not drawn out in a parenthetical	10			
11	manner like this, but in a way that any POSA would	11			
12	have immediately drawn that conclusion.	12			
13	Q. I'm just saying that the structure is not	13			
14	shown in Tsien, is it?	14			
15	A. The structure is not drawn out.	15			
1					
16	Q. Okay.	16 17			
18	MR. SCHWARTZ: I would like to go off the record if I could. I have to mark up some	17			
1	•				
19 20	exhibits. THE VIDEOGRAPHER: Off the record at	19 20			
21	3:38 p.m.	21			
23	(Recess taken.) THE VIDEOGRAPHER: On the record at	22			
		23			
24	3:48 p.m. This concludes today's deposition of	24			
25	Floyd Romesberg Ph.D. We are off the record at	25			
1	Page 219 I, the undersigned, a Certified Shorthand	1	Page 220 DECLARATION UNDER PENALTY OF PERJURY		
2	Reporter of the State of California, do hereby	2	Case Name: Illumina, Inc. vs. The Trustees of Columbia		
3	certify:		University in the City of New York		
4	That the foregoing proceedings were taken	3	Date of Deposition: 09/19/2018		
5	before me at the time and place herein set forth;	4	Job No.: 10047002		
6	that any witnesses in the foregoing proceedings,	5			
7	prior to testifying, were duly sworn; that a	6	I, FLOYD ROMESBERG, PH.D. hereby certify		
8	verbatim record of the proceedings was made by me	7	that I have read the transcript of my deposition and that		
9	using machine shorthand which was thereafter	8	this transcript accurately states the testimony given by me,		
10	transcribed under my direction; that the foregoing	9	with the changes or corrections, if any, as noted on an errata		
11	transcript is a true record of the testimony given.	10	sheet or errata sheets.		
12	Further, that if the foregoing pertains to the	11	Executed this day of		
13	original transcript of a deposition in a Federal	12	, 2018, at		
14	Case, before completion of the proceedings, review	13			
15	of the transcript [X] was [] was not requested.	14			
16	I further certify I am neither financially	15	FLOYD ROMESBERG, PH.D.		
17	interested in the action nor a relative or employee	16	NOTARIZATION (If Required)		
18	of any attorney of party to this action.	17	State of		
19	IN WITNESS WHEREOF, I have this date	18	County of		
20	subscribed my name.	19	Subscribed and sworn to (or affirmed) before me on		
21	Dated: September 21, 2018	20	this day of, 20,		
22	PUR, Q. Q.	21	by, proved to me on the		
		22	basis of satisfactory evidence to be the person		
23			who approved hefere we		
	PATRICIA Y. SCHULER	23	who appeared before me.		
23 24 25	PATRICIA Y. SCHULER CSR NO. 11949	23 24 25	who appeared before me. Signature: (Seal)		

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Job No.: 10047002 6 Page Line Reason 4 Reason Codes: 1. To clarify the record. 2. To conform to the facts. 8 Page Line Reason 5 3. To correct transcription errors. 9 From to 10 Page Line Reason 6 Page Line Reason 10 Page Line Reason 7 From to 11 From to 12 Page Line Reason 9 From to 13 From to 13 From to 11 From to 15 From to 16 Page Line Reason 17 From to 18 Page Line Reason 19 From to 12 Page Line Reason 10 Page Line Reason 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10		Name of Witness: Floyd Romesberg, Ph.D.	4	Page _	Line	Reason	
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8 Page Line Reason 11 From 12 Page Line Reason 12 Page Line Reason 13 From to 13 From to 14 Page Line Reason 14 Page Line Reason 13 From to 14 Page Line Reason 13 From to 14 Page Line Reason 13 From to 14 Page Line Reason 14 Page Line Reason 13 From 14 Page Line Reason 14 Page Line Reason 15 From 16 Page Line Reason 16 Page Line Reason 17 From 10 16 Page Line Reason 19 From 10	6		10	Page _	Line	Reason	
9 From	7		11	From _		t	to
9 From			12	Page _	Line	Reason	
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12 Page Line Reason 16 Page Line Reason 17 From to 17 From to 18 Page Line Reason 18 Page Line Reason 19 From to 19 From to 10 Page Line Reason 11 Page Line Reason 11 Page Line Reason 11 Page Line Reason 11 Page Line Reason 12 Page Line Reason 12 From 12 Subject to the above changes, I certify that the transcript is true and correct 12 12 Page Line Reason 12 12 No changes have been made. I certify that the transcript is true and correct. 12 12 Page <			15				
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20 Page Line Reason transcript is true and correct 21 From to 23 No changes have been made. I certify that the transcript is true and correct. 22 Page Line Reason 24 23 From to 24			22		_ Subject to	the above of	changes, I certify that the
21 From					transcript	is true and	d correct
22 Page Line Reason transcript is true and correct. 23 From			23		_ No changes	have been r	made. I certify that the
23 From to 24 24 Page Line Reason 24					transcript	is true an	nd correct.
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	25		25			FLOYD RO	OMESBERG, PH.D.

Floya Rolliesberg, Fli.D.			y in the City of New York
·	PREVIOUSLY MARK	00385 16:6,25	1012 12:9,10,13 13:5
Exhibits	ED EXHIBIT 2035	00517 58:23	33:9 80:16,20 208:6,
	4:15 21:17,21 27:1 49:21 58:22 72:2	00678 17:16	9
	EXHIBIT 2053 5:8	012 166:22,23	1013 15:24 16:2,5,6 17:21 19:11 43:19
PREVIOUSLY MARK ED EXHIBIT 1001 6:6	108:1	06678 17:15	47:10 57:16 83:4
204:15,19 216:4,8	EXHIBIT 2057 5:23	083 187:16,17,25	100:22 207:17
PREVIOUSLY MARK	190:22	188:11,16,18	208:15
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10			10				
11	VIDEOTAPED DEPOSITION OF FLOYD	ROMESBERG, PH.D.	11	Videotaped dep	osition of Floyd Ro	mesberg, Ph.D.,	
12 13	VOLUME II SAN DIEGO, CALIFORI	ΝΤΔ	12	Volume II, tal	en on behalf of the	Patent Owner at	5
14	SAN DIEGO, CALIFOR SEPTEMBER 20, 201		13	12790 El Camin	no Real, San Diego,	California,	
15		10	14	beginning at	9:04 a.m. and ending	at 10:41 a.m.,	on
16			15	-	2018, before PATRIC		
17			16	Certified Sho	thand Reporter No.	11949.	
18			17 18				
19	Reported By:		18				
20	PATRICIA Y. SCHULER CSR No. 11949		20				
20 21	Job No. 10047003		21				
22	505 NO. 1004/005		22				
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1 2	APPEARANCES: FOR PETITIONER:		1 2	WITNESS:	I-N-D-E-X	EXAMIN	NOTTON
3	KNOBBE MARTENS - SAN DIEG	~	3		RG, PH.D., VOLUME I		
4	BY: NATHANAEL R. LUMAN, H		4		BY MR. SCHWARTZ		228
5	12790 El Camino Real	-9 2 .	5		BY MR. ZIMMERMAN		270
6	San Diego California 9213		6				
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/	-	30	7		E-X-H-I-B-I-T-	5	
	(858) 707-4000	30	8	PATENT OWNER'	S		PAGE
8	(858) 707-4000 nate.luman@knobbe.com	30			S Document entitled	"A Mild and	PAGE 232
9	(858) 707-4000 nate.luman@knobbe.com - and -		8 9		S Document entitled Rapid Regeneration	"A Mild and of Alcohols	
9 10	(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO	DN DC	8		S Document entitled Rapid Regeneration from their Allylic	"A Mild and of Alcohols Ethers by	
9 10 11	(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN	DN DC , ESQ.	8 9		S Document entitled Rapid Regeneration	"A Mild and of Alcohols Ethers by ane/Sodium	
9 10 11 12	(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue,	DN DC , ESQ.	8 9 10	Exhibit 1037	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsia	"A Mild and of Alcohols Ethers by ane/Sodium t al"	232
9 10 11 12 13	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006</pre>	DN DC , ESQ. Suite 900	8 9 10 11	Exhibit 1037	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsia Iodide" by Kamal et	"A Mild and of Alcohols Ethers by ane/Sodium z al" "DNA polymerase	232
9 10 11 12 13 14	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com</pre>	DN DC , ESQ. Suite 900	8 9 10 11 12 13	Exhibit 1037 Exhibit 2031	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsil Iodide" by Kamal en Document entitled fluorescent substra reversible 3' tags	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with	232
9 10 11 12 13 14 15	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com FOR PATENT OWNER:</pre>	DN DC , ESQ. Suite 900	8 9 10 11 12	Exhibit 1037 Exhibit 2031	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsil Iodide" by Kamal e Document entitled fluorescent substra reversible 3' tags Document entitled	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with	232
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9 10 11 12 13 14 15 16 17 18	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com FOR PATENT OWNER: FITZPATRICK, CELLA, HARPEI BY: ROBERT S. SCHWARTZ, H BY: ZACHARY L. GARRETT, H</pre>	DN DC , ESQ. Suite 900 R & SCINTO ESQ. ESQ.	8 9 10 11 12 13 14 15	Exhibit 1037 Exhibit 2031	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsila Iodide" by Kamal en Document entitled fluorescent substra reversible 3' tags Document entitled "Palladium(II)-Cata Oxidation of Aldehy Ketones. 1. Carbony	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with " alyzed ydes and ylation of n Monoxide dium(II)	232
9 10 11 12 13 14 15 16 17 18 19	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com FOR PATENT OWNER: FITZPATRICK, CELLA, HARPEN BY: ROBERT S. SCHWARTZ, M BY: ZACHARY L. GARRETT, M 1290 Avenue of the America</pre>	DN DC , ESQ. Suite 900 R & SCINTO ESQ. ESQ.	8 9 10 11 12 13 14 15 16	Exhibit 1037 Exhibit 2031 Exhibit 2089	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsila Iodide" by Kamal en Document entitled fluorescent substra reversible 3' tags Document entitled "Palladium(II)-Cata Oxidation of Aldehy Ketones. 1. Carbony Ketones with Carbon Catalyzed by Pallad	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with " alyzed ydes and ylation of n Monoxide dium(II) ol"	232 235 238
9 10 11 12 13 14 15 16 17 18 19 20	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com FOR PATENT OWNER: FITZPATRICK, CELLA, HARPEN BY: ROBERT S. SCHWARTZ, H BY: ZACHARY L. GARRETT, H 1290 Avenue of the America New York, New York 10104</pre>	DN DC , ESQ. Suite 900 R & SCINTO ESQ. ESQ.	8 9 10 11 12 13 14 15 16 17	Exhibit 1037 Exhibit 2031 Exhibit 2089	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsili Iodide" by Kamal en Document entitled fluorescent substra reversible 3' tags Document entitled "Palladium(II)-Cata Oxidation of Aldehy Ketones. 1. Carbony Ketones with Carbon Catalyzed by Pallad Chloride in Methan	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with " alyzed ydes and ylation of n Monoxide dium(II) ol"	232 235 238
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9 10 11 12 13 14 15 16 17 18 19 20 21 22	<pre>(858) 707-4000 nate.luman@knobbe.com - and - KNOBBE MARTENS - WASHINGTO BY: WILLIAM R. ZIMMERMAN, 1717 Pennsylvania Avenue, Washington, D.C. 20006 bill.zimmerman@knobbe.com FOR PATENT OWNER: FITZPATRICK, CELLA, HARPEN BY: ROBERT S. SCHWARTZ, M BY: ZACHARY L. GARRETT, M 1290 Avenue of the America New York, New York 10104- zgarrett@fchs.com Also Present: Dr. Steven Menchen John White, Esq.</pre>	DN DC , ESQ. Suite 900 R & SCINTO ESQ. ESQ. as -3800	8 9 10 11 12 13 14 15 16 17 18 19 20 21	Exhibit 1037 Exhibit 2031 Exhibit 2089	S Document entitled Rapid Regeneration from their Allylic Chlorotrimethylsili Iodide" by Kamal en Document entitled fluorescent substra reversible 3' tags Document entitled "Palladium(II)-Cata Oxidation of Aldehy Ketones. 1. Carbony Ketones with Carbon Catalyzed by Pallad Chloride in Methana	"A Mild and of Alcohols Ethers by ane/Sodium c al" "DNA polymerase ates with " alyzed ydes and ylation of n Monoxide dium(II) ol"	232 235 238
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	yd Romesberg, Ph.D.		
1	Page 227 SAN DIEGO, CALIFORNIA; THURSDAY, SEPTEMBER 20, 2018	1	Page 228 representing Columbia. And at the end of the table
2	9:04 a.m.	2	is John Murnane, also representing Columbia from
3	THE VIDEOGRAPHER: Good morning. We are	3	the Fitzpatrick firm.
	now on the record. Today's date is Thursday,	4	MR. ZIMMERMAN: Bill Zimmerman of Knobbe
	September 20, 2018, and the time is 9:04 a.m.	5	Martens Olson & Bear representing Illumina and the
6	This is the video deposition of	6	witness, Dr. Romesberg. With me is my partner Nate
	Floyd Romesberg, Ph.D., Day II, being taken in the	7	Luman and Marcus Burch of Illumina.
	matter of Illumina, Inc. versus the Trustees of	8	THE VIDEOGRAPHER: The court reporter
	Columbia University in the City of New York,	9	today is Patricia Schuler, and she may now swear in
10	pending in the United States Patent and Trademark	10	the witness.
11	Office before the Patent and Appeal Board, Case	11	
12	Nos. IPR2018-00291, IPR2018-00318, IPR2018-00322,	12	FLOYD ROMESBERG, PH.D.,
13	and IPR2018-00385.	13	having been administered an oath, was examined and
14	We are at Knobbe Martens in San Diego,	14	testified as follows:
	12790 El Camino Real. My name is Daniel Bermudez	15	
16	of Aptus Court Reporting located at 600 West	16	FURTHER EXAMINATION
17	Broadway, Suite 300, San Diego, California 92101.	17	BY MR. SCHWARTZ:
18	Will counsel please identify yourselves	18	Q. Good morning, Dr. Romesberg.
19	and state whom you represent.	19	A. Good morning.
20	MR. SCHWARTZ: I'm Robert S. Schwartz	20	Q. Is there any reason you can't give
21	representing Columbia, from Fitzpatrick, Cella,	20	truthful testimony today?
	Harper & Scinto. With me is Zack Garrett, also	22	A. No.
23	from Fitzpatrick. Sitting next to him is	22	Q. Did you consult with your attorneys
24	Dr. Steven Menchen. Sitting next to him is	23	regarding the substance of your testimony
	John Light from the Cooper and Dunham firm also	24	yesterday?
20	Sour Light norm the obsper and Danham min also	23	yesterday :
1	Page 229	1	Page 230
1	A. No, I did not.	1	molecules may occur has been derived from
2	A. No, I did not.Q. And did you consult with your attorneys	2	molecules may occur has been derived from experiments on the reaction of mercury chloride
2 3	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate 	2 3	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA."
2 3 4	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? 	2 3 4	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that?
2 3 4 5	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. 	2 3 4 5	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do.
2 3 4 5 6	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to 	2 3 4 5 6	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant,
2 3 4 5 6 7	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was 	2 3 4 5 6 7	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct?
2 3 4 5 6 7 8	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was Exhibit 1013. Please draw your attention to 	2 3 4 5 6 7 8	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct? A. It denatures DNA, yes.
2 3 4 5 6 7 8 9	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was Exhibit 1013. Please draw your attention to Page 24, and look at Line 29. 	2 3 4 5 6 7 8 9	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct? A. It denatures DNA, yes. Q. Going back to the allyl ethers in Tsien,
2 3 4 5 6 7 8 9 10	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was Exhibit 1013. Please draw your attention to Page 24, and look at Line 29. And Tsien writes "Allyl ethers are 	2 3 4 5 6 7 8 9 10	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct? A. It denatures DNA, yes. Q. Going back to the allyl ethers in Tsien, the cleavage method for the allyl ethers cited by
2 3 4 5 6 7 8 9 10 11	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was Exhibit 1013. Please draw your attention to Page 24, and look at Line 29. And Tsien writes "Allyl ethers are cleaved by treatment with mercury II in acetone 	2 3 4 5 6 7 8 9 10 11	molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct? A. It denatures DNA, yes. Q. Going back to the allyl ethers in Tsien, the cleavage method for the allyl ethers cited by Tsien is based on the Gigg and Warren reference,
2 3 4 5 6 7 8 9 10 11 12	 A. No, I did not. Q. And did you consult with your attorneys regarding the substance of what you anticipate discussing today? A. No, I did not. Q. I am going to start out by asking you to go back to the Tsien reference. That was Exhibit 1013. Please draw your attention to Page 24, and look at Line 29. And Tsien writes "Allyl ethers are cleaved by treatment with mercury II in acetone water." 	2 3 4 5 6 7 8 9 10 11 12	 molecules may occur has been derived from experiments on the reaction of mercury chloride with DNA." Do you see that Kitt writes that? A. Yes, I do. Q. And mercury II is a DNA denaturant, correct? A. It denatures DNA, yes. Q. Going back to the allyl ethers in Tsien, the cleavage method for the allyl ethers cited by Tsien is based on the Gigg and Warren reference, correct?
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	ya Romesberg, Ph.D.		University in the City of New York
1	Page 231 strong basic conditions, and then the vinyl ethers	1	Page 232 You saw that, right?
2	are cleaved by mercury II, correct?	2	A. Yes.
3	A. Yes. I am not reading the first step.	3	Q. And that denatures DNA, correct? And
4	If I could take a second to read this?	4	that cannot be used for SBS, correct?
5	Q. Sure.	5	A. Certainly, at high concentrations
6	A. That seems to be the case, yes.	6	potassium t-butoxide would denature DNA. I don't
7	Q. The structure of allyl ethers is	7	know exactly at what concentration that would
8	different from that of vinyl ethers, correct?	8	occur, but that is a base, and, you know, I think
9	A. Yes.	9	at sufficiently basic conditions, DNA is damaged.
10		10	Q. So a person reading Tsien along with Gigg
11		11	and Warren would not use Gigg's condition for
12		12	cleavage of allyl ethers for SBS because there
13		13	
14	•	14	
15		15	A. I think that is fair.
16		16	(Exhibit 1037 was previously marked for
17		17	
18		18	,
19	· ·	19	Q. I am going to put another exhibit in
20		20	front of you because I know you don't have enough
21		21	of them already. I am putting in front of you
22		22	Exhibit 1037, which is a paper entitled "A Mild and
23	-	23	Rapid Regeneration of Alcohols from their Allylic
24		24	
25	-	25	
1	Page 233 Have you seen this paper before.	1	Page 234 Q. And protecting groups for use in SBS must
1	Have you seen this paper before,	1	Q. And protecting groups for use in SBS must
2		1 2 3	Q. And protecting groups for use in SBS must be removable under aqueous conditions. We went
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FIO	/d Romesberg, Ph.D.		University in the City of New York
1	Page 235 the DNA. And you would dehydrate the DNA and cause	1	Page 236 reasoned that."
	it to denature.	2	Do you see that?
3	Q. And Kamal's allyl deprotection method	3	A. I do.
4	uses chlorotrimethylsilane, correct?	4	Q. Canard writes "We reasoned that ether
5	A. It does.	5	bonds would be hard to cleave under mild conditions
6	Q. And chlorotrimethylsilane is not	6	compatible with DNA chemical stability."
7	compatible with water, correct?	7	That's what Canard writes, correct?
8	A. No, it is not.	8	A. That is what is written.
9	(Exhibit 2031 was previously marked for	9	Q. And the Canard paper was published in
10	identification.)	10	1994, correct?
11	BY MR. SCHWARTZ:	11	A. That is correct.
12	Q. I am putting in front of you	12	Q. And the Tsien application was published
13	Exhibit 2031, which is a paper entitled "DNA	13	in 1991, correct?
14	Polymerase Fluorescent Substrates with Reversible	14	A. That is correct.
15	3'-Tags" by Canard and Sarfati, published in 1994.	15	Q. So even years after Tsien, it was
16	Please turn to page 4. You've seen this	16	understood that ether bonds are hard to cleave
17	paper before, haven't you, Dr. Romesberg?	17	under mild conditions compatible with DNA chemical
18	A. I have.	18	stability.
19	Q. Please draw your attention to the right	19	That is what Canard says, right?
20	side of the page, and please look at the bottom	20	A. Canard says that they reasoned that
21	where there is a section D.	21	that ether bonds would be hard to cleave.
22	Do you see that?	22	Q. But that reasoning was three years after
23	A. I do.	23	the publication of Tsien?
24	Q. Please draw your attention to the third	24	A. Yes. Canard, in this paper, suggested
25	sentence in that section that starts with "We	25	that they thought that ethers would be difficult to
1	Page 237 Cleave.	1	Page 238 identification.)
2	Q. And in your declarations that you	2	BY MR. SCHWARTZ:
3	submitted in the current IPRs, you don't cite to	3	Q. I'm putting in front of you a paper
4	any reference prior to October 2000 that provides	4	entitled "Palladium(II)-Catalyzed Oxidation of
5	data showing that a nucleotide with a 3'-O-allyl	5	Aldehydes and Ketones. 1. Carbonylation of Ketones
6	protecting group is cleaved under mild conditions	6	with Carbon Monoxide Catalyzed by Palladium(II)
7	compatible with DNA chemical stability, do you?	7	Chloride in Methanol" by Hamed et al, which was
8	A. I provided examples of reactions that	8	published in 2001.
9	were very close, that a POSA would have expected,	9	Please turn to page 182. And please draw
10	and had high expectation of success that the one	10	your attention to the left side of the page towards
11	difference of methanol versus water would not have	11	the top of the page where there is a reaction
12	been a problem, and that would have provided	12	scheme entitled "Alkoxy Palladium Root."
13	conditions where a 3'-O-allyl protected nucleoside	13	Do you see that?
14	could have been allyl deprotected. I did not find	14	A. I do.
15	examples of precisely that reaction in water.	15	Q. The reaction scheme showing
16	Q. And sitting here today, are you aware of	16	palladium (II)reacting with an olefin plus an
17	any examples of that precise reaction occurring in	17	alcohol to generate a palladium (II)complex with
18	water?	18	the olefin with the generation of acid."
19	A. No, I did not find any last night.	19	That is what is the reaction scheme
20	Q. Did you look for them last night?	20	shows, correct?
20	A. I did not.	20 21	A. It does show that.
22	MR. SCHWARTZ: I am going to go to	21 22	Q. An allyl protecting group is an example
22	another exhibit, please. This one will require a	22	of an olefin, correct?
23	new marking, Court Reporter, 2089.	23 24	A. Yes.
	new marking, court neporter, 2003.	47	A. 100.
24	(Exhibit 2089 was marked for	25	Q. So a POSA reading I'm sorry, strike

	ya Romesberg, Ph.D.		University in the City of New Fork
1	Page 239 that.	1	Page 240 Lindahl.
2	A POSA understood that reacting in allyl	2	Do you recall that?
3	protecting group with palladium (II) will generate	3	There were two papers, Lindahl 1993 and
4	acid, correct?	4	Lindahl and Nyberg 1972, and those papers reported
5	A. Yes.	5	that acid catalyzes depurination of nucleotides.
6	MR. ZIMMERMAN: Object to the form of the	6	Do you remember that?
7	question. And just so you know my objection, you	7	A. Yes.
8	made clear yesterday that a POSA was in 2002	8	Q. And conditions that result in
9	THE WITNESS: 2000.	9	depurination of nucleotides are not compatible with
10	MR. ZIMMERMAN: 2000. And you are asking	10	SBS, correct?
11	him about something dated after that and assuming a	11	A. If you're depurinating during sequencing
12	POSA would know.	12	reaction, that would not be compatible with
13	So the timing in your question is	13	sequencing by synthesis.
14	nonsensical. So I am objecting as to form.	14	Q. And please go back to the Qian reference
15	MR. SCHWARTZ: Okay.	15	we looked at yesterday, Exhibit 1036, and go back
16	BY MR. SCHWARTZ:	16	to Page 2185 where there was Scheme 2. And please
17	Q. A POSA understood that reacting in allyl	17	look again at the figure legend. The deprotection
18	protecting group, palladium (II) generates acid,	18	of the allyl takes place with palladium (II) and
19	correct?	19	methanol, correct?
20	A. Yes.	20	A. It does.
21	Q. So please	21	Q. And that reaction would be expected to
22	A. Stoichiometric acid, yeah.	22	produce acid, correct?
23	Q. Please go back to the first page of	23	A. It would produce stoichiometric amounts,
24	Hamed strike that.	24	5
25	Yesterday we looked at some papers by	25	compared to the adding solvent at millimolar
	Page 241		Page 242
1	concentrations of the sort that we talked about	1	
2	yesterday, and of the sort that Lindahl talked	2	to the abstract. And the abstract reads I'm not
3	about, this would generate the amounts of acid	3	going to read the whole thing. I am going to ask
4	that one proton for every time you deprotected	4	you to look at the second sentence that starts with
5	an allyl. And that's common, and that would be	5	"The results."
6	handled under sequencing by synthesis methods,	6	Do you see that?
7	because I assume that is what we are interested in.	7	"The results indicate that palladium (II)
8	And that would be handled by buffering the system.	8	interacts with both the phosphate and bases of
9	That would be well within the buffer power and	9	DNA."
10 11	standard for any POSA to deal with.	10 11	That is what the authors write, correct? A. That is correct.
12	Q. But I'm not talking about a different reaction condition. I'm talking specifically about	12	
13	Qian's reaction condition.	12	
14	So in Qian's reaction condition that would	14	•
15	be expected to produce acid, there is not a buffer	15	
16	there, correct?	16	
17	A. It would produce stoichiometric acid.	17	intro they're talking about palladium II, and the
18	MR. SCHWARTZ: I am going to mark a new	18	
19	exhibit. Court reporter, this will require a new	19	
20	number, 2094.	20	
21	(Exhibit 2094 was marked for	21	consistent with that.
22	identification.)	22	
23	BY MR. SCHWARTZ:	23	
24	Q. I'd like to direct you to a document	24	
25	-	25	

 chloride, figure 4." chloride, figure 4." hydrogen bonded double helical system has bean disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the regions where binding of disturbed at least in the authors write, correct? A. That is what the authors write, correct? A. That is what the authors write, correct? disturbed at least in the authors write, correct? disturbed a		yu Romesberg, Ph.D.		University in the City of New For
3 into the record, but let me just ask you to go to the record to the paragraph that starts with "The ultraviolet." 3 palladium (II) takes place." 4 next paragraph that starts with "The ultraviolet." WR. ZIMMERMAN: Object norm. 6 A. 1do. G. That is what it withen. 7 O. It reads "The ultraviolet spectrum 6 Q. That is what the authors write. correct? 6 A. 1do. C. A POSA understood from Pillal that 9 the polyme, thus affecting the shape and magnitude 9 10 electron effects." Cites a reference. 1 Witston Crick hydrogen bondd 13 of the cotton effects." Cites a reference. 12 and the fidelity of the Watson Crick hydrogen 14 "Unstacking of the bases of 12 and the fidelity of the Watson Crick hydrogen 14 polynucleotides may give rise to a ref serence. 13 of the cotton effects." And it cites a 15 polynucleotides may give rise to a ref serence. 14 A. There's no evidence that the duplex is 16 palpha heily (1) to NA results in reduction of ORD 20 Satistis frequction of NA 22 palladium (II) to bards accompanied by red shifts. These 14 If ou want the exoplan a lifte bit abou	1	Page 243 chloride, figure 4."	1	Page 244 hydrogen bonded double helical system has been
4 next paragraph that starts with "The ultraviolet." 4 MR. ZIMMERMAN: Object to norm. 5 Dy ou see that? 5 BY MR. SCHWARTZ: 7 Q. It reads "The ultraviolet spectrum 6 Q. That is what its written. 8 (figure 1), indicates binding of palladium (II) to 9 A. That is what its written. 9 the bases of DNA. This is expected to modify the 9 palladium chloride cannot be used in a reaction 10 electronic state of the bases, and will be 10 with DNA, because palladium chloride disturbs the 11 reference. "Similarly disruption of the bases of 11 Watson Cick hydrogen bonded double helical system. 13 of the coth effects." Cits a reference. 13 bonded double helical system is essential for DNA 14 regrence. "Similarly disruption of hydrogen bonds 14 sequencing including SBS. 14 polynucleotides may give rise to a red shift in the 20 exequencing by synthesis here. 15 polynucleotides may give rise to a red shift in the 20 Sequencing by synthesis here. 14 regreads and dB show that binding of 21 HG/NO wat me to explain alittle bindou 12 reladium (II) to DNA results	2	And I'm not going to read the whole thing	2	disturbed at least in the regions where binding of
5 Do you see that? 6 BY MR. SCHWARTZ: 6 A. 1 do. 6 Q. That is what is what is what or write, correct? 7 Q. It reads "The ultraviolet spectrum 8 (figure 1), indicates binding of palladium (il) to 3 Q. A POSA understood from Pillai that 8 (figure 1), indicates binding of palladium (il) to 3 A. That is what is witten. Q. A POSA understood from Pillai that 9 electronic state of the bases, and will be 10 with DNA, because palladium chloride disturbs the 11 reflected in the interaction of their dipoles in 11 with DNA, because palladium chloride disturbs the 12 the polymer, thus affecting the shape and magnitude 10 with DNA, because palladium chloride disturbs the 13 of the cotton effects." Cites a reference. 12 and the fidelity of the Watson Crick hydrogen 14 sequencing, by synthesis here. 20 Sequencing by synthesis here. 20 21 reference. 11 13 A. There's no evidence that the duplex is 15 polyucleotides is thought to result in drininshed 19 disrupted in any maner that would prevent 20 sequencing by synthesis	3	into the record, but let me just ask you to go to th	3	palladium (II) takes place."
 A. 1do. C. It reads "The ultraviolet spectrum G. It reads "The ultraviolet spectrum (figure 1), indicates binding of palladium (II) to the bases of DNA. This is expected to modify the electronic state of the bases, and will be electronic state of the bases, and will be electronic state of the bases, and will be if reference. "Similarly disruption of their dipoles in the cotton effects." Cites a reference. "Justacking of the bases of polynucleotides is thought to result in diminished magnitudes of cotton effects." And it cites a reference. "Similarly disruption of hydrogen bonds between the base pairs of double-stranded polynucleotides may give rise to a red shift in the position of the maxima." And it cites a reference. "Figure 4A and 4B show that binding of patient of the cotton effects." And it cites a reference. "Figure 4A and 4B show that binding of patient and the diston and Crick polynucleotides may give rise to a red shift in the position of the maxima." And it cites a reference. "Figure 4A and 4B show that binding of and CD bands accompanied by red shifts. These reduced considerably, and the Watson and Crick store the considerably, and the Watson and Crick People don't understand the origins of Soft words lystems. We say, okay, an and help are only way that we use these signals is by model system. Ma and a protein and beta show that, you can't say that its denatured the small changes happen all the time, the cost. It sub the way shat its denatured and they are only understandable in the cortext of a an an CRD signal. Suggesting that the suble is sub discusts. ORD and CD. Its - the only way that we use these signals is by model system. Ma an ordenis sub this. Show that	4	next paragraph that starts with "The ultraviolet."	4	MR. ZIMMERMAN: Object to norm.
7 A. That is what is written. 8 (figure 1), indicates binding of palladium (ii) to the bases of DNA. This is expected to modify the the bases of DNA. This is expected to modify the the bases of DNA. This is expected to modify the the bases of DNA. This is expected to modify the the bases of DNA. This is expected to modify the the bases of DNA. This is expected to modify the the bases of DNA. This is expected to the bases of the magnitudes of cotton effects." Cites a reference. 10 with DNA, because palladium chloride disturbs the the bases of the magnitudes of cotton effects." Cites a reference. 16 magnitudes of cotton effects." And it cites a magnitudes of cotton effects." And it cites a polynucleotides may give rise to a red shift in the polynucleotides may give rise to a red shift in the polynucleotides may give rise to a red shift. These 2 10 A. May don't you agree? 17 C. And why don't you agree? 11 14 You want me to explain a little bit abou 2 20 CRO, CRO stands for optical rotatory dispersion. CD 2 20 CRO, CRO stands for optical rotatory dispersion. CD 2 20 CRO polarize light. 2 20 CRO and right polarize light. 2 2	5		5	BY MR. SCHWARTZ:
8 (figure 1), indicates binding of pailadium (II) to 9 8 Q. A POSA understood from Pillai that 9 9 the bases of DNA. This is expected to modify the 10 10 10 11 reflected in the interaction of their dipoles in 3 10 10 With DNA, because palladium chloride disturbs the 10 12 the polymer, thus affecting the shape and magnitude 3 of the cotton offects." Cites a reference. 13 10 with DNA, because palladium chloride disturbs the sequencing, including SBS. 15 polynucleotides is thought to result in diminished 16 16 No. 17 reference. "Similarly disruption of hydrogen bonded 17 16 A. No. 18 between the base pairs of double-stranded 19 17 Nerefreence. 14 16 19 polynucleotides may give rise to a red shift in the 20 20 And My don't you agree? 1 14 regues duiges that stacking interactions are 24 11 11 14 14 20 palladium (II) to DNA results in reduction of ORD 24 20 11 11 14 24 results sugges that stacking interacting interacting 24 results suggesting stacking interacting and 25	6	A. Ido.	6	Q. That is what the authors write, correct?
8 (figure 1), indicates binding of pailadium (II) to 9 8 Q. A POSA understood from Pillai that 9 9 the bases of DNA. This is expected to modify the 10 10 10 11 reflected in the interaction of their dipoles in 3 10 10 With DNA, because palladium chloride disturbs the 10 12 the polymer, thus affecting the shape and magnitude 3 of the cotton offects." Cites a reference. 13 10 with DNA, because palladium chloride disturbs the sequencing, including SBS. 15 polynucleotides is thought to result in diminished 16 16 No. 17 reference. "Similarly disruption of hydrogen bonded 17 16 A. No. 18 between the base pairs of double-stranded 19 17 Nerefreence. 14 16 19 polynucleotides may give rise to a red shift in the 20 20 And My don't you agree? 1 14 regues duiges that stacking interactions are 24 11 11 14 14 20 palladium (II) to DNA results in reduction of ORD 24 20 11 11 14 24 results sugges that stacking interacting interacting 24 results suggesting stacking interacting and 25	7	Q. It reads "The ultraviolet spectrum	7	
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10electronic state of the bases, and will be10with DNA, because palladium chloride disturbs the11reflected in the interaction of their dipoles in11Watson Crick hydrogen bonded double helical system,13of the cotton effects." Cites a reference.12and the fdelity of the Watson Crick hydrogen14"Unstacking of the bases of13bonded double helical system is essential for DNA15polynucleotides is thought to result in diminished15Do you agree?16magnitudes of cotton effects." And it cites a16A. No.17reference. "Similarly disruption of hydrogen bonded16A. No.18between the base pairs of double-stranded16A. Muhy don't you agree?19polynucleotides may give rise to a red shift in the19disrupted in any manner that would prevent20and CD bands accompanied by red shifts. These23stands for cipcial rotatory dispersion. CD23and CD bands accompanied by red shifts. These24chiral molecules, when you polarize light,24results suggest that stacking interactions are24chiral molecules, when you polarize light,25reduced considerably, and the Watson and Crick1thing, and sodium is perfectly compatible.2vou get differences in the ORD and CD signals.5some respects it's - there's no evidence here that3and CD. Lit's - the only way that we use these3introgens of the base, it still has a CD an3and they are only understand the origins ofCRD and the ORD that s	9		9	palladium chloride cannot be used in a reaction
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	ya Romesberg, Ph.D.		University in the City of New York
1	Page 247 say, correct?	1	Page 248 MR. ZIMMERMAN: You said palladium not
2	A. I don't see that they relate that to	2	palladium II.
3	recognition by a polymerase in any manner.	3	MR. SCHWARTZ: Okay. I will correct it
4	Q. They are simply saying that the presence	4	by saying palladium II.
5	of palladium results in stacking interactions	5	THE WITNESS: That is what they say. And
6	reduced considerably they use the word	6	I think a POSA would have understood that those
7	"considerably" and that the Watson Crick	7	changes that number one, that there was no link
8	hydrogen bonded double helical system is disturbed	8	to any sequencing by synthesis to any polymerase
9	where the regions of palladium binding are. That	9	recognition for that matter at all. And a POSA
10	is what they say. That is what they teach, right?	10	certainly would have still understood that there
11	A. Could you, again, point me to exactly	11	was still a helical structure there.
12	where they say that?	12	
13	Q. At the end of the paragraph: "These	13	whatever extent they're reduced considerably, they
14	results suggest that stacking interactions are	14	
	reduced considerably." Those are their words.	15	0
15	-		Q. But a POSA reading this would understand
	"And the Watson and Crick hydrogen bonded double	16	
17	helical system has been disturbed, at least in the	17	that the palladium causes stacking interactions to
18	regions where binding of palladium takes place."	18	be reduced considerably?
19	That was the authors' interpretation of	19	MR. ZIMMERMAN: Objection. Asked and
20	their own data, correct?	20	answered.
21	MR. ZIMMERMAN: Objection.	21	THE WITNESS: Those are the words there.
22	Mischaracterizes the document. And you read it	22	BY MR. SCHWARTZ:
23	incorrectly.	23	Q. What you were talking about before when
24	MR. SCHWARTZ: Where did I read it	24	you gave your long explanation of the ORD and the
25	incorrectly?	25	CD, that was speculation on your part on whether or
	Page 249		
1		1	
	not these were reduced considerably, correct?	1	And then there is a partial attorney
2	not these were reduced considerably, correct?A. I don't understand the question. What	2	And then there is a partial attorney statement, and the question continues with
2 3	not these were reduced considerably, correct?A. I don't understand the question. Whatpart are you asking me if it was speculation?	2 3	And then there is a partial attorney statement, and the question continues with 3'-oxygen protecting groups. The question that was
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1 10 y	u Romesberg, Fil.D.		Oniversity in the City of New Ton
1	Page 251 MR. ZIMMERMAN: Objection. Incomplete.	1	Page 252 Q. Take your time.
	BY MR. SCHWARTZ:	2	A. Okay. Could you repeat the question?
3	Q. I said the portions that I read were what	3	Q. Sure. So my question was you testified
	you testified, correct?	4	that as of 2002 people did not believe that they
5	A. Those are the words written, yes.	5	could modify or engineer polymerases to improve
6	Q. And you testified that as of 2002, people	6	their ability to incorporate nucleotides with 3'-O
-	did not believe that they could modify or engineer	7	protecting groups?
	polymerases to improve their ability to incorporate	8	A. I think that's too general of a
	nucleotides with 3'-O protecting groups, correct?	9	statement. That is certainly not what I intended
10	MR. ZIMMERMAN: Objection.	10	to say in this declaration. What I intended to say
11	Mischaracterizes.	11	was that it would have been viewed by in
12	THE WITNESS: Could you repeat the	12	general, as being very difficult.
13	question?	13	Q. But you didn't actually say that. You
14	BY MR. SCHWARTZ:	14	said that people would not believe that you could?
15	Q. You testified that as of 2002 people did	15	A. No, I did say that. If you read on, I
16	not believe that they could modify or engineer	16	said people would not have considered that a likely
17	polymerases to improve their ability to incorporate	17	possibility. A handful of groups including
18	nucleotides with 3'-O protecting groups?	18	people were starting to make progress. In 2002,
19	MR. ZIMMERMAN: Objection.	19	people were starting to make progress. A handful
20	Mischaracterizes.	20	of groups, including mine, sure. No one thought it
21	BY MR. SCHWARTZ:	21	was easy. To this day it is not easy.
22	Q. That is what you testified, correct?	22	So I certainly meant that it was difficult
23	A. I would like to read this.	23	and that people would have viewed it as difficult, n
24	Q. Sure.	24	doubt.
25	A. One second please.	25	Q. And in 2000, it would have been even more
	·		
1	Page 253 difficult, correct?	1	Page 254 "Answer: Yes.
2	A. There are no significant differences	2	"Question: And in the discussions you've
	between 2000 and 2002, in my opinion, in my	3	had with Dr. Tsien, have you ever discussed
	recollection. I don't know of anything that would	4	modified nucleotides?
	have changed my answer for 2000.	5	"Answer: Surprisingly, no."
6	Q. But again, in the testimony that we read,	6	I'm not going to read the whole
	your response to the question at that deposition	-	
	was, no, people did not believe that you could.	8	you can look at any part that you want. I'm going
9	A. No, I believe I tried to clarify that and	9	to ask that you skip ahead to Page 46 and draw your
10	explain that people would have thought it was very	10	attention to Line 8. Starting at Line 8, the
11	difficult. But there were groups working on it.	11	transcript reads:
12	So if there were groups working on it,	12	"Answer: And to tell you the truth, I
13	people some people believed that you could,	13	never really knew that he did this work until I
14	including my group. So I believed we could.	14	read this patent."
15	But what I was trying to say then is that -	15	And the portions of the transcript I read
16	and this certainly is still the case today, is that	16	is what you testified, correct?
17	it's not easy.	17	A. That is what is written, yes.
18	Q. I am going to ask you to turn back to	18	Q. And the patent which you were referring
19	Exhibit 2080, which is your deposition transcript	19	to in your testimony is the Tsien patent
20	in IPR2013-00128. Please turn to Page 45.	20	application, which is marked as Exhibit 1013,
21	A. I am there.	21	correct?
22	Q. Please draw your attention to Line 21,	22	A. It is the Tsien patent. It is marked as
23	and starting at Line 21 the transcript reads:	23	1013.
		24	Q. And you've been doing scientific research
24			
24 25	"Question: So then you've met Dr. Tsien?	25	on modified nucleotides for over 25 years; is that

Floy	/d Romesberg, Ph.D.		University in the City of New York
1	Page 255 right?	1	Page 256 A. No.
2	A. No. I've been doing work on modified	2	Q. And you testified in 2014 in the
3	nucleotides since 19 the end of 1998, so	3	transcript that we just looked at that you never
4	Q. About 20 years?	4	even knew that Tsien did any work on modified
5	A. About 20 years.	5	nucleotides until you became involved as an expert
6	Q. And you've published many research papers	6	for Illumina in the 2013 time frame, correct?
	on modified nucleotides during that time, right?	7	A. I guess I'm not positive I said that. I
8	A. I have published a number of papers on	8	certainly was unaware that he had worked on the
-	that topic, yes.	9	sequencing aspect of modified nucleotides. I mean,
10	Q. Did you ever cite to Tsien's patent	10	he did lots of work on RNA and DNA, and there were,
11	application in any of your papers?	11	for example, linkers in that. And I would have to
12	A. No.	12	rethink that carefully as to I've never cited
13	Q. Do you recall ever citing to Tsien's	13	it, but I am sort of peripherally aware that he did
14	patent application in any of your review articles?	14	that work, but I believe what would be a factual
15	A. No.	15	statement and I would have to read this to make
16	Q. Do you ever recall citing to Tsien's	16	sure that this is what I said, but the content of
17	patent application in any of your grant	17	the patent that is under discussion, I was unaware
18	applications?	18	that he had been interested in that.
19	A. No, I did not.	19	MR. SCHWARTZ: I'm just looking at the
20	Q. Do you ever recall citing to Tsien's	20	record.
21	patent application in any presentations that you've	21	THE WITNESS: While you do that, could I
22	made?	22	request a bathroom break?
23	A. No.	23	MR. SCHWARTZ: Sure.
24	Q. I'm talking about scientific	24	THE VIDEOGRAPHER: We're off the record
25	presentations.		at 9:46 a.m.
20		20	
1	Page 257 (Recess taken.)	1	O-modified nucleotides. Page 258
2	THE VIDEOGRAPHER: We are on the record	2	Q. So a person of ordinary skill who is
	at 9:49 a.m.	3	developing nucleotides for use in SBS in the 2000
-	BY MR. SCHWARTZ:	4	time frame, it's fair to say that they would give
5	Q. I asked you this yesterday,	5	more weight to Metzker's paper that does provide
-	Dr. Romesberg, and I think you agree that Tsien	-	data than Tsien's prophetic application that is
	doesn't provide any data showing that any 3' block	7	completely devoid of data, correct?
8	nucleotide works for DNA sequencing, correct?	8	A. On certain issues that data certainly
9	MR. ZIMMERMAN: Objection. Asked and	9	is important. And I think that there are people
10	answered by your own admission.	10	would have weighted that data differently and
11	THE WITNESS: Yes. I believe we had that	11	certainly more impactfully. They would have
12	discussion. And I believe I agreed with that.	12	seeing is believing.
13	BY MR. SCHWARTZ:	13	So I just don't want to say in any way that
14	Q. And Tsien's application is entirely	14	they would have then dismissed a good idea, whether
15	prophetic, right?	15	you viewed Tsien's idea as good or not.
16	A. On this subject, yes. I'd have to go	16	The downside of data is that it is a
17	back and I did not have that in mind as I read	17	specific question, and you can only get answers to
18	it, so it is possible that there were spots that	18	specific questions.
19	had data. I did look at the patent in regards to	19	So it is difficult to interpret data that i
20	this particular issue, and there is no data.	20	negative as being it is difficult to know how
21	Q. And in contrast to Tsien's prophetic	21	general a negative result is.
22	patent application, Metzker 1994 does provide data	22	Q. I think my question may have been a
	for a number of 3' block nucleotides for use in DNA	23	little bit different, which is, given that Metzker
23			
23 24		24	actually provided data for 3'-O protecting groups
-	sequencing, right? A. Metzker provides data for, I think, eight		actually provided data for 3'-O protecting groups for use in DNA sequencing, is it fair to say that a

Flo	yd Romesberg, Ph.D.		University in the City of New York
	Page 259	4	Page 260
1	skilled artisan would give more weight to a paper with data in terms of understanding how these 2' O	1	if you have a situation where you have a paper and suppose that you are interested in developing 3'-O
2	with data in terms of understanding how those 3'-O	2	
3	protecting groups would work for DNA sequencing	3	protecting groups for DNA sequencing, you have a
4	than another application that is entirely	4	paper that actually tests 3'-O protecting groups?
5	prophetic?	5	A. See, that is the problem. So you can
6	A. I mean, I understand what you are saying,	6	make the suggestion that you could use 3'-O
7	and in one perspective I agree with that because,	7	protecting groups, and you cannot address that
8	like I said, data speaks louder than suggestions.	8	question in a single paper. You cannot fully
9	Seeing is believing. But at the same time, there	9	address that question.
10	is always the caveat because, as I said,	10	5
11	experimental data always is a specific experiment.	11	one paper and you can look at the specific
12	It answers a specific question.	12	
13	And you may look at that and say, well, tha	13	
14	is not the question exactly I wanted, or I think tha	14	
15	I could optimize this. There are a variety of	15	•
16	responses.	16	5
17	In general, certainly seeing is	17	
18	believing, and you want to see data. That is	18	
19	always good.	19	
20	So, yeah, what you're saying is certainly	20	
21	sort of true, but at the same time, a good idea is a	21	you could use different polymerases or whatever, but
22	good idea. And sometimes people want to pursue it	22	51
23	more than someone else might.	23	
24	Q. I think again, I'm not sure that you	24	, , , , , , , , , , , , , , , , , , , ,
25	are responding directly to my question, which is,	25	But to the extent that data answers
	Page 261		Page 262
1	specific questions and only those specific	1	Q. The IPR2013-00517. It is Exhibit 2007.
2	specific questions and only those specific questions, then you might not want to dismiss an	2	Q. The IPR2013-00517. It is Exhibit 2007. A. Yes.
2 3	specific questions and only those specific questions, then you might not want to dismiss an entire idea based on one particular set of data.	2 3	 Q. The IPR2013-00517. It is Exhibit 2007. A. Yes. Q. Please turn to Page 23. Please look at
2 3 4	specific questions and only those specific questions, then you might not want to dismiss an entire idea based on one particular set of data. Q. Please go back to Exhibit 2035, your	2 3 4	 Q. The IPR2013-00517. It is Exhibit 2007. A. Yes. Q. Please turn to Page 23. Please look at the first full sentence. And the sentence reads
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loyd Romesberg, Ph.D.		University in the City of New Yor
Page 263 1 Mischaracterizes.	1	Page 26 Q. And so you wrote it?
2 THE WITNESS: That is not what I intended	2	A. It's written. Two hours with 80 percent
3 to write.	3	yield is very different than two hours
4 BY MR. SCHWARTZ:	4	quantitative.
5 Q. Did you write it?	5	So the time is inseparable from the yield.
6 MR. ZIMMERMAN: Objection.	6	Those are related things. So because I know whe
7 Mischaracterizes.	7	you are going with this, just be clear that my
8 THE WITNESS: I wrote the words that are	8	criteria for two hours, that is influenced by the
9 there. You are interpreting it differently than I	9	fact that it is 90 percent yield. So just to be
10 intended.	10	clear.
11 BY MR. SCHWARTZ:	11	Q. So if the yield is not satisfactory, two
12 Q. Do you not say that any one of these	12	hours is certainly not satisfactory with that
13 considerations would have led one of ordinary skill	13	yield. Is that what you are saying?
14 to have expected stannous chloride to be	14	A. I'm sorry. Could you repeat that?
15 incompatible with Tsien's and Ju's requirements for	15	Q. If the deprotection yield is not
16 sequencing by synthesis?	16	satisfactory, then certainly two hours with that
A. That is what I wrote. And that is what I	17	deprotection yield is not satisfactory?
18 meant.	18	A. Certainly.
19 Q. And any one of those requirements	19	Q. And that would not be compatible with
20 included the two hours?	20	Tsien's method?
21 A. Can I explain it?	21	A. For any efficiency component of that. As
22 Q. Did you write that?	22	we talked about yesterday, you can still do
23 A. I would like to explain it.	23	sequencing by synthesis, just not in any sort of
24 Q. Can you answer my question first?	24	efficient way that I believe Tsien required.
A. You can read it. That is correct.	25	Q. And that's true for SBS methods in
Page 265 1 general. It's not just for Tsien, correct?	1	Page 26 Q. And because of the natural dAPT
A. For anything to be efficient, if you have	2	contaminant you would end up with read through; is
3 a deprotection time that's part of the procedure	3	that correct?
4 then, yes, that would add time and make it	4	A. It would prevent termination.
5 effectively unrealistic to perform.	5	Q. And that would result in read through?
6 Q. I am going to ask you to go back to the	6	A. It would prevent termination. So I think
7 Metzker paper. The Metzker 1994 paper, I should	7	we're using I think we're saying the same thing.
8 say. It is Exhibit 1016.	8	Q. So please turn to Page 4263, right
9 A. I have it.	9	column, second full paragraph. Halfway through
10 Q. And please go back to Table 2 on	10	that paragraph Metzker says in Figure 3B do you
11 Page 4263. Yesterday we discussed how Table 2		
12 reports termination asterisk for the 3'-O-allyl	11	see that?
	11 12	see that? A. I do.
13 dAPT with the Vent polymerase.		
	12	A. I do.
13 dAPT with the Vent polymerase.	12 13	A. I do. Q. Metzker writes in Figure 3B "The O-allyl
13 dAPT with the Vent polymerase.14 Do you remember that?	12 13 14	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O
 13 dAPT with the Vent polymerase. 14 Do you remember that? 15 A. I remember the discussion. 	12 13 14 15	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT.
 dAPT with the Vent polymerase. Do you remember that? A. I remember the discussion. Q. And the asterisk indicates that there was 	12 13 14 15 16	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT. In addition to termination, some read
 dAPT with the Vent polymerase. Do you remember that? A. I remember the discussion. Q. And the asterisk indicates that there was incomplete termination, correct? 	12 13 14 15 16 17	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT. In addition to termination, some read through was also observed due to the presence of
 dAPT with the Vent polymerase. Do you remember that? A. I remember the discussion. Q. And the asterisk indicates that there was incomplete termination, correct? A. That is what is written. 	12 13 14 15 16 17 18	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT. In addition to termination, some read through was also observed due to the presence of contaminating dATP."
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 dAPT with the Vent polymerase. Do you remember that? A. I remember the discussion. Q. And the asterisk indicates that there was incomplete termination, correct? A. That is what is written. Q. And if I understand your position from yesterday, the incomplete termination is the result of the natural dAPT contaminate being incorporated, 	12 13 14 15 16 17 18 19 20 21	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT. In addition to termination, some read through was also observed due to the presence of contaminating dATP." That is what Metzker writes, correct? A. That is what is written. Q. So the read through reported for the 3'-O
 dAPT with the Vent polymerase. Do you remember that? A. I remember the discussion. Q. And the asterisk indicates that there was incomplete termination, correct? A. That is what is written. Q. And if I understand your position from yesterday, the incomplete termination is the result of the natural dAPT contaminate being incorporated, which allows synthesis to continue. 	12 13 14 15 16 17 18 19 20 21 22	 A. I do. Q. Metzker writes in Figure 3B "The O-allyl template gel also shows the incorporation of 3'-O methyl dAPT, compound one, by AMV RT. In addition to termination, some read through was also observed due to the presence of contaminating dATP." That is what Metzker writes, correct? A. That is what is written. Q. So the read through reported for the 3'-O methyl dATP for AMV indicates that there was

Floyd Romesberg, Ph.D.	University in the City of New York
Page 267	Page 268
1 thing.	1 that you still have I believe what termination
2 Q. And so let's look at Table 2 and focus on	2 means here is that you produced a banding pattern
3 the 3'-O methyl dAPT with the AMV reverse	3 the same as the dideoxy.
4 transcript, please.	4 Now, if there was a little bit of read
5 Do you see that?	5 through but effectively you produced exactly that
6 A. Yes.	6 same banding pattern, I believe that Metzker would
7 Q. And Metzker reports that there was	7 have called that complete termination. And that's
8 complete termination, correct?	8 simply because the contaminating 1 percent dAPT just
9 A. That is correct.	9 didn't effectively prevent all those bands from bein
10 Q. And Metzker reports that there was	10 formed due to termination.
11 complete termination even though there was read	11 And I assume that's because the 3'-O methyl
12 through, correct?	12 is very competitive with the dAPT. And the 3' allyl
13 A. Some read through he reports.	13 would have been less competitive with the dAPT.
14 Q. So the presence of read through caused by	14 So I think that's consistent with the
15 the natural dAPT contaminant did not lead Metzker	15 O-methyl being more competitive.
16 to report incomplete termination for the 3'-O	16 MR. SCHWARTZ: I would like to take a
17 methyl dAPT with that polymerase, correct?	17 break now and discuss whether I have any further
18 A. That is what is written. And what that	18 questions.
19 would mean is that the O-methyl is even better	19 THE VIDEOGRAPHER: We are off the record
20 incorporated.	20 at 10:04 a.m.
21 Q. But despite the presence of the read	21 (Recess taken.)
22 through, Metzker reports that as complete	22 THE VIDEOGRAPHER: We are on the record
23 termination?	23 at 10:25 a.m.
	,
25 there, but it might have been such a small amount	25 I have no further questions on this
Page 269	Page 270
1 cross-examination.	1 EXAMINATION
1 cross-examination.	1 EXAMINATION 2 BY MR. ZIMMERMAN:
1 cross-examination. 2 /// 3	 EXAMINATION BY MR. ZIMMERMAN: Q. Dr. Romesberg, I appreciate you being
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LIO	yd Romesberg, Ph.D.		University in the City of New York
1	Page 271 A. Yes.	4	Page 272
2	Q. If you look at Column 8 of Ju at Line 26.	1 2	Q. And how many carbons does MOM have?A. It has two carbons.
3	A. Yes.	2	
4	Q. Do you see there is preference to, quote,	4	Q. Leaving aside hydrogen, how many atoms does MOM have?
5	"A small chemical moiety such as."	5	A. Three.
6	A. I'm sorry. Where are you?	6	Q. And leaving aside hydrogen, how many
7	Q. Line 25.	7	atoms does allyl have?
8	A. Yes.	8	A. Three.
9	Q. Column 8. You will see it reads "A small	9	Q. If we could turn to the Tsien patent,
10	chemical moiety such as CH2OCH3 or CH2CH-double	10	
11	bond-CH2?	11	
12	A. Yes.	12	
13	Q. What is the second compound there?	13	-
14	A. That is the allyl.	14	
15	Q. And what is the first compound?	15	-
16	A. That is the MOM.	16	
17	Q. And does that show MOM as having three	17	, ,
18	-	18	· ·
19	A. Did I say three carbons? I apologize. I	19	0
20		20	
21	Q. So let me ask you the question again. Do	21	will recall, about the concentrations in that
22		22	
23	A. No, I'm sorry.	23	• •
24	Q. How many carbons does allyl have?	24	5 / 1
25	A. Allyl has three carbons.	25	-
	, ,		
1	Page 273 Q. And then there was comparisons of those	1	Page 274 Do vou recall a number of discussions
1	Q. And then there was comparisons of those	1	Do you recall a number of discussions
2	Q. And then there was comparisons of those concentrations to other references?		Do you recall a number of discussions during your examination about a POSA?
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	/d Romesberg, Ph.D.		University in the City of New York
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1	Page 275	1	Page 276 compound or multiple compounds?
2	A. No.	2	
3			A. That should be a single compound as he
-	Q. I would like you to look at the Metzker	3	has shown it here. He shows the structure of that
4	'94 article, which is Exhibit 1016.	4	compound in Figure 2.
5	Do you have that?	5	Q. And is that a substituted allyl?
6	A. Ido.	6	A. No, that is just the canonical allyl
7	Q. So between the publication of this paper	7	group.
8	and October of 2000, did Dr. Metzker publish any	8	Q. Are there any other 3'-O allyls in the
9	retraction or correction of this paper, to your	9	Metzker '94 paper?
10	knowledge?	10	A. No, there's not.
11	A. Not to my knowledge.	11	Q. Does he describe a genus of 3'-O allyls?
12	Q. And are you aware of any publication by	12	A. No, he does not.
13	Metzker or anyone in his group or anyone who	13	Q. Does he describe a substituted
14	reviewed this, that indicated the data in this	14	3'-O allyl?
15	publication was incorrect between '94 and	15	A. Not to my knowledge, no.
16	October 2000?	16	
17	A. No, I do not believe there were any such	17	
18	publications.	18	
19	Q. Now, if you look at Table 2 on Page 4263.	19	
20	A. Yes, I'm there.	20	
		20	•
21	Q. Entry No. 3.		
22	A. Yes.	22	
23	Q. 3'-O-allyl modified dATP?	23	5
24	A. That is correct.	24	
25	Q. Can you tell me if that is a single	25	(Recess taken.)
	Page 277		Page 278
1	THE VIDEOGRAPHER: We're on the record at	1	I, the undersigned, a Certified Shorthand
2	10:41 a.m.	2	Reporter of the State of California, do hereby
3	MR. SCHWARTZ: Dr. Romesberg, we don't	3	certify:
4	have any recross. Thank you for your time today.	4	That the foregoing proceedings were taken
5	THE WITNESS: You are welcome.	5	before me at the time and place herein set forth;
6	MR. ZIMMERMAN: Thank you.	6	that any witnesses in the foregoing proceedings,
7	THE VIDEOGRAPHER: This concludes today's	7	prior to testifying, were duly sworn; that a
8	deposition of Floyd Romesberg, Ph.D. We are off	8	
9			verbatim record of the proceedings was made by me
1 -		9	verbatim record of the proceedings was made by me using machine shorthand which was thereafter
10	the record at 10:41 a.m.	9 10	using machine shorthand which was thereafter transcribed under my direction; that the foregoing
10	the record at 10:41 a.m. (The videotaped deposition of		using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given.
11	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10	using machine shorthand which was thereafter transcribed under my direction; that the foregoing
11 12	the record at 10:41 a.m. (The videotaped deposition of	10 11	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal
11 12 13	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the
11 12 13 14	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal
11 12 13 14 15	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review
11 12 13 14 15 16	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested.
11 12 13 14 15 16 17	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially
11 12 13 14 15 16 17 18	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee
11 12 13 14 15 16 17 18 19	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action.
11 12 13 14 15 16 17 18 19 20	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date
11 12 13 14 15 16 17 18 19 20 21	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19 20	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date subscribed my name.
11 12 13 14 15 16 17 18 19 20 21 22	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19 20 21	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date subscribed my name.
11 12 13 14 15 16 17 18 19 20 21	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19 20 21 22	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date subscribed my name.
11 12 13 14 15 16 17 18 19 20 21 22	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19 20 21 22	<pre>using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date subscribed my name. Dated: September 21, 2018</pre>
11 12 13 14 15 16 17 18 19 20 21 22 23	the record at 10:41 a.m. (The videotaped deposition of FLOYD ROMESBERG, PH.D., VOLUME II,	10 11 12 13 14 15 16 17 18 19 20 21 22 23	using machine shorthand which was thereafter transcribed under my direction; that the foregoing transcript is a true record of the testimony given. Further, that if the foregoing pertains to the original transcript of a deposition in a Federal Case, before completion of the proceedings, review of the transcript [X] was [] was not requested. I further certify I am neither financially interested in the action nor a relative or employee of any attorney of party to this action. IN WITNESS WHEREOF, I have this date subscribed my name. Dated: September 21, 2018 MAMAA PATRICIA Y. SCHULER

	Dore 270	1				Page 280
1	Page 279 DECLARATION UNDER PENALTY OF PERJURY	1	DEPOSI	TION ERRATA	SHEET	Page 200
2	Case Name: Illumina, Inc. vs. The Trustees of Columbia	2	Case Na	ame: Illumi	.na, Inc. vs.	. The Trustees of Columbia
						City of New York
	University in the City of New York		Name of		Floyd Romesh	•
3	Date of Deposition: 09/20/2018	3			on: 09/20/201	5,
4	Job No.: 10047003			.: 10047003		
5		4			To clarify	the record.
6	I, FLOYD ROMESBERG, PH.D., hereby certify				-	to the facts.
7	that I have read the transcript of my deposition and that	5				transcription errors.
8	this transcript accurately states the testimony given by me,	6	Page		Reasor	-
9	with the changes or corrections, if any, as noted on an errata	7				to
10	sheet or errata sheets.	8			Reasor	
		9				 to
11	Executed this day of	10			Reasor	
12	, 2018, at	11	From			 to
13		12			Reasor	
14		13				 to
15	FLOYD ROMESBERG, PH.D.	14			Reasor	
16	NOTARIZATION (If Required)	15				to
17	State of	16			Reasor	
18	County of	17				to
19	Subscribed and sworn to (or affirmed) before me on	18			Reasor	
		19				 to
20	this day of, 20,	20			Reasor	
21	by, proved to me on the	21				 to
22	basis of satisfactory evidence to be the person	22			Reasor	
23	who appeared before me.	23				 to
24	Signature: (Seal)	24			Reasor	
25		25				 to
1	DEPOSITION ERRATA SHEET Page 281					
2	Page Line Reason					
3	From to					
4						
5	Page Line Reason From to					
6						
7	Page Line Reason					
8	From to Page Line Reason					
9	From to					
10	Page Line Reason					
11	From to					
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	Page Line Reason					
15	From to					
16	Page Line Reason					
17	From to					
18	Page Line Reason					
19 20	From to					
	Page Line Reason					
21	From to					
22	Subject to the above changes, I certify that the					
	transcript is true and correct					
23	No changes have been made. I certify that the					
24	transcript is true and correct.					
24						
1		1				
25	FLOYD ROMESBERG, PH.D.					

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