IN THE FOR THE M	UNITED STAT 4IDDLE DISTR	ES DISTRICT COURT ICT OF PENNSYLVANIA
TAMMY KITZMILLER, v. DOVER AREA SCHOOL	et al DISTRICT,	: CASE NO. 4:04-CR-002688 :
et al TR	RANSCRIPT OF BENCH	: PROCEEDINGS TRIAL
	MORNING	SESSION
BEFORE	E: HON. JOH	N E. JONES, III
DATE	: October 9:00 a.m	18, 2005
PLACE	: Courtroo Federal Harrisbu	m No. 2, 9th Floor Building rg, Pennsylvania
ВҮ	: Wendy C. U.S. Off	Yinger, RPR icial Court Reporter
APPEARANCES:		
ERIC J. ROTHSCHILD WITOLD J. WALCZAK, STEPHEN G. HARVEY, RICHARD B. KATSKEE THOMAS SCHMIDT, ES For the Plainti PATRICK T. GILLEN, RICHARD THOMPSON, ROBERT J. MUISE, E	D, ESQUIRE ESQUIRE ESQUIRE SQUIRE Effs ESQUIRE ESQUIRE ESQUIRE ESQUIRE	

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FOR THE DEFENDANTS	DIRECT	CROSS	REDIRECT	RECROSS
Michael Behe By Mr. Muise	3			

THE COURT: Good morning to all. Mr. Muise, 1 if it's Tuesday, we must be on the blood clotting. 2 3 MR. MUISE: We will be getting to blood clotting, immunity systems, and many more complex 4 5 systems, Your Honor. 6 THE COURT: All right. You may proceed. 7 MR. MUISE: Thank you. (Whereupon, Michael Behe, Ph.D., resumed the 8 9 stand and testimony continued.) DIRECT EXAMINATION (CONTINUED) 10 BY MR. MUISE: 11 12 Good morning, Dr. Behe. Q. 13 Α. Good morning. 14 Before we do get to the blood clotting, I need to Ο. 15 circle back to sort of cover one housekeeping matter. 16 MR. MUISE: If I may approach the witness, Your Honor? 17 THE COURT: Yes. 18 19 BY MR. MUISE: 20 0. Sir, I've handed you what has been marked as 21 Defendants' Exhibit No. 237, which is an article from Saier, correct? 22 23 Α. That's right. 24 Q. Is that one of the articles that you referenced 25 during your testimony and appeared on one of the slides

1	regarding the type III secretory system?
2	A. Yes, it is.
3	Q. Okay. Thank you, sir. Sir, yesterday, just to
4	sort of recap and bring us to where we need to begin
5	this morning, I had asked you if some scientists had
6	argued that there is experimental evidence that complex
7	biochemical systems can arise by Darwinian processes,
8	and I believe you indicated there were two that are
9	offered, correct?
10	A. That's right.
11	Q. And the first one was the lac operon?
12	A. Yes.
13	Q. And we discussed that yesterday?
14	A. Yes.
15	Q. And what is the second one?
16	A. The second one concerns what's called the blood
17	clotting cascade, the system for clotting blood in
18	animals. And I should say that, emphasize again that
19	this is the second example of an experimentally an
20	experimental result that was offered as evidence against
21	some of the arguments that I made in Darwin's Black Box.
22	In this one, this is directed more to the
23	question of irreducible complexity than to the question
24	of whether Darwinian processes can put together a
25	complex system.

1	Q. Now, sir, we've put up on the slide a figure,
2	6-5, that appears on page 142 in the Pandas text. Can
3	you explain what we see here?
4	A. That's right. This is an electron micrograph of
5	some red blood cells caught in a meshwork of a protein
6	called fibrin, which forms a blood clot. And most
7	people, when they think about blood clotting, if they
8	think about it at all, it appears to be a simple
9	process.
10	When somebody cuts themself, a minor cut slows
11	down, stops, and heals over, and it doesn't seem like
12	it doesn't seem like much at all. But thorough
13	investigation over the past 40 to 50 years has shown
14	that the blood clotting system is a very intricate
15	biochemical system. And I believe there's an
16	illustration of it on the next slide.
17	Q. Now you referred to, I believe, a blood clotting
18	cascade, is that correct?
19	A. That's right.
20	Q. Can you explain a little bit to us as you're
21	explaining what we see here on this particular diagram?
22	A. Okay, sure. Yeah, this is a figure of the blood
23	clotting cascade taken from the biochemistry textbook by
24	Voet and Voet, which is widely used in colleges and
25	universities around the country. You see all these

names of things and arrows. The names of things are 1 2 very complex proteins of the complexity or sometimes more complex than the hemoglobin that I showed 3 4 vesterday. 5 In blood clotting, the material that forms the clot cannot, of course, be in its solid clotted form 6 7 during the normal -- during the normal life of an animal or all of the blood would be clotted, and that would be 8 9 inconsistent with its life. So the material of the clot 10 that actual eventually forms the clot exists as 11 something called fibrinogen, which is actually a soluble 12 pre-cursor to the clot material. 13 It floats around in your bloodstream during 14 normal times. But when a cut occurs, fibrinogen is 15 transformed into something called fibrin, and that happens when another protein comes along and cuts off a 16 17 small piece of fibrinogen, a specific piece which 18 exposes a sticky site on it, sticky in the sense of 19 those two proteins yesterday that I saw that -- that I 20 showed you that had complimentary surfaces. 21 It exposes a sticky site on the surface of the 22 fibrinogen, which allows the many copies of fibrinogen, 23 now turned into fibrin, to aggregate and stick to each 24 other, forming the blood clot. But what is the component that cuts fibrinogen 25

and activates it? Well, the component is another protein called thrombin. But now we've got the same problem again. If thrombin were going around cutting fibrinogen and turning it into fibrin, all the blood would clot, and that would congeal the blood and kill the animal.

So thrombin itself is an inactive form called prothrombin, so it has to be activated when a cut occurs. And that's the responsibility of another protein. And that protein exists in an inactive form, and it's -- the activation of that is the responsibility of another protein.

So in the blood -- it's called a blood clotting 13 14 cascade because one component acts on the next which acts on the next which acts on the next and so on. Now 15 notice that the blood clotting cascade actually has what 16 are called two branches. There is one in this box up 17 18 here is labeled the intrinsic pathway. And this is labeled the extrinsic pathway. So there are actually 19 20 two branches to this blood clotting cascade. 21 I believe this section is addressed in the 0. 22 textbook Pandas, correct? 23 A. Yeah, that's correct. On the left is a figure 24 from Of Pandas and People illustrating the blood 25 clotting cascade. And that was drawn after the

1	illustration from the textbook by Voet and Voet. On the
2	right-hand side is the illustration for the blood
3	clotting cascade that appears in Darwin's Black Box.
4	I discussed the blood clotting cascade in one
5	chapter of that of my book, and the illustration is
6	very similar to the one in Pandas.
7	Q. I believe the diagram in Pandas is found on page
8	143?
9	A. Yes, that's right.
10	Q. Now these two diagrams, the one that appears in
11	Darwin's Black Box and one of the blood clotting cascade
12	appear, to my eye, to be virtually similar or almost
13	exactly similar?
14	A. Yeah, they are very similar, except for the color
15	in Pandas and so on. And that's because I wrote the
16	discussion in Pandas and, of course, also in my own
17	book. So the figures are very similar between the two.
18	Q. Now you testified yesterday that you coined the
19	term irreducible complexity in Darwin's Black Box, which
20	was published in 1996, is that correct?
21	A. Yes.
22	Q. So that book was published actually three years
23	after Pandas was written, is that accurate?
24	A. Yes, that's correct.
25	Q. Is it accurate to say then that the concept of

1	irreducible complexity was not fully developed when you
2	had written that section in Pandas on blood clotting in
3	1993?
4	A. Yes, that's right. I was still contemplating the
5	idea.
6	Q. Does Pandas, however, discuss the complexity of
7	this system, the blood clotting system?
8	A. Yes, it does. It elucidates all the parts of the
9	system.
10	Q. Is that discussion consistent with your
11	discussion in Darwin's Black Box?
12	A. Yes, it introduces the concept of the purposeful
13	arrangement of parts and says that's how we perceive
14	design.
15	Q. That's introduced in the Pandas book?
16	A. Yes, uh-huh.
17	${\tt Q}$ . When you talk about the purposeful arrangement of
18	parts, that's similar to what you were discussing
19	yesterday in your testimony, is that correct?
20	A. Yes.
21	Q. So is the scientific explanation of the blood
22	clotting system similar to the the discussion in
23	Pandas similar to the blood clotting cascade scientific
24	explanation in Darwin's Black Box?
25	A. That's right, they're essentially the same. I

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1	think it's more detailed in Darwin's Black Box
- -	O In fact way did was the similar diagrams?
2	Q. In fact, you did use the similar diagrams?
3	A. Yes, that's correct.
4	Q. To explain the two?
5	A. Yes, uh-huh.
6	Q. I believe the next slide we have is, this is from
7	your you discussed this and treated this as well in
8	your book Debating Design, is that correct?
9	A. That's right. When I wrote Darwin's Black Box,
10	and when Darwin's Black Box was subsequently reviewed by
11	people, some of them looked at the argument about the
12	blood clotting cascade and argued against what I had
13	written in Darwin's Black Box.
14	And I thought that the counterarguments were
15	themselves flawed, and so I answered some of those
16	arguments in a variety of cites, but most recently in
17	the chapter in that book, Debating Design, published by
18	Cambridge University Press from the year 2004.
19	I wrote The Blood Clotting Cascade. Having dealt
20	with some common misconceptions about intelligent
21	design, I will examine two systems that were proposed as
22	serious counterexamples of my claim of irreducible
23	complexity. One of them discussed in that article is
24	the blood clotting cascade.
25	Q. If you could then, explain to us how you refute

1	the claims that are made that the blood clotting cascade
2	is experimental evidence to refute irreducible
3	complexity?

Okay. In the next slide, I believe that shows an 4 Α. 5 excerpt from an article written by a man named Russell 6 Doolittle entitled A Delicate Balance, which appeared in 7 a publication called the Boston Review in 1997. Now Russell Doolittle is a very eminent scientist, a 8 9 professor of biochemistry at the University of 10 California, San Diego.

11 He's a member of the National Academy of 12 Sciences, and has worked on the blood clotting system for the past 45 years or so. And this article was a 13 14 part of the symposium organized by Boston Review, which again is published by MIT, and contained contributions 15 from a number of academics, scientists discussing my 16 17 book and discussing a book that had been recently 18 published by Richard Dawkins of Oxford University.

Participants included myself, Russell Doolittle, James Shapiro, who is a professor of microbiology at the University of Chicago, Alan Orr, who is a professor of evolutionary biology at the University of Rochester, Robert DiSilvestro, who is a professor of biochemistry at Ohio State, and a number of other people as well. And in his essay, Professor Doolittle argued

that, in fact, there was experimental evidence showing 1 2 that the blood clotting system was not irreducibly complex. And he said the following. Let me read the 3 quote. Quote, Recently the gene for plaminogen (sic) --4 5 and that's actually a typo. There should be an S there. 6 The gene for plaminogen (sic) was knocked out of mice -which means that it was destroyed by molecular 7 biological methods -- and predictable, those mice had 8 9 thrombotic complications because fibrin clots could not 10 be cleared away.

11 Let me stop a second and explain that plasminogen 12 is a protein that acts as a chemical scissors which cuts up and removes blood clots once the clot has finished 13 14 its job. Let me resume the quote from Russell 15 Doolittle. Not long after that, the same workers knocked out the gene for fibrinogen in another line of 16 17 mice. Again, predictably, these mice were ailing, 18 although in this case, hemorrhage was the problem.

Let me stop again and explain that fibrinogen, remind you, is the pre-cursor of the clot material itself, the pre-cursor of those fibers. And what do you think happened when these two lines of mice were crossed? For all practical purposes, the mice lacking both genes were normal.

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Contrary to claims about irreducible complexity,

the entire ensemble of proteins is not needed. 1 Music 2 and harmony can arise from a smaller orchestra. So Professor Doolittle's point, if I just might briefly 3 4 say, was that, if you knock out one component of the 5 blood clotting cascade, yes, those mice have problems. If you knock out a different component in a 6 7 different line of mice, yes, those mice have problems, too. But if you make a string of mice in which both of 8 9 those components were missing, then the mice are normal and the blood clotting cascade is okay. And so 10 11 presumably then, that shows that the blood clotting 12 cascade is not irreducibly complex. 13 Q. Was there a particular study that Professor 14 Doolittle is referring to? 15 Yes, it's shown on the next slide. This is the Α. article that he was referencing in his own essay. 16 It's 17 entitled Loss of Fibrinogen Rescues Mice from the 18 Pleiotropic Effects of Plasminogen Deficiency. Now if we could go to the next slide. 19 20 Now because of the phrase, rescues mice, in the 21 title, Professor Doolittle thought that the mice missing 22 both components were normal. But it turns out, that was 23 a misreading of the article. 2.4 In the abstract of the article itself, the 25 authors write, quote, Mice deficient in plasminogen and

1 fibrinogen are phenotypically indistinguishable from 2 fibrinogen deficient mice. Now translated that into 3 English on the next slide.

That means that mice missing both components have 4 5 all the problems that mice missing fibrinogen only have. 6 Their blood does not clot. They hemorrhage. Female 7 mice die during pregnancy. They are not normal. Thev are not promising evolutionary intermediates. So if we 8 9 look at this table of the symptoms of the various strings of mice, we can see what the authors meant by 10 11 that phrase, rescues mice.

Lacking plasminogen, mice can't remove blood clots once their job is done and their blood circulation gets interfered with and they develop problems such as thrombosis, ulcers, and so on. Lacking fibrinogen, they can't clot blood in the first place, and they have a different suite of symptoms.

18 When they lack both, they have been rescued from the symptoms of plasminogen deficiency, but only to 19 20 suffer the symptoms of fibrinogen deficiency. And if 21 you think about it for just a minute, it's easy to understand what is going on. When an animal lacks 22 23 plasminogen, it can't remove blood clots and its 2.4 circulation becomes impeded and it suffers problems. 25 Lacking fibrinogen, it can't make clots in the

1 first place, and so hemorrhage is a problem. Lacking 2 both, it doesn't matter that it's lacking plasminogen, 3 because the plasminogen's job is to remove blood clots 4 after the job is finished. But the mouse missing both 5 components can't form clots in the first place. So 6 there are no clots to remove.

Q. Has subsequent work verified those results?
A. Yes, here's a table of not only the work that was
9 cited in this discussion here on plasminogen fibrinogen,
10 but also subsequent work by the same group of scientists
11 who knocked out other components of the blood clotting
12 cascade, including something called prothrombin and
13 something else called tissue factor.

And if you look at the -- under the column labeled effect, in each case the blood clotting cascade is broken. They suffer hemorrhage. They cannot clot their blood. And that is exactly the result you would expect if, in fact, the blood clotting cascade were irreducibly complex, as I had written.

Q. So Professor Doolittle's refutation of your claims was based on a misreading of the study, is that correct?

A. That's right. He misread the original paper that he pointed to. And if I could make a couple of points based on this. As I said, this study, or this essay by Professor Doolittle and the one I discussed yesterday by Professor Miller were the two examples which offered experimental evidence that either irreducible complexity was not correct or that random mutation and natural selection could explain complex biochemical systems.

6 But if you look at the exact studies that were 7 offered as support for Darwinian evolution, and you look 8 at them closely, in reality, they highlight the 9 difficulties for Darwinian evolution. So I think this 10 is an illustration of how a scientist's preconceptions 11 about the truth of a theory or the validity of a theory 12 can affect his reading of the evidence.

And one more point is that, Professor Doolittle, of course, is a very eminent scientist. Professor Miller is, too. And they're quite capable of surveying the entire scientific literature for studies that they think are problems for my argument for intelligent design.

And nonetheless, when they surveyed the whole literature, and they seemed to be motivated to look for counterexamples to intelligent design, when they do so, they offer studies such as this, which are, at best, very problematic and none of which, I would say, are arguments against intelligent design.

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So in my mind, I conclude that since highly

motivated capable scientists who could advance arguments
or who could point to studies that have created problems
for intelligent design, that they have failed to do so,
makes me confident that intelligent design is a good
explanation.
Q. Now these article findings, the actual findings
in these articles, is that what you would expect to find
for an irreducibly complex system?

9 A. Yes, that's right. This is completely consistent with my expectations. 10

11 Q. As far as you know, has Professor Doolittle ever 12 acknowledged that he misread that paper?

A. Yes, he has.

0. And if I could --

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15 MR. ROTHSCHILD: Objection. Hearsay, Your Honor. I would move to strike. 16

MR. MUISE: Your Honor, he just -- he has an 17 18 understanding that Professor Doolittle has indicated he 19 has misread this paper.

20 MR. ROTHSCHILD: If he has a basis, I'd like 21 to see it.

22 THE COURT: Well, it's his understanding, 23 and I'm take it for that. I won't take it as a matter 24 of fact. His understanding is, he didn't quote 25 something that Professor Doolittle said. It's simply,

1	I'll take it as his understanding, and you're free to
2	cross-examine him and present rebuttal evidence, if you
3	see fit. So it's overruled.
4	BY MR. MUISE:
5	Q. Dr. Behe, I'd ask you to look at the exhibit
6	binder that I had provided you yesterday. It's at your
7	table in front of you. If you go to tab 17, please.
8	A. Yes.
9	Q. You'll see an exhibit marked Defendants' Exhibit
10	272. Is that the article by Russell Doolittle that
11	you've been referring to here in your testimony?
12	A. Yes, that's correct. This is a web version.
13	MR. ROTHSCHILD: Objection, Your Honor. I
14	want to make clear, I think that's not the
15	acknowledgment of the mistake, it's just the article
16	that's being referred to. I just want to clarify that.
17	MR. MUISE: I think the question was pretty
18	clear.
19	BY MR. MUISE:
20	Q. That's the article in the Boston Review that
21	you're referring to?
22	A. Yes, this is Russell Doolittle's article in the
23	Boston Review.
24	THE COURT: Does that resolve the objection?
25	MR. ROTHSCHILD: Yes. I just want to

1	clarify, this was not Dr. Doolittle's acknowledgment of
2	a mistake.
3	THE WITNESS: Yes.
4	THE COURT: All right.
5	BY MR. MUISE:
6	Q. Dr. Behe, does anyone else know how the blood
7	clotting cascade can be explained in Darwinian fashion
8	and other proposed examples or explanations?
9	A. No, that's one of the very nice things about
10	science is that, if there is no explanation in the
11	science library in scientific literature, and if leaders
12	in the field do not know how something could have come
13	about, and presumably they know the literature very,
14	very well, then one can be confident that not only do
15	they not know how something could have been done, but
16	nobody else in the world knows how that could have been
17	done as well. And that's important to keep in mind
18	because some people claim that nonetheless.
19	Q. And that's my next question. There have been
20	individuals that nonetheless have made such claims, and
21	do you have some slides to bring that up?
22	A. Yes, that's correct. On the next slide is an
23	excerpt from an article by a man named Michael Ruse.
24	Michael Ruse is a professor of philosophy of science
25	currently at Florida State University. And in

1 particular, he's a philosopher interested in Darwinian 2 thought.

And he's written many books on Darwin, his ideas, the history around them, and so on. And several years after my book came out in 1998, Professor Ruse wrote an article entitled Answering the Creationists, Where They Go Wrong and What They're Afraid Of, and had it published in a magazine called Free Inquiry. And he said the following in the article.

Quote, For example, Behe is a real scientist, but this case for the impossibility of a small-step natural origin of biological complexity has been trampled upon contemptuously by the scientists working in the field. They think his grasp of the pertinent science is weak and his knowledge of the literature curiously, although ventsly, outdated.

17 For example, far from the evolution of clotting 18 being a mystery, the past three decades of work by 19 Russell Doolittle and others has thrown significant 20 light on the ways in which clotting came into being. 21 More than this, it can be shown that the clotting 22 mechanism does not have to be a one-step phenomenon with 23 everything already in place and functioning. One step 24 in the cascade involves fibrinogen, required for 25 clotting, and another, plaminogen -- there's that typo,

missing the S -- required for clearing clots away. 1 2 And he goes on in his article to quote that passage from Russell Doolittle's Boston Review essay 3 that I showed on the slide a couple slides ago. So this 4 5 excerpt, in my view, shows that Professor Ruse relies completely on Professor Doolittle's explanation for the 6 7 blood clotting cascade and has no independent knowledge of his own. 8

9 As a matter of fact, the fact that the same typo, the same misspelling of plasminogen occurs in Professor 10 Ruse's essay makes me think that he relied on Professor 11 12 Doolittle even for the spelling of the components of the cascade. So the point is that, even though Professor 13 14 Ruse is a prominent academic concerned with Darwin and 15 Darwinian thought, he has no knowledge that Professor 16 Doolittle does not have concerning the blood clotting cascade. 17

18 Do you have another example, sir? 0. 19 Yes, another person has written on this, a man Α. 20 named Neil Greenspan, who is a professor of pathology at 21 Case Western Reserve University, and he wrote an article 22 in a magazine called The Scientist in the year 2002 23 entitled Not-so-intelligent Design. In the article, he 24 writes the following. Quote, The Design advocates also ignore the accumulating examples of the reducibility of 25

biological systems. As Russell Doolittle has noted in 1 2 commenting on the writings of one ID advocate -- and perhaps I can be forgiven if I think he means me -- mice 3 genetically altered so they lack either thrombin or 4 5 fibrinogen have the expected abnormal hemostatic 6 phenotypes. However, when the separate knockout mice 7 are bred, the double knockouts apparently have normal hemostasis, reducible complexity after all, at least in 8 9 the laboratory.

10 So the reasoning here exactly mimics the reasoning of Russell Doolittle in his Boston Review 11 12 article. And let me just point out here that he talks about thrombin or fibrinogen, but the study was actually 13 14 on plasminogen and fibrinogen. So again, I think this illustrates that even a scientist has -- even a 15 scientist writing publicly on this topic, even a 16 17 scientist writing publicly on this topic in order to 18 arque against intelligent design has no more knowledge of this than Professor Doolittle has. 19

And once more, I think this speaks to the point of how firmly a theory can guide persons' thinking. I think the fact that Professor Ruse relied so heavily on Professor Doolittle, and Professor Greenspan did, too, and apparently they did not even go back and read the article on blood clotting that was being disputed, shows

1	that they are so confident in Darwinian evolution that
2	they don't think they have to, you know, check the
3	facts.
4	They can rely on the authority of a person like
5	Professor Doolittle. So I think that shows the grip of
6	a theory on many people's thinking.
7	Q. Do you have an additional example?
8	A. Yes, one other excerpt here. In 1999, the
9	National Academy of Sciences issued a booklet called
10	Science and Creationism. And in it, they write the
11	following, quote, The evolution of complex molecular
12	systems can occur in several ways. Natural selection
13	can bring together parts of a system for one function at
14	one time, and then at a later time, recombine those
15	parts with other systems of components to produce a
16	system that has a different function.
17	Genes can be duplicated, altered, and then
18	amplified through natural selection. The complex
19	biochemical cascade resulting in blood clotting has been
20	explained in this fashion.
21	Let me make a comment on this. Professor
22	Doolittle is a member of the National Academy of
23	Sciences. There is no other member of the National
24	Academy who knows anything more about blood clotting
25	than Professor Doolittle. But if Professor Doolittle

1	does not know how Darwinian processes could have
2	produced the blood clotting cascade, as I think is
3	evident from his pointing to an inappropriate paper in
4	his attempt to refute a challenge to Darwinian
5	evolution, then nobody in the National Academy knows
6	either. I should also well, I'll
7	Q. Do they cite any papers or experiments to support
8	this claim, the National Academy of Sciences, in this
9	particular booklet?
10	A. No. That's a very interesting point. They
11	simply assert this. They do not cite any paper in any
12	journal to support this. And it's an interesting point,
13	if I may say so. I've heard said earlier in this trial
14	that not every utterance by a scientist is a scientific
15	statement.
16	And that's something that I entirely agree with.
17	And it's also true that not every utterance by a
18	scientist even on science is a scientific statement.
19	And it's also true that not even, not every
20	proclamation, or not every declaration by a group of
21	scientists about science is a scientific statement.
22	Scientific statements have to rely on physical
23	evidence. They have to be backed up by studies. And
24	simply saying that something is so does not make it so.
25	In fact, this statement of the National Academy is

simply an assertion. It is not a scientific statement.
Q. Does the National Academy of Sciences, in this
document that you referenced, give any other examples of
complex biochemical systems that have been explained?
A. This is the only example that they point to.
Q. In his testimony, Dr. Miller has pointed to the
work of, I believe, you pronounce is Jiang, J-i-a-n-g
A. Yes.
Q and Doolittle and Davidson, et al, to argue
against the irreducible complexity of the blood clotting
system. Do you agree with his assessment of those
studies?
A. No, I do not.
Q. And you have some diagrams to explain this
further, sir?
A. Yes, I do. This is a slide from Professor
Miller's presentation showing work from Jiang and
Doolittle. And he also shows a diagram of the blood
clotting cascade. And notice again, it's a branched
pathway with the intrinsic pathway and the extrinsic
pathway.
And Professor Miller makes the point that in DNA
sequencing studies of something called a puffer fish,
where the entire DNA of its genome was sequenced, and
scientists looked for genes that might code for the

1 first couple components of the intrinsic pathway, they
2 were not found.

And so Professor Miller demonstrated that by -if you could push to start the animation -- Professor Miller demonstrated that by having those three components blanked out in white. Nonetheless, puffer fish have a functioning clotting system. And so Professor Miller argued that this is evidence against irreducible complexity.

But I disagree. And the reason I disagree is that I made some careful distinctions in Darwin's Black Box. I was very careful to specify exactly what I was talking about, and Professor Miller was not as careful in interpreting it.

In Darwin's Black Box, in the chapter on blood 15 clotting cascade, I write that, a different difference 16 17 is that the control pathway for blood clotting splits in 18 two. Potentially then, there are two possible ways to 19 trigger clotting. The relative importance of the two 20 pathways in living organisms is still rather murky. 21 Many experiments on blood clotting are hard to do. And 22 I go on to explain why they must be murky.

And then I continue on the next slide. Because of that uncertainty, I said, let's, leaving aside the system before the fork in the pathway, where some

details are less well-known, the blood clotting system 1 fits the definition of irreducible complexity. 2 And I noted that the components of the system 3 beyond the fork in the pathway are fibrinogen, 4 5 prothrombin, Stuart factor, and proaccelerin. So I was 6 focusing on a particular part of the pathway, as I tried 7 to make clear in Darwin's Black Box. If we could go to the next slide. 8 Those 9 components that I was focusing on are down here at the 10 lower parts of the pathway. And I also circled here, 11 for illustration, the extrinsic pathway. It turns out 12 that the pathway can be activated by either one of two directions. And so I concentrated on the parts that 13 14 were close to the common point after the fork. 15 So if you could, I think, advance one slide. Ιf you concentrate on those components, a number of those 16 17 components are ones which have been experimentally 18 knocked out such as fibrinogen, prothrombin, and tissue 19 factor. 20 And if we go to the next slide, I have red arrows 21 pointing to those components. And you see that they all 22 fall in the area of the blood clotting cascade that I 23 was specifically restricting my arguments to. And if 24 you knock out those components, in fact, the blood 25 clotting cascade is broken. So my discussion of

1	irreducible complexity was, I tried to be precise, and
2	my argument, my argument is experimentally supported.
3	Q. Now just by way of analogy to maybe help explain
4	further. Would this be similar to, for example, a light
5	having two switches, and the blood clotting system that
6	you focus on would be the light, and these extrinsic and
7	intrinsic pathways would be two separate switches to
8	turn on the system?
9	A. That's right. You might have two switches. If
10	one switch was broke, you could still use the other one.
11	So, yes, that's a good analogy.
12	Q. So Dr. Miller is focusing on the light switch,
13	and you were focusing on the light?
14	A. Pretty much, yes.
15	Q. I believe we have another slide that Dr. Miller
16	used, I guess, to support his claim, which you have some
17	difficulties with, is that correct?
18	A. Yes, that's right. Professor Miller showed these
19	two figures from Davidson, et al, and from Jiang, et al,
20	Jiang and Doolittle, and said that the suggestions can
21	be tested by detailed analysis of the clotting pathway
22	components.
23	But what I want to point out is that whenever you
24	see branching diagrams like this, especially that have
25	little names that you can't recognize on them, one is

1 talking about sequence comparisons, protein sequence 2 comparisons, or DNA nucleotide sequence comparisons. As 3 I indicated in my testimony yesterday, such sequence 4 comparisons simply don't speak to the question of 5 whether random mutation and natural selection can build 6 a system.

7 For example, as I said yesterday, the sequences of the proteins in the type III secretory system and the 8 9 bacterial flagellum are all well-known, but people still 10 can't figure out how such a thing could have been put together. The sequences of many components of the blood 11 12 clotting cascade have been available for a while and were available to Russell Doolittle when he wrote his 13 14 essay in the Boston Review.

And they were still unhelpful in trying to figure out how Darwinian pathways could put together a complex system. And as we cited yesterday, in Professor Padian's expert statement, he indicates that molecular sequence data simply can't tell what an ancestral state Was. He thinks fossil evidence is required.

So my general point is that, while such data is interesting, and while such data to a non-expert in the field might look like it may explain something, if it's asserted to explain something, nonetheless, such data is irrelevant to the question of whether the Darwinian

1	mechanism of random mutation and natural selection can
2	explain complex systems.
3	Q. So is it your opinion then, the blood clotting
4	cascade is irreducibly complex?
5	A. Yes, it is.
6	Q. Now Professor Pennock had testified that he was
7	co-author on a study pertaining to the evolution of
8	complex features. Does this study refute the claim of
9	irreducible complexity?
10	A. No, it does not.
11	Q. And I believe we put up a slide indicating the
12	paper that was apparently by Lenski and Pennock,
13	correct?
14	A. That's right. Richard Lenski, and Professor
15	Pennock was co-author, and several other co-authors as
16	well. This is the first page of that article. Let me
17	reemphasize that the last two systems that I talked
18	about, the lac operon and the blood clotting cascade
19	were ones in which experiments were done on real
20	biological organisms to try to argue against intelligent
21	design and irreducible complexity.
22	This study of Lenski is a computer study, a
23	theoretical study not using live organisms, one which is
24	conducted by writing a computer program and looking at
25	the results of the computer program.

If I could have the next slide. This is an 1 2 excerpt from the abstract of that paper. Let me read 3 parts of it. It says, quote, A long-standing challenge to evolutionary theory has been whether it can explain 4 5 the origin of complex organismal features, close quote. 6 Let me just stop there to emphasize that these workers 7 admit that this has been a long-standing problem of 8 evolutionary theory. 9 MR. ROTHSCHILD: Objection. This 10 mischaracterizes the document. 11 THE COURT: Elaborate on that objection. 12 MR. ROTHSCHILD: I'm sorry? 13 THE COURT: Elaborate on the objection. You 14 say he's mischaracterizing --15 MR. ROTHSCHILD: This is a long-standing 16 challenge not a long-standing problem. 17 THE COURT: Well, I think he's 18 characterizing something and not necessarily reading 19 from it. What are you objecting to? MR. ROTHSCHILD: I think he's 20 21 mischaracterizing it. That's my objection. 22 THE COURT: Again, you'll have him on cross. This is direct examination. I'll overrule the 23 24 objection. You may proceed. 25 BY MR. MUISE:

1	
1	O Dr Bobo just for reference the article you are
T	Q. DI. Bene, just for reference, the article you are
2	referring to is published in 2003, is that correct?
3	A. That's correct, yes.
4	Q. Continue, please.
5	A. So apparently, this had not been explained up
6	until at least the publication of this paper. The
7	authors continue, quote, We examined this issue using
8	digital organisms, computer programs that
9	self-replicate, mutate, compete and evolve. Let me
10	close quotes there.
11	You have to remember that the labeling of these
12	things as organisms is just a word. These things are
13	not flesh and blood. These things are little computer
14	programs. There are strings of instructions. And a
15	comparison of these to real organisms is kind of like
16	comparing an animated character in some movie to a real
17	organism.
18	So the authors go on. And the next slide,
19	please. And this is the first figure on the first page
20	of their article. And I just want to emphasize, this is
21	just an illustration emphasizing that these there are
22	computer instructions. Each one of these are little
23	computer instructions; swap, nand, nand, shift R. They
24	have no similarity to biological features, biological
25	processes. You see over here little strings of ones and

1 zeroes.

2	These are characters in a computer memory. These
3	are not anything biological. Let me say that,
4	theoretical studies of biology can oftentimes be very
5	useful. And I'm certainly not denigrating the use of
6	computer in studying biology. But one has to be
7	careful, very careful that one's model, computer model
8	mimics as closely as possible a real biological
9	situation. Otherwise, the results one obtains really
10	don't tell you anything about real biology.
11	And I think that the Lenski paper, it does not
12	mimic biology in the necessary way. And that's shown on
13	the next slide.
14	Q. Let me just, to clarify. So a crucial question
15	is whether or not it's a good model for biological
16	process, is that correct?
17	A. Yes, that's right.
18	Q. And you don't believe this is one?
19	A. No, I think it misses the point and it assumes
20	what should be proven instead. And let me try to
21	explain that with an excerpt from the article itself.
22	The authors write in their discussion, quote, Some
23	readers might suggest that we stacked the deck by
24	studying the evolution of a complex feature that could
25	be built on simpler functions that were also useful,

1 close quote.

2	Let me stop there to comment that, yes, that is
3	exactly what I would suggest, that they stacked the
4	deck. They built a model in which there was a
5	continuous pathway of functional Features very close
6	together in probability, which is exactly the question
7	that's under dispute in real biological organisms. Is
8	there such a pathway in real biological organisms?
9	So to assume that in your computer model is
10	stacking the deck. Let me go back to the abstract.
11	They continue, quote, However, that is precisely what
12	evolutionary theory requires. Now I'll close quote
13	there, and let me comment on that.
14	Just because your theory requires something does
15	not mean it exists in nature. James Clerk Maxwell's
16	theory required ether. Ether does not exist. So just
17	because a theory requires it is no justification for
18	saying that building a model shows something about
19	biology.
20	Q. Dr. Behe, if you could, just so we're clear on
21	the record, because I'm not sure if we have it that
22	clear, can you identify the title and the specifics of
23	this article, so we're clear on what specific article
24	you're referring to?
25	A. Yes, this is an article by Lenski, Ofria,

Pennock, and Adami published in the year 2003. 1 The 2 title is The Evolutionary Origin of Complex Features published in the journal Nature, volume 423, pages 139 3 to 144. 4 5 Thank you. And the authors go on to say in their Q. discussion, indeed, our experiments showed that the 6 7 complex feature never evolved when simpler functions 8 were not rewarded. This is not surprising to me. This 9 shows the difficulty of irreducible complexity. If you do not have those closely stacked functional states, if 10 you have to change a couple things at once before you 11 12 get a selectable property, then I have been at pains to 13 explain, that's when Darwinian theory starts to fail, 14 not when you have things close together. 15 And to build them into your model is, again, begging the question. The fact that when they do not 16 17 build that into their model, they run into problems that 18 complex features then don't evolve. That is exactly what I would expect. I would cite this as evidence 19 20 supporting my own views. 21 Have other scientists made similar criticisms? Ο. 22 Α. Yes. A couple years ago, there was an article 23 published by two scientists named Barton and Zuidema 24 published in a journal called Current Biology. The 25 title of the article is Evolution, The Erratic Path

1 Towards Complexity.

2	And much of the article is a commentary on the
3	work by Lenski and co-workers. And if I could just read
4	a couple excerpts from that article. They make a couple
5	interesting points. The authors say, complex systems,
6	systems whose function requires many interdependent
7	parts, that is irreducible complexity systems in my
8	view, are vanishingly unlikely to arise purely by
9	chance.
10	Darwin's explanation of their origin is that
11	natural selection establishes a series of variants, each
12	of which increases fitness. This is an efficient way of
13	sifting through an enormous number of possibilities,
14	provided there is a sequence of ever-increasing fitness
15	that leads to the desired feature, close quote.
16	So that's the exact that's the big question.
17	Is there such a pathway, or is it, as it certainly
18	appears, that one has to make large numbers of changes
19	before one goes from a functional selectable state to a
20	second functional selectable state? And Barton and
21	Zuidema continue in their discussion.
22	They say, in Lenski's artificial organisms, the
23	mutation rate per site is quite high. So, in other
24	words, if I might make my own comment, they are using
25	they are using factors which are not common for
1 biological organisms.

Now picking up with the paper again. So that favorable pairs can be picked up by selection at an appreciable rate. This would be unlikely in most real organisms because, in these, mutation rates at each locus are low. In other words, again, they are building into the model exactly the features they need to get the result they want.

9 But building it into your model does not show 10 that that's what exists in nature. And Barton and 11 Zuidema comment further, quote, Artificial life models 12 such as Lenski, et al's, are perhaps interesting in 13 themselves, but as biologists, we are concerned here 14 with the question of what artificial life can tell us 15 about real organisms.

It's -- it can be productive and it can be interesting to do such studies as Lenski, et al, did. But the big question is, do they tell us anything about real organisms? And I am very skeptical that this study does so.

21 Q. Now have you done some work yourself that's 22 somewhat similar?

A. Yes, indeed. A year ago, as I mentioned earlier
in my testimony, David Snoke and myself published a
paper in the journal Protein Science entitled Simulating

Evolution by Gene Duplication of Protein Features that 1 Require Multiple Amino Acid Residues. 2 In this, we also -- it was essentially a 3 theoretical study using computer programs to try to 4 5 mimic what we thought would occur in biology. But we 6 tried, as closely as possible, to mimic features of real 7 proteins and real mutation rates that the professional 8 literature led us to believe were the proper reasonable 9 values. 10 And when we used those values, the short, the 11 gist of the matter is that, once -- if there is not a 12 continuous pathway, if one has to make two or three or 13 four amino acid changes, those little changes from that 14 figure of two interacting proteins that I talked about 15 yesterday, if one has to make several changes at once, then the likelihood of that occurring goes -- drops 16 17 sharply in the length of time, and the number of 18 organisms in a population that one would need to have 19 that goes up sharply. 20 Q. Would it be fair to say that your model is closer 21 to biological reality? 22 Α. Well, I certainly think so. Now Dr. Miller testified that the immune system 23 0. 24 is being explained by Darwinian theory. Do you agree 25 with that?

1	
1	A No I do not
⊥	A. No, I do not.
2	Q. And so i'd ask you ii you could explain why not?
3	A. Yes. On the next slide is a is the first
4	slide of Professor Miller's discussion of this topic and
5	his presentation simply showing a model of an
6	immunoglobulin protein. And here is kind of a little
7	cartoon version of the same thing, the immunoglobulin
8	protein.
9	He goes on the next slide to take an excerpt from
10	my book where in a chapter where I discussed the immune
11	system and argue that, in fact, it is not well-explained
12	by Darwinian processes but, in fact, is better explained
13	by design.
14	Q. Can you explain that Sisyphus reference?
15	A. Yeah, okay. Sisyphus. I said, Sisyphus himself
16	would pity us. That was just a literary flourish there.
17	Sisyphus is a figure from mythology who was doomed for
18	eternity to have to roll a bolder up a hill, and
19	whenever he got to the top of the hill, the bolder would
20	roll back, and he would have to start all over again.
21	This was meant to indicate frustration. And I
22	argued that Darwinian attempts at explanations would be
23	similarly frustrating.
24	Q. I just want to make a point clear. You said
25	there were two examples where those who claim that

irreducible complexity does not work or is not a valid 1 2 explanation, they use experimental evidence, and that was the blood clotting system and the lac operon. How 3 does the immunity system, is that experimental evidence 4 or is that a theoretical claim? 5 No, this is mostly a theoretical claim. There is 6 Α. 7 no experimental evidence to show that natural selection 8 could have produced the immune system. And I think 9 that's a good example of the different views that people 10 with different theoretical frameworks bring to the 11 table. 12 If we could show the next slide. Professor Miller shows this slide from a reference that he cited 13 14 by Kapitonov and Jurka, and he has titled Summary, Between 1996 and 2005, each element of the transposon 15 hypothesis has been confirmed. He has this over this 16 17 diagram. 18 But again, as I mentioned previously, whenever you see diagrams like this, we're talking about sequence 19 20 data, comparison of protein, sequences, or gene 21 sequences between organisms. And such data simply can't speak to the question of whether random mutation and 22 23 natural selection produced the complex systems that

24 we're talking about.

25

So Professor Miller -- so, in my view, this data

does not even touch on the question. And yet Professor 1 2 Miller offers as compelling evidence. And one more time, I view this as the difference between two people 3 with two different expectations, two different 4 5 theoretical frameworks, how they view the same data. And I'd like to take a little bit of time to 6 7 explain why such studies do not impress me. And I'll do 8 so by looking at one of the papers that Professor Doolittle -- I'm sorry, Professor Miller, that's his 9 name, cited in his presentation, Kapitonov and Jurka, 10 11 that was published this year. 12 I just want to go through, and just kind of as a quick way to show why I am not persuaded by these types 13 14 of studies. I want to excerpt some sentences from this study to show what I consider to be the speculative 15 nature of such studies. 16 17 For example, in this excerpt, the authors say, 18 something indicates that they may be important. This 19 may indicate. It may be encoded. It might have been 20 added. If so, it might have been derived. 21 Alternatively, it might have been derived from a 22 separate unknown transposon. It was probably lost. And 23 we have a lot more of those, one more slide at least. 24 It says, we cannot exclude the possibility. Ιn any case, the origin appears to be a culmination of 25

1 earlier evolutionary processes. If so, this might have 2 been altered. Again, without going into the detail of 3 the article, I just wanted to emphasize those phrases to 4 point out what I consider to be the very speculative 5 nature of such papers.

6 Here's what I view to be the problem. The 7 sequence of the proteins are there. The sequence of the 8 genes are experimentally determined. And the question 9 is, what do we make of that information? People like 10 Professor Miller and the authors of this paper working 11 from a Darwinian framework simply fit that data into 12 their framework.

But to me, that data does not support their framework. It does not offer experimental evidence for that framework. They're simply assuming a background of Darwinian random mutation and natural selection and explaining it -- or fitting it into that framework, but they're not offering support for it. Q. Dr. Behe, is there another paper that scientists

20 point to for the support that the immune system can be 21 explained by this Darwinian process?

A. Yes, there is. There is one more that I have to
discuss. Here is a recent paper, again the year 2005,
by Klein and Nikolaidis entitled The Descent of the
Antibody-Based Immune System by Gradual Evolution. And

1	on the next slide is an excerpt from the initial part of
2	their discussion where they say, quote, According to a
3	currently popular view, the Big Bang hypothesis, the
4	adaptive immune system arose suddenly, within a
5	relatively short time interval, in association with the
6	postulated two rounds of genome-wide duplications.
7	So these people, Klein and Nikolaidis, are going
8	to argue against what is the currently popular view
9	among immunologists and people who study the immune
10	system on how that system arose.
11	Q. And what is the Big Bang hypothesis that's
12	referred to here?
13	A. Well, that's kind of a label that they put on to
14	kind of indicate the fact that the immune system appears
15	in one branch of animals, the vertebrates, and any
16	obvious pre-cursors or functional parts of such a system
17	do not appear to be obvious in other branches of
18	animals.
19	So it seems like the immune system arose almost
20	complete in conjunction with the branching of
21	vertebrates from invertebrate.
22	Q. Do scientists acknowledge that or treat that as a
23	problem for Darwin's theory?
24	A. Well, in my experience, no, nobody treats such a
25	thing as a problem for Darwin's theory.

1	Q. Do you consider it a problem?
2	A. I certainly consider it a problem. But other
3	scientists who think that Darwinian evolution simply is
4	true don't consider much of anything to be a problem for
5	their theory.
6	Q. Why do you consider it a problem?
7	A. Because the as Darwin insisted, he insisted
8	that adaptations had to arise by numerous successive
9	slight modifications in a very gradual fashion. And
10	this seems to go against the very gradual nature of his
11	view.
12	Q. Now has this paper been held up by scientists as
13	refuting claims against intelligent design?
14	A. Yes, it has. As a matter of fact, Professor
15	Miller cited it in his expert report, although he didn't
16	refer to it in his testimony. Additionally, I attended
17	a meeting on evolution at Penn State in the summer of
18	2004 where one of the authors, Juan Kline, spoke on his
19	work, and he interpreted it in those terms.
20	Q. Now we have some quotes, I believe, from this
21	paper that you want to highlight?
22	A. Yes. Again, I want to pull out some excerpts
23	from that paper just to show you why I regard this as
24	speculative and unpersuasive. For example, they start
25	with, by saying, quote, Here, we sketch out some of the

1 changes and speculate how they may have come about. We 2 argue that the origin only appears to be sudden. They 3 talk about something as probably genuine.

It probably evolved. Probably would require a 4 5 few substitutions. It might have the potential of signaling. It seems to possess. The motifs presumably 6 7 needed. One can imagine that a limited number. Ιt might have been relatively minor. Quote, The kind of 8 9 experimental molecular evolution should nevertheless 10 shed light on events that would otherwise remain 11 hopelessly in the realm of mere speculation. They're 12 talking about experiments that have yet to be done.

Next slide, I have even more such quotations.
These factors are probably genuine. Nonetheless. They
might have postdated. Nevertheless. Albeit. It seems.
This might have been. These might represent. They
might have been needed. This might have functioned.
This might have. And this might have contributed.

So again, this is just a shorthand way of trying to convey that, when I read papers like this, I do not see any support for Darwin's theory. I read them as speculative and -- but nonetheless, people who already do believe in Darwin's theory fit them into their own framework.

25

Q. Now Dr. Miller cited numerous papers in his

testimony to support his claims on irreducible 1 2 complexity, the type III secretory system, and so forth. Have you done a review of those papers and have some 3 comments on them that you prepared slides for? 4 A. Yes, I did. I went through many of the papers 5 that Professor Miller cited, as many as I could, and 6 simply, as a shorthand way of trying to indicate or 7 8 trying to convey why I don't regard any of them as persuasive, I simply did a search for the phrases, 9 random mutation, which is abbreviated here in this 10 11 column, RM, and the phrase, natural selection. 12 Random mutation, of course, and natural selection are the two elements of the Darwinian mechanism. 13 That. 14 is what is at issue here. And so this is, you know, 15 this is, of course, a crude and perhaps shorthand way, but nonetheless, I think this illustrates why I do not 16 17 find any of these papers persuasive. 18 When I go through the papers that Professor Miller cited on the blood clotting cascade, Semba, et 19 20 al, Robinson, et al, Jiang and Doolittle, there are no 21 references to those phrases, random mutation and natural 22 selection.

Q. Some of your indications on this slide, you have of with asterisks and some without. Is there a reason for that?

A. Yes. The papers that have asterisks, I scanned 1 2 I read through them visually. Ones that do not by eye. have an asterisk, I was able to do a computer search for 3 those phrases because they are on the web or in computer 4 readable form. I have a number of other such tables. 5 On the next one are references that Professor 6 7 Miller cited on the immune system. And again, none of these references contain either those phrases, random 8 9 mutation and natural selection. There were a couple 10 more references on the immune system that Professor 11 Miller cited, and they didn't contain those phrases 12 either. 13 In references for the bacterial flagellum and the 14 type III secretory system, there was one paper by Hauch, 15 a review in 1998 that did use the phrase natural selection. However, that phrase did not occur in the 16 17 body of the paper. It was in the title of one of the 18 references that Hauck listed. 19 And on the next slide, I think there are papers 20 cited by Professor Miller on common descent of 21 hemoglobin. And again, those phrases are not there. Ι 22 think there's another slide or two, if I'm not mistaken. 23 This is the one on what he described as molecular trees, 24 Fitch and Margoliash, from 1967. And I didn't find the 25 phrase there either. So again, this is a shorthand way

of showing why I actually considered these off-the-point 1 2 and unpersuasive. Q. So all these papers that are being used to 3 provide evidence for Darwin's theory of evolution, in 4 5 particular, the mechanism evolution of natural selection, yet they don't mention random mutation or 6 7 natural selection in the body of the works? A. That's correct. 8 Could you summarize the point then, Dr. Behe, 9 Q. that you are making with, referring to these studies and 10 the comments you made about the speculative nature of 11 12 some of these studies? 13 A. Yes. Again, much of these studies, in my view, 14 are speculative. They assume a Darwinian framework. 15 They do not demonstrate it. And certainly, you know, certainly scientists should be free to speculate 16 17 whatever they want. You know, science usually starts 18 with speculation, but it can't end with speculation. 19 And a person or, and especially a student, should 20 be able to recognize and differentiate between 21 speculation and actual data that actually supports a 22 theory. Q. So it would be beneficial to point this sort of 23 24 feature that you just described, point that out to 25 students?

I very much think so. 1 Α. 2 MR. MUISE: Your Honor, we're going to be 3 moving again into another subject, and it appears to be close to the time for a break. 4 THE COURT: Yeah, why don't we take a break 5 at this point. I think that makes good sense. We'll 6 7 break for 20 minutes at this juncture, and we'll return 8 and pick up direct examination at that point. 9 (Whereupon, a recess was taken at 10:11 a.m. 10 and proceedings reconvened at 10:36 a.m.) 11 THE COURT: All right. Mr. Muise, you may 12 continue. 13 MR. MUISE: Thank you, Your Honor. 14 BY MR. MUISE: 15 Dr. Behe, Dr. Miller severely criticized Pandas 0. for its treatment of the topic of protein sequence 16 17 similarity. Do you agree with his assessment? 18 No, I don't. Α. 19 And I would ask you to explain why not? Q. 20 Α. On the next slide, we see one of Professor 21 Miller's slides, the first, I think, in his sequence 22 where he very severely criticized the book Of Pandas and 23 People for its treatment of the question of why similar 24 proteins in separate organisms have the differences in 25 their sequence that they do.

And on the next slide, this is again a slide from 1 2 Professor Miller. He reproduces a figure from Pandas 3 which shows -- it's hard to read on here -- that the difference in the number of amino acids of a protein 4 5 called cytochrome c, which is a small protein which is 6 involved in energy metabolism and which has about 100 7 amino acids in it, the difference between that protein which occurs in fish is about 13 percent. 8

9 About 13 amino acids differ between the fish cytochrome C and frog cytochrome C; and about 13 or so 10 between bird and fish cytochrome C; and about 13 between 11 12 mammalian cytochrome C and fish cytochrome C. So that 13 remarkably, the proteins in these different organisms 14 all seem to have roughly the same number of differences, 15 although the differences are not the same differences, 16 but they have the same number of differences from fish 17 cytochrome C.

18 And Pandas discusses this in their text. And 19 Professor Miller -- Professor Miller takes Pandas to 20 task because he says that, in fact, this is a 21 well-studied and a problem that has been solved by evolutionary theory. For example, he says, in fact, 22 23 these sequence differences confirm that each of these 24 organisms is equi-distant from a common ancestor, which is the actual prediction of evolutionary theory. 25

1	He has a little tree diagram there, too. But one
2	has to realize that, in fact, Professor Miller is
3	mistaken. Evolutionary theory does not predict that.
4	Or one could say, evolutionary theory predicts that in
5	the same sense that evolutionary theory predicted that
6	the vertebrate embryos, as drawn by Haeckle, should be
7	very, very similar to it; or the prediction of
8	evolutionary theory after newer results came out, that
9	vertebrate embryos could vary by quite a bit; or the
10	prediction of evolutionary theory that the type III
11	secretory system would be a good pre-cursor for the
12	flagellum; or the prediction of evolutionary theory that
13	the flagellum or that the type III secretory system
14	might be derived easily from a flagellum.
15	So, in fact, what we have, I will try to make
16	clear, is an instance where experimental science comes
17	up with data, and the data is attempted to be fit into a
18	framework. But this data was not predicted by any
19	evolutionary theory.
20	Q. How was Pandas' treatment of this compared with
21	what Dr. Miller found?
22	A. In my view, Pandas' treatment of this topic is
23	actually much more accurate than Professor Miller's
24	discussion of the same topic in his testimony here.
25	Professor Miller, in his discussion, where he says that,

1	evolutionary theory predicts this remarkable amount of
2	difference, is referring to something, although he does
3	not call it such, something called the molecular clock
4	hypothesis.
5	And notice that, in fact, in Pandas, on the page
6	opposite to the figure that Professor Miller used in his
7	presentation, there is a section entitled A Molecular
8	Clock where they go through and discuss some issues with
9	it, which I will talk about later on.
10	${\tt Q}$ . Just to be clear for the record, the diagram,
11	figure 9 that you've been referring to that Dr. Miller
12	cited in his testimony, appears on page 38 of Pandas, is
13	that correct?
14	A. Yes.
15	Q. And the discussion of the molecular clock
16	appearing on the subsequent page appears on page 39 of
17	Pandas, as indicated in this slide, is that correct?
18	A. That's correct.
19	${\tt Q}$ . Do you have some slides and discussion as to how
20	this molecular clock problem is treated in the science
21	community?
22	A. Yes, I do, and it will probably take about 10
23	minutes or so to go through it. So please be patient.
24	But here is a cover of the Biochemistry textbook that I
25	referred to frequently here by Voet and Voet, which is

1 used in many universities and colleges across the 2 country.

And they have a section on the molecular clock hypothesis and on cytochrome C in which they discuss these issues. Let's imagine -- I'm going to try to explain a molecular clock. Let's imagine that these lengths of time -- these lines represent time. And down at the bottom of the screen is a time -- a distant time g ago, and up at the top is modern time.

And the branches here represent events in the course of life where a population of organisms split into two -- split into two, and one branch went off to form one group of organisms and another group went off to form a different type of organisms.

Q. If I might just interrupt briefly. You're referring to a phylogenetic tree that has vertical lines that branch off to each other, and that's what you're referring to the vertical lines running, two at the top of the diagram, and then they branch off into different sections?

A. That's correct. That's exactly right.
Q. Could you continue, please?
A. Yes. So, for example, at this branch, a
population of organisms split off that went on to become
plants, and at this branch, a population split off which

1 went on to become animals.

2	Now I suppose that before any split in the
3	population, the pre-cursor population organisms had a
4	cytochrome c with a certain sequence. We'll say there
5	was a hundred letters. Just think of a string of a
6	hundred letters; Z, Q, A, L, W.
7	Now, however, when we get to this branch point,
8	we have a group of organisms going off to form the
9	animals, another going off to form the plants. They no
10	longer interbreed, and so that string of a hundred
11	letters representing cytochrome c can't accumulate
12	mutations in it separately.
13	So, for example, suppose once every year or so,
14	the cytochrome c in the branch that is forming the
15	plants suffered a mutation, so that one of those letters
16	changed from what it had been. And similarly, in the
17	branch going off to form the animals, once every hundred
18	years or so, one of those letters changed into
19	something.
20	Not necessarily the same. Maybe a different one.
21	So that after a while, those two sequences would be
22	different. And suppose every hundred years, that
23	happened, one change, one change, one change, and so on.
24	After a while, you'd start to accumulate a number of
25	changes.

Now further suppose that along the line to 1 2 animals, the population of animals split into two, one line leading to, say, insects, and another line leading 3 to mammals. Now you could have the same thing with the 4 5 cytochrome c sequence that had been mutating all along, 6 but now they split into two populations, and now these 7 two populations also begin to accumulate mutations independently. 8

9 But notice here, they start right at the branch 10 point with the same sequence. But after, say, a hundred 11 years, this will have one difference with what it had at 12 the beginning. This one will have one difference, too. 13 And they don't necessarily have to be the same 14 difference.

15 So they'll start to accumulate differences with 16 each other between, say, the branch leading to the 17 insects and the branch leading to the mammals. Now 18 here's the point. Any sequence along this branch should 19 have accumulated the same number of sequences between 20 any sequence on this branch.

21 So that the number of differences between insects 22 and plants should be roughly the same between, as that 23 between mammals and plants. Any animal and any plants 24 should have roughly the same number of differences. 25 Whereas between subgroups of animals that have split off

1	from each other earlier than animals did from plants,
2	they will have had less time to accumulate differences
3	in their amino acid sequences. And so they will have
4	so they will have fewer differences.
5	Q. You mean, if they split off later. You said,
6	earlier. They were split off later, correct?
7	A. Thank you. Yes, later. So Professor Miller has,
8	I believe, this sort of model in mind, which is commonly
9	which is a common way of thinking of these things in
10	science.
11	So the idea is that, since fish branched off from
12	those other groups of vertebrates, mammals, birds, and
13	so on, the fish, under this model, would be expected to
14	have the same number of differences in their amino acid
15	sequences between themselves and all those other
16	vertebrate groups.
17	Q. So here you have plants splitting off at the same
18	time as the insects or you have the same you have the
19	same connection between insects and plants as plants and
20	mammals?
21	A. That's right. So the critical point is that, the
22	difference between animals, any animal group like
23	mammals and plants and insects and plants, they should
24	have the same difference between animals and plants, no
25	matter what the subgroup of animals.

But between animals which branch off -- groups of animals which branched off at an earlier -- or from each other earlier to the current time, they would have less time to accumulate differences. And I believe this is what Professor Miller had in mind.

However, this model has some difficulties with it which are well recognized and have been discussed in the literature for over 40 years. For example, I said, suppose every hundred years or so, a mutation occurred. Okay. Well, suppose that in this branch, every hundred years or so, a mutation occurred. But in this branch, suppose a mutation occurred every 50 years.

And suppose when these split, the mutation rate again changed somewhat. Now you would not expect this nice, neat pattern to occur. Now you would expect a jumble. It's not quite clear what one might expect. And it turns out, that's a real problem because it's thought that most mutations accumulate in a lineage when an organism reproduces.

When an organism reproduces, the DNA in it has to be replicated, and that gives a chance for mutations to come into the DNA. But different organisms can reproduce at greatly differing rates. For example, a fruit fly might have a generation time of two weeks, and an elephant might have a generation time of 20 years. 1 So if the number of mutations that a protein or 2 gene underwent was proportional to the number of 3 generations, you might expect a lineage with quickly 4 reproducing organisms to accumulate mutations much more 5 quickly, and the one with slowly reproducing organisms 6 to accumulate more slowly.

7 And I believe this is -- on the next slide, there shows discussion from the Biochemistry textbook 8 9 explaining exactly that point. Let me quote from it. 10 Quote, Amino acid substitutions in a protein mostly 11 result from single base changes in the gene specifying 12 the protein. If such point mutations mainly occur as a 13 consequence of errors in the DNA duplication process, 14 then the rate at which a given protein accumulates 15 mutations would be constant with respect to numbers of cell generations. 16

Not with time. With numbers of cell generations.
If, however, the mutations process results from a random
chemical degradation of DNA, then the mutation rate
would be constant with absolute time. So here's this
complication. If most mutations occur during
replication, you wouldn't expect this difference that we
see in cytochrome c.

If, for some reason, mutations occurred constant with time, well, then you might expect that. But the

1	problem is, we know of no reason why that necessarily
2	that has to be so, why a mutations have to would have
3	to occur constant in time.
4	Q. Is there a problem in addition to this
5	generational rate change?
6	A. Yes, that's one complication, but there's another
7	one as well. And that's that, this so-called molecular
8	clock seems to tick at different rates in different
9	proteins. And this is an illustration again from the
10	Biochemistry textbook that applies to this point.
11	On the bottom, the X axis, this is time. This is
12	200 million, 400 million, a billion years, and so on.
13	This is number of or percent amino acid sequence
14	difference. And the idea is that, here's the line for
15	cytochrome c.
16	Organisms which diverge about 200 million years
17	ago have these many sequence differences; about 400
18	million years ago, have these many, and so on. Look at
19	how nice and neat that is. However, for another
20	protein, hemoglobin, the molecular clock seems to tick
21	faster. For the same amount of time, hemoglobin has
22	maybe twice as many mutations.
23	Another region of a protein called a
24	fibrinopeptide seems to accumulate mutations extremely
25	rapidly. And a fourth protein, if you can look at the

bottom of the figure, it's hard to see, for something called histone H4, barely accumulates any mutations at all. Organisms in very widely separated categories have virtually identical histone H4's.

5 Now to resolve this problem, it was postulated 6 that perhaps this has to do with the number of amino 7 acid residues in a protein that are critical for its 8 function. Perhaps in some proteins, you know, most of 9 the amino acid residues cannot be changed or it destroys 10 the function and would destroy the organism.

11 And in others, maybe some can be changed, but not 12 others. And so you can change those. And perhaps in another group, almost all of them can be changed without 13 14 really affecting the function. And so that's an interesting idea. But there are also difficulties with 15 that because, under that model, you would predict that 16 17 if you changed the amino acid sequence of histone H4, 18 then that should cause problems for an organism, because all of its, or most of its, or practically all of its 19 20 amino acids are critical for function. But 21 experimentally, that is not supported, as shown on the next slide. 22 23 Q. Is this -- so you've done work in this area with 24 the histone H4 and the molecular clock? 25 A. Yes, uh-huh. I've written this commentary in

1 1990 in a journal called Trends in Biochemical Sciences, 2 commenting on the work of somebody else who 3 experimentally took an organism called yeast into the 4 lab and altered its histone H4 and actually chopped off 5 a couple amino acids at the beginning portion of that 6 protein.

7 And when he looked, it seems that it didn't make 8 any difference to the organism. The organism grew just 9 as well without those mutations, which is surprising, 10 which is not what you would expect if all of those 11 residues were critical for the function of that protein, 12 histone H4.

Later on, in the year 1996, I and a student of mine, Sema Agarwal, we were interested in this problem of histone H4 and molecular clock, and so we experimentally altered some amino acid residues into protein and changed them into different amino acids, with the expectation that these might destroy the function of the protein. But it turned out not to.

These positions, these amino acids could be substituted just fine, which is unexpected, and which kind of complicates our interpretation of the molecular clock hypothesis. So there are two complications; complications upon complications.

25

One, we would expect the number of mutations to

accumulate with generation time, but it seems to
accumulate, for some unknown reason, with absolute time.
And the second is that, proteins accumulate mutations at
different rates. We would expect that it would have to
do with how vulnerable they are to mutations, and
mutations might destroy the function of one protein that
evolved slowly, but that is not experimentally
supported.
Q. Now has this problem been discussed in the
scientific literature?
A. Yes, this has been continuously discussed ever
since the idea of the molecular clock hypothesis was
first proposed in the early 1960's by two men named
Emile Zuckerkandl and Linus Pauling. And here are a
couple of papers which deal with the difficulties of the
molecular clock hypothesis.
Here's a recent one, Gillooly, et al, published
in the Proceedings in the National Academy of Sciences,
entitled The Rate of DNA Evolution, Effects of Body Size
and Temperature on the Molecular Clock. In this
publication, they say that, in fact, the size of an
organism and temperature can affect how fast or how slow
this clock might tick.
Francisco Ayala has written on this frequently.
Here's one from 1997. And I should say, Francisco Ayala

is a very prominent evolutionary biologist. He wrote an
 article in 1997 entitled Vagaries of the Molecular
 Clock. And I think the title gets across the idea that
 there are questions with this hypothesis.

And in 1993, a researcher named Tomoka Ohta published an article in the Proceedings of the National Academy of Sciences entitled An Examination of the Generation-time Effect on Molecular Evolution in which she considers exactly that complication that the textbook Voet and Voet pointed out, this generation-time effect.

You know, why shouldn't organisms that reproduce more quickly accumulate more mutations. I have another slide just from one more recent paper. This paper by Drummond, et al, is entitled Why Highly Expressed Proteins Evolve Slowly. And it's referring to the sequence evolution that I've been discussing.

It was published in the Proceedings of the National Academy of Sciences, and this was from an online version. This is so recent that I don't think it has yet appeared in print. The point I want to make with this is that, these people treat this question as a currently live question.

They start off by saying, a central problem in molecular evolution is why proteins evolve at different

rates. So that question I was trying to illustrate with 1 2 histone H4, why does one protein tick faster and another one tick more slowly, that's still -- that is still 3 unknown. 4 And I think I will skip the rest of this slide 5 and go to the next slide and just point out a couple 6 7 words here. Drummond, et al, say, Surprisingly, the best indicator of a protein's relative evolutionary rate 8 is the expression level of the encoding gene. 9 The only point I want to make with this is that, 10 11 they are reporting what is a surprise, what was not 12 expected, which was not known, you know, 40 years ago, which has only been seen relatively recently. And they 13 14 say, quote, We introduce a previously unexplored hypothesis, close quote. 15 And the point I want to emphasize is that, here 16 17 in this paper published, you know, weeks ago, that they 18 are exploring new hypotheses to try to understand why proteins have the sequences that they do. 19 20 Q. So in summary, this protein sequence, the fact 21 that the equi-distant from a common ancestor is not what 22 evolutionary theory would actually predict? 23 Α. That's right. Evolutionary theory makes no firm 24 prediction about this anymore than it makes a firm prediction about the structure of vertebrate embryos. 25

It's a common understood problem that biologists 1 Q. 2 are trying to resolve at this point? A. Yes, within the community of scientists who work 3 on this. People have been working on it for decades. 4 5 Is this a problem that an American Biology 0. teacher should be aware of? 6 7 A. Yes, an American Biology teacher should be aware of it, because an article on this very topic was 8 9 published in the magazine, American Biology Teacher, a couple years ago, which is put out by the National 10 11 Association of Biology Teachers. 12 And the article is entitled Current Status of the 13 Molecular Clock Hypothesis. And one of the first --14 this is a red arrow that I added to the figure. One of the first subsections of the article is entitled How 15 Valid is the Molecular Clock Hypothesis? And if you'll 16 17 advance to the next slide, let me just read the last 18 line from the paper. The author says, The validity of a molecular 19 20 clock, except in closely related species, still remains 21 controversial. So the point is that, extrapolating 22 across wide biological distances, such as from fish to 23 other vertebrates, that is controversial. 2.4 Maybe similar species, species of mice or some 25 such thing, okay. But when you try to extrapolate

further, the model is guite controversial. 1 2 How does Pandas then address this issue? 0. Well, I have here the section from Pandas 3 Α. entitled The Molecular Clock where they discuss exactly 4 5 all these things. They discuss the molecular clock, the 6 standard molecular clock model, the naive molecular 7 clock model, and then they discuss complications with it. 8

9 Let me just read this section from Pandas on the molecular clock. They write, quote, Some scientists 10 have suggested that the idea of a molecular clock solves 11 12 the mystery. The explanation they advance is that there is a uniform rate of mutation over time, so quite 13 14 naturally, species that branched off from a common 15 ancestor at the same time in the past will now have the same degree of divergence in their molecular sequences. 16

There are some serious shortcomings, however, with this explanation. First, mutation rates are thought to relate to generation times, with the mutation rates for various molecules being the same for each generation.

The problem comes when one compares two species of the same taxon, say two mammals, with very different generation times. Mice, for instance, go through four to five reproductive cycles a year. The number of mutations, therefore, would be dramatically higher than,
 say, those of an elephant.

Thus, they should not reflect similar percent sequence divergences for comparable proteins. Besides that, the rates of mutations are different for different proteins even of the same species. That means that, for the molecular clock idea to be correct, there must be not one molecular clock, but thousands.

9 So let me point out here that, in this section, 10 Pandas describes the simple molecular clock idea that was proposed 40 years ago by Zuckerkandl and Pauling, 11 12 and then talks about the two complications for the model, which are common knowledge and are taught in 13 14 basic science texts that deal with this issue, the generation time problem and the fact that different 15 proteins accumulate mutations at different rates. 16

And as I have shown from the literature I just cited, that continue to be live issues in the scientific community.

Q. In that section you read from on the molecular clock from Pandas are found on page 39, is that correct? A. Yes, that's correct.

Q. Again, returning to that slide that Dr. Miller
presented in his testimony?

25

A. Yes. I just wanted to go back to that slide

where Dr. Miller says -- again, I should say that, in his testimony, which I attended, he, you know, excoriated Pandas on this point. And he says -- on his slide, he says, in fact, the information we have confirms that each of these organisms is equidistant from a common ancestor, which is the actual prediction of evolutionary theory.

And that's simply is incorrect. And in my view, 8 9 Pandas is treating problems that Professor Miller, treating real live problems that Professor Miller shows 10 11 no signs of being aware of. So I think a student 12 reading this section would actually get a better appreciation for this subject than otherwise. 13 14 Q. Dr. Behe, in Dr. Miller's testimony, he also 15 criticized another example found in Pandas that had a message such as, quote, John loves Mary, written on the 16 17 beach, would be a sure sign of intelligence.

18 He claimed that any philosopher, any logician would spot the mistake in logic, because we know a human 19 20 made that message, and probably made it with a stick, 21 because we have seen such things happen in our own 22 experience. Do you agree with this reasoning? 23 No, I disagree with Professor Miller's reasoning. Α. 24 Q. And if I can just say, the example that John 25 loves Mary, and we have a slide up, that's on page 7 of

1 Pandas, correct?

25

2 A. Yes, that's right.

3 Q. Again, could you explain why you disagree with 4 this reasoning?

The inference from the -- the inference 5 Α. Yes. from the existence of designed objects in the -- in our 6 7 world of experience to the conclusion of design in life is an example of an inductive inference. And I think I 8 9 explained earlier that, in an inductive inference, one 10 always infers from examples of what we know to examples 11 of what we don't know.

12 And the strength of the inference depends on 13 similarities between the, between the inference in 14 relevant properties. For example, in the Big Bang 15 hypothesis, scientists extrapolated, or used inductive 16 reasoning of their knowledge of explosions from our 17 everyday world from things like fireworks and canon 18 balls and so on.

19 They extrapolated from their experience that the 20 motion of objects away from each other bespeaks an 21 explosion. They extrapolated from our common everyday 22 experience to something that nobody had ever seen 23 before, an entirely new idea, that the universe itself 24 began in something like a giant explosion.

Nonetheless, they were confident that this was a

1 good idea because they thought the relevant property,
2 the parts moving rapidly away from each other, was what
3 we understand from an explosion. And that's how science
4 often reasons.

5 In the same way, the purposeful arrangement of parts in our everyday experience bespeaks design. 6 7 Pandas is exactly right, that if we saw such a message on the beach, we could conclude that it had been 8 designed. And William Paley is exactly right, that if 9 we stumbled across a watch in a field, that we would 10 conclude that it was designed, because in each case 11 12 there is this strong appearance of design from the 13 purposeful arrangement of parts.

14 Now we have found purposeful arrangement of parts 15 in an area where we didn't expect to, in the very cellular and molecular foundation of life, in the cell. 16 17 The cell again was not understood in Darwin's day. And 18 it is much better understood now. And from the new information we have, again, we see this purposeful 19 20 arrangement of parts, and it's -- by inductive 21 reasoning, we can apply our knowledge of what we see in 22 our everyday world to a different, completely different 23 realm.

And so that sort of inference has been done in science throughout the history of science, and it's a

1	completely valid inference for Pandas to make.
2	Q. Now we've heard some testimony throughout the
3	course of this trial of a program called SETI, S-E-T-I,
4	a project, I believe, that stands for the search for
5	extraterrestrial intelligence?
6	A. Yes.
7	Q. Are you familiar with that project?
8	A. Yes, I am.
9	Q. Whose project is that?
10	A. The search for extraterrestrial intelligence is a
11	project that was, for a while, was sponsored by the
12	federal government. It involved scientists scanning the
13	skies with detectors to see if they could detect some
14	electromagnetic signal that might point to intelligence.
15	Q. Is there a comparison with that project to the
16	discussion you had in here with the John loves Mary on
17	the beach?
18	A. Yes. Again, if they detected something that
19	seemed to have a purposeful arrangement of parts, if
20	they saw something that bespoke a message, then even
21	though we have had no experience with other entities
22	from off the Earth trying to send us a message,
23	nonetheless, we could still be confident that an
24	intelligent agent had designed such a message.
25	And again, whenever we see John things like

John loves Mary, we can be confident of that. And when 1 2 we see the purposeful arrangement of parts in the cell, the argument is that, we can be confident of that, that 3 4 that bespeaks design as well. I want to bring this discussion somewhat down to 5 0. the molecular level, and ask you whether or not new 6 7 genetic information can be generated by Darwinian 8 processes. And I want to be more specific and ask 9 whether new genetic information can be generated by known processes such as gene duplication and exon 10 11 shuffling? 12 A. Well, that's a topic about which you have to be very careful and make distinctions. 13 14 Q. Okay. Let's start with the gene duplication. Ιf 15 you could explain what that is in the context of generating new genetic information? 16 17 A. Well, gene duplication is a process whereby a 18 segment of DNA gets copied twice or gets duplicated and replicated so that where one gene was present before, a 19 20 second copy of the exact same gene is now present in the 21 genome of an organism. Or sometimes larger segments can 22 be duplicated, so you can have multiple copies of 23 multiple genes. 2.4 Q. Are you saying, duplication, like photocopying, is just making another copy of the gene that was 25
1 originally existing?

2	A. Yeah, that's a good point. It's important to be
3	aware that gene duplication means that you simply have a
4	copy of the old gene. You have not done anything new.
5	You've just taken the same gene and copied it twice. So
6	it would be like, like photocopying a page. And now you
7	have two pages, but it's just a copy of the first one,
8	it's not something fundamentally new.
9	It would be like saying, the example of Pandas
10	here with John loves Mary. If you walked down the sand
11	another five yards or something, and you came across
12	another message that says, John loves Mary, well, that's
13	interesting, but you don't have anything fundamentally
14	new.
15	Q. Can there be variations though in the duplication
16	of those genes?
17	A. Well, once a gene has duplicated, then the idea
18	goes that, perhaps one of those two copies can continue
19	to perform the function that the single copy gene
20	performed before the duplication, and the other one is
21	sort of a spare copy.
22	Now it's available to perhaps undergo mutation,
23	and mutation accumulate changes, and perhaps Darwinian
24	theory postulates. Perhaps it can go on to develop
25	brand new properties.

Q. Does this generate new information? And if you 1 2 use that John loves Mary example to help explain 3 perhaps? A. Well, again, you have to be careful. Nobody 4 disputes that random mutation and natural selection can 5 6 do some things, can make some small changes in 7 pre-existing systems. The dispute is over whether that explains large complex functional systems. 8 9 And to leave the world of proteins for a second, to look at John loves Mary, suppose we're looking at the 10 spare copy, and the first copy was continuing to fulfill 11 12 the function of conveying that information. Well, you know, suppose you changed a letter. Suppose you changed 13 14 the final n in the word John to some other, some other letter, like r. That would not spell a name in the 15 English language. 16 17 So that's kind of an analogy to saying that, you 18 might lose the function of the message in the terms. Ιn

19 the terms of protein, the protein might no longer be 20 functional. But you might get to closeby. You might 21 get to closeby messages. For example, if you deleted 22 the r and the y from the end of Mary, you might get to 23 John loves Ma, or some such thing. But you're not going 24 to get anything radically different from that.

Q. So you are operating with the copy. The copy is

1	operating with those same letters, the John loves Mary,
2	or some variation or deletions of that subset?
3	A. That's right. A copy is a copy. It's
4	essentially the same thing. And now the big problem
5	that Darwinian processes face is, now what do you do?
6	How do you generate a new complex function?
7	Q. And that's with gene duplication that we just
8	talked about. Could you explain a little bit about exon
9	shuffling in the context of generating new complex
10	information?
11	A. Yes, exon shuffling is a little bit more
12	involved. It turns out that the gene for a protein can
13	contain regions of DNA that actually code for regions of
14	a protein interrupted by regions of DNA that don't code
15	for regions of a protein. And the regions that code for
16	the part of the protein are called exons.
17	Now it turns out that, in cellular processes,
18	similar to gene duplication and other processes, too,
19	one can duplicate separate exons and sometimes transfer
20	them to different places in the genome and other such
21	processes. But to make it more understandable, we can
22	go back to the analogy of John loves Mary.
23	And in this sense, exon shuffling might be
24	expected to generate something like, instead of John
25	loves Mary, perhaps Mary loves John, or John Mary loves,

1	
1	
Ţ	or something like that. But again, it's kind of a
2	mixture of pre-existing properties, and we're not
3	generatesing something fundamentally new.
4	Q. So, for example, you couldn't generate Brad loves
5	Jen from exon shuffling using your beach example?
6	A. No, I hope not.
7	Q. Do these concepts, particularly gene duplication,
8	exon shuffling, do they have any impact on the concept
9	of irreducible complexity that you've been discussing
10	quite a bit throughout your testimony?
11	A. Yes. In fact, there is an important point to
12	recognize here. Russell Doolittle knew all about the
13	processes of gene duplication and exon shuffling. And
14	as a matter of fact, in the blood clotting cascade, many
15	proteins look similar to each other, and they're often
16	times pointed to as examples of exon shuffling.
17	But nonetheless, that knowledge did not allow him
18	to explain how the blood clotting system might have
19	arisen. Again, these are sequence comparisons. And
20	such information simply does not speak to the question
21	of random mutation and natural selection being able to
22	build complex new biochemical structures.
23	In the same way, the people who are investigating
24	the type III secretory system and the bacterial
25	flagellum know all about gene duplication and exon

1 shuffling. And nonetheless, that information has not 2 allowed them to explain the origin of either of those 3 structures.

So those are interesting processes. And people 4 who are convinced of Darwinian theory include those 5 6 processes in their theory, but they do not explain --7 they do not explain where new complex systems come from. And it's an example of somebody accommodating this 8 9 information to an existing theory rather than getting information that actually experimentally supports the 10 11 theory.

12 Q. So can random mutation and natural selection 13 generate new information?

14 Well, again, that's -- you have to be careful. Α. 15 You can make small changes in pre-existing systems. And that's clearly the case. One can clearly do that. 16 But there has been no demonstration to show that such 17 18 processes can give rise to new complex systems such as we've been suggesting. And there are many reasons to 19 20 think that it would be extremely difficult to do so. 21 Q. Have you prepared some slides with a couple --22 several quotes that make this point? 23 A. Yes, I do. This first one is an excerpt from a 24 paper from John Maynard Smith, which I spoke about 25 earlier, from 1970 entitled Natural Selection and the

Concept of a Protein Space. Let me read the first 1 2 excerpt. Quote, It follows that if evolution by natural 3 selection is to occur, functional proteins must form a 4 continuous network which can be traversed by unit 5 6 mutational steps without passing through nonfunctional 7 intermediates, close quote. Again, let me explain. If you can remember the figure of two proteins 8 9 binding to each other that I showed in -- I showed yesterday, he is speaking of unit mutational steps in 10 11 terms of one of those interactions, maybe a plus charge and a minus charge or a hydrophobic group and another 12 13 hydrophobic group. 14 And so to get two proteins to -- or proteins to 15 start change into something new and different with different properties, each one of those changes would 16 17 have to be a beneficial one, or at least not cause any 18 difficulties for the problem. And actually, seeing how that could happen is extremely difficult. 19 20 And continuing on this slide. I'm sorry. Could 21 you back up one slide? Thank you. The bottom part of 22 the quotation, he says, quote, An increase in the number 23 of different genes in a single organism presumably 24 occurs by the duplication of an already existing gene 25 followed by divergency. So here, he's kind of

describing the standard scenario which -- scenario,
which is standard in Darwinian thinking, that one has
gene duplication and then divergence of the sequence of
a gene, and that gives a brand new interesting and
complex protein.

6 But notice that I, of course, underlined and 7 bolded the word presumably. Well, presumably, you know, 8 is a presumption. And it may be true, and it may not. 9 But presumptions are not evidence. And so in order to 10 support this idea, one needs more than the presumption 11 that it occurs.

Q. Do you have another citation to a science text?
A. Yes, I do. Here's an excerpt from an article by
a man named Alan Orr, who is an evolutionary biologist
at the University of Rochester. And again, this speaks
to the same consideration, that you have to be able to
have a pathway that step by tiny step could lead from
one functional protein to another.

He says, quote, Given realistically low mutation rates, double mutants will be so rare that adaptation is essentially constrained to surveying, and substituting, one mutational step neighbors. Thus, if a double mutant sequence is favorable, but all single amino acid mutants are deleterious, adaptation will generally not proceed. Again, this makes the point that, if you only

need to change one little step, Darwinian evolution 1 2 works fine. But if you need to change two things before you get to an improved function, the probability of 3 Darwinian processes drops off dramatically. 4 If you need three things, it drops off, you know, 5 even more dramatically. And nonetheless, as I showed in 6 that figure of interacting proteins, even to get two 7 proteins to stick together, multiple groups are 8 9 involved. Did you write about something similar in a paper? 10 Ο. 11 Yes. The paper that I published with David Snoke Α. 12 last year speaks exactly to this topic. It's entitled Simulating Evidence by Gene Duplication of Protein 13 14 Features that Require Multiple Amino Acid Residues. 15 And in this theoretical study, we showed that, again, if you need one change, that's certainly doable. 16 17 If you need two amino acid changes before you get a 18 selectable function, the likelihood of that drops 19 considerably. Three or more, now you're really in the 20 very, very improbably range. So again, gene duplication 21 is not the answer that it's often touted to be. 22 Can you make an analogy here at all to -- you Q. 23 talked about Maxwell and the ether theory? A. Yes. When Darwinian -- adherence to Darwinian 24 25 theory, when they view that there are similar genes in

1 different -- in the same organism, and they infer a
2 process of gene duplication, it is simply their
3 theoretical framework, which is saying, such a process
4 must be important in generating new and complex
5 structures.

6 That has not been demonstrated. Just like James 7 Clerk Maxwell knew that light was a wave and inferred 8 from his theory that there must be an ether, modern 9 Darwinists infer from something we know, the existence 10 of gene copies to an unproved role of such a process in 11 generating complex biochemical systems.

Q. Now Dr. Miller says that Pandas necessarily rejects common descent, and points to a figure -- I believe it was 4.4 on page 99 -- showing separate lines representing categories of animals rather than a branching tree. Do you regard that as ruling out common descent?

A. No, I don't. And here's a figure that I made up in the upper right-hand corner. It's figure 4.4 from Pandas, which is the figure that Professor Miller showed, which shows straight lines instead of a branching tree, which is the traditional representation of how -- of the fossil record.

Nonetheless, here I regard this as simply tryingto describe the data without a theoretical framework,

without the branched lines in between. One has to
 realize that these lines do not occur in the fossil
 record. These are theoretical constructs.

And how one groups things together is theory building rather than data itself. I viewed this as Pandas trying to describe the data without the framework of the existing theory. And I might add that, this was figure 4.4. And earlier, a couple pages earlier, Pandas describes the traditional interpretation of the fossil record in terms of a branching tree.

And in this section, section 96 through 100, the 11 12 meaning of gaps in the fossil record, Pandas describes the traditional tree diagram for the fossil record, and 13 14 then points to statements by biologists, saying that there seem to be difficulties in this sort of 15 representation, and then goes on to discuss what 16 17 interpretations, what ideas have been offered to try to 18 account for the form of the fossil record.

19 Pandas writes, Several interpretations have been 20 offered to resolve this problem. That is, that the tree of life doesn't seem to be as continuous as one might 21 expect. Number 1, they say, imperfect record. 22 That is, 23 maybe not all organisms left representative of 24 fossilized specimens. Number 2, incomplete search. And 25 that is, maybe we simply haven't looked in the right

1 places or looked in all the places on the Earth, and 2 maybe when we do, then we will find what we expect to be 3 there.

Number 3, what they call jerky process, or which 4 5 has been called punctuated equilibrium, which was an 6 idea advanced by Steven J. Gould and Niles Eldredge in 7 the 1970's, whereby it said that the mode or the tempo of evolution is one in which a species or a branch of 8 9 life stays pretty much constant for a long period of 10 time, and then within a relatively short period of time, 11 large changes occur.

And then fourth, they say, well, perhaps -- they suggest something called the sudden appearance or face value interpretation, saying that, well, maybe if we see the sudden appearance of some feature or organism in the fossil record, then that, in fact, might be what happened.

18 Nonetheless, as I say, they discuss all of these possibilities, including the standard interpretation. 19 20 And at the end of the section, they write that, 21 scientists should not accept the face value 22 interpretation of the fossil record without also 23 exploring the other possibilities, and even then, only 2.4 if the evidence continues to support it. 25 So as I read this, Pandas is telling students

1	that they should follow the data where the data lead.
2	And if the data lead from this model to another model,
3	or from that model to a second model, then a scientific
4	attitude toward the problem is to follow the data, where
5	the data go.
6	Q. Dr. Behe, does intelligent design necessarily
7	rule out common descent?
8	A. No, it certainly does not.
9	Q. Now we've heard testimony from several witnesses
10	claiming that the theory of evolution is no different
11	than, say, the germ theory of disease, so there's no
12	reason to pay any special attention to it. Do you agree
13	with that?
14	A. No, I disagree.
15	Q. And why?
16	A. Well, in a number of ways, evolutionary theory is
17	unique. It's been my experience that students have a
18	number of misconceptions about the theory. They confuse
19	facts with theoretical interpretations. They do not
20	make distinctions between the components of evolutionary
21	theory.
22	And perhaps, most strikingly, a number of people
23	have made very strong extra-scientific claims for the
24	implications of evolutionary theory.
25	Q. Now I just want to return to something you had

1	said about your experience with students. You testified
2	that you teach a course called popular arguments on
3	evolution, is that correct?
4	A. Yes, that's right.
5	Q. And you've been teaching that for 12 years?
6	A. Roughly, yes.
7	Q. Now are there some standard misconceptions that
8	you can point to about the theory of evolution that you
9	find your students bringing to the class?
10	A. Yes. In my experience, a number of students come
11	in thinking that, in fact, evolution is completely true;
12	that is, they don't make a distinction between fact and
13	theory, they don't think it will be falsified, or they
14	don't think there's a possibility of it being falsified.
15	They also confuse various components of
16	evolutionary theory. For example, you can ask a
17	student, you know, why they think Darwinian evolution is
18	correct? And they'll say, you know, because, you know,
19	because of the dinosaurs. And they're mistaking change
20	over time with the question of natural selection. And
21	they will assume that the existence of animals in the
22	past necessarily means that animals in the present were
23	derived from them by random mutation and natural
24	selection.
25	Oftentimes also, students think that utterly

unsolved problems, such as the origin of life, have, in 1 2 fact, been solved by science. I had students tell me that, gee, it's true, right, that science has shown 3 genes being produced in origin of life experiments. 4 So in my experience, students bring a number of 5 6 misconceptions to this issue. 7 Q. One of the first ones you indicated is that they believe that Darwin's theory of evolution is a fact as 8 9 opposed to a scientific theory? 10 That's right. Α. 11 Ο. Does intelligent design seek to address some of 12 these misconceptions? 13 A. Yes. Yes, it does. One way is -- one way to 14 address the problem of students not understanding that 15 the distinction between fact and theory is to at least have at least one more theoretical framework in which to 16 treat facts. 17 18 If a student has only one theory and a group of facts to think of, it's extremely difficult to 19 20 distinguish what is theory and what is fact. The little 21 lines connecting various points on, say, a protein 22 sequence comparison are theory, but students can often 23 confuse them, confuse them to be facts. 2.4 Q. Do you believe these students will be better 25 prepared if they had learned that Darwin's theory of

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1	evolution was not a fact and that gaps and problems
2	existed within this theory?
ر ۲	$\lambda$ Vos I cortainly do They would see that in
5	A. Tes, I certainly do. They would see that, in
4	fact, if you can look at the data in a couple ways, then
5	they'll more easily distinguish data from interpretation
6	or from theory. And if they are aware that there are
7	problems in a theory, then perhaps they won't expect
8	they won't, again, confuse it with a fact, they'll
9	understand that there are some problems that are
10	unresolved.
11	Q. Now you made some indication previously in your
12	answer to my question that there are claims made about
13	the theory that go beyond biology, is that true?
14	A. Yes, that's certainly true.
15	Q. And do you have some slides to demonstrate some
16	of those examples?
17	A. Yes, I have a couple of slides, four slides over
18	that point to this. For example, in the high school
19	textbook Biology, which was written by Professor Kenneth
20	Miller and his co-author, Joseph Levine, this is the
21	1995 version, I think, the third edition, in a section
22	entitled The Significance of Evolutionary Theory, the
23	authors write, quote, The influence of evolutionary
24	thought extends far beyond biology. Philosopher J.
25	Collins has written that, quote, there are no living

1 sciences, human attitudes, or institutional powers that 2 remain unaffected by the ideas released by Darwin's 3 work, close quote.

In another example of the implications, the 4 5 profound implications beyond biology that some people 6 see for Darwin's theory, there's a section in his book, 7 Finding Darwin's God, A Scientist's Search for Common Ground Between God and Evolution, where Dr. Miller 8 9 writes that, quote, God made the world today contingent upon the events of the past. He made our choices 10 11 matter, our actions genuine, our lives important. Ιn 12 the final analysis, He used evolution as the tool to set us free. 13

14 So here is a scientific theory which is being 15 used to support the idea that we are free, we are free, in apparently some metaphysical sense, because of the 16 17 work of Darwin. In another example -- it's just that --18 for example, the expert, Professor John Hauck, the 19 theologian from Georgetown University, has written a 20 number of books, including God After Darwin, a Theology 21 of Evolution.

Further example, in -- the evolutionary biologist, Richard Dawkins, in his book, The Blind Watchmaker, writes, Darwin made it possible to be an intellectually-fulfilled atheist.

If I could have the next slide. Thank you. 1 The 2 Darwinian philosopher, Daniel Dennett, who's at Tufts University, has described Darwinism as a universal acid 3 4 that destroys our most cherished beliefs. And he says, 5 quote, Darwin's idea had been born as an answer to 6 questions in biology, but it threatened to leak out, 7 offering answers, welcome or not, to questions in 8 cosmology, going in one direction, and psychology, going 9 in the other direction.

10 If the cause of design in biology could be a 11 mindless, algorithmic process of evolution, why couldn't 12 that whole process itself be the whole product of 13 evolution, and so forth, all the way down? And if 14 mindless evolution could account for the breathtakingly 15 clever artifacts of the biosphere, how could the products of our own real, quote, unquote, minds be 16 17 exempt from an evolutionary explanation? Darwin's idea 18 thus also threatened to spread all the way up, dissolving the illusion of our own authorship, our own 19 20 divine spark of creativity and understanding.

21 So again, Professor Dennett sees implications for 22 Darwin's theory that are profound and that extend well 23 beyond biology. Another philosopher by the name of Alex 24 Rosenberg, who's at Duke University, published an 25 article a few years ago in the journal Biology and

Philosophy that, quote, No one has expressed the destructive power of Darwinian theory more effectively than Daniel Dennett. Others have recognized that the theory of evolution offers us a universal acid, but Dennett, bless his heart, coined the term.

In short, it, that is Darwin's idea, has made Darwinians into metaphysical Nihilists denying that there is any meaning or purpose to the universe, close quote. So again, a number of philosophers, a number of scientists, and so on, see very, very profound implications in Darwin's theory.

12 Two more quotations on this last slide on this 13 topic. Larry Arnhart is a professor of political 14 science at Northern Illinois University. He wrote a 15 book entitled Darwinian Natural Right, The Biological Ethics of Human Nature. And in it, he writes -- and in 16 17 it, he writes the following, that, quote, Darwinian 18 biology sustains conservative social thought by showing how the human capacity for spontaneous order arises from 19 20 social instincts and a moral sense shaped by natural 21 selection in human evolutionary history.

So let me emphasize that he sees implications for politics from Darwin's theory. And the same -- and a Princeton University philosopher by the name of Peter Singer has written a book entitled A Darwinian Left,

Politics, Evolution, and Cooperation. And in it, he 1 2 writes that we should try to incorporate a Darwinian ethic of cooperation into our political thought. 3 So the gist of Professor Singer's book is that, 4 5 Darwinian ideas support a liberal political outlook. 6 And he argues for that. So, again, these -- all of 7 these people see profound implications for Darwin's theory well far beyond biology. 8 9 Ο. These are non-scientific claims, correct? Yes, that's correct. 10 Α. 11 Have you come across any similar claims made 0. 12 about, say, the germ theory of disease? I have never seen the germ theory of disease 13 Α. 14 argued to say how we should conduct our political life. How about atomic theory? 15 Ο. I have never seen atomic theory used in such 16 Α. 17 profound senses either. So my point then is that, it is 18 perfectly rationale to treat a scientific theory, which so many people have claimed such profound implications 19 20 for, to treat it differently from other scientific 21 theories for which such far-reaching implications have not been claimed. 22 23 It might be very important, and I think a school 24 district would be very justified to say that, since this 25 particular theory seems to reach far beyond its

providence, then we should take particular care in explaining to our students exactly what the data is for this theory, exactly what is the difference between theory and fact, exactly what is the difference between theory and interpretation. And so I think such an action would be justified.

Q. Sir, I want to ask you some questions about creationism as it relates to intelligent design. First of all, let me ask you, does creationism have a popular meaning or is there a popular understanding of that term?

A. Well, again, you have to be careful, because many words in these discussions can have multiple meanings. And if you're not very careful about your definitions, you'll easily become confused.

Creationism -- creationist has sometimes been used, as John Maddox, the editor of Nature, used it, simply to mean somebody who thinks that nature was begun by a supernatural act, by God, and the laws of nature perhaps were made of God, and unfolded from there nonetheless.

Q. That would be similar to Dr. Miller's view towards evolution that he had written in his book Finding Darwin's God?

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A. Yes, that seems to be consistent with what he

1	wrote. But nonetheless, in the popular useage,
2	creationism means creationist means somebody who
3	adheres to the literal interpretation of the first
4	several books or first several chapters of the Book
5	of Genesis in the Bible, somebody who thinks that the
6	Earth is relatively young, on the order of, say, 10,000
7	years, that the major groups of plants and animals and
8	organisms were created ex-nihilo in a supernatural acts
9	by a supernatural being, God, that there was a large
10	worldwide flood which is responsible for major features
11	of geology, and so on.
12	Q. Now we've heard different terms; young-earth
13	creationism, old-earth creationism, and special
14	creationism. And you have familiarity with those terms,
15	is that correct?
16	A. Yes, that's right.
17	Q. Is intelligent design creationism, whether you
18	call it young-earth creationism, old-earth creationism,
19	or special creationism?
20	A. No, it is not.
21	Q. And why not?
22	A. Creation creationism is a theological concept,
23	but intelligent design is a scientific theory which
24	relies exclusively on the observable, physical,
25	empirical evidence of nature plus logical inferences.

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1	It is a scientific idea.
2	Q. Is it special creationism?
3	A. No, it is not special creationism.
4	Q. Again, why not?
5	A. Again, for the same reason. Creation is a
6	theological religious concept. And intelligent design
7	is a scientific idea, which is based exclusively on the
8	physical, observable evidence plus logical processes.
9	${\tt Q}$ . Dr. Miller has made a claim that if the bacterial
10	flagellum, for example, was designed, then it had to be
11	created, and is, therefore, special creationism. Is
12	that accurate?
13	A. No, that is inaccurate. The reason it's
14	again, creation is a theological concept. It is a
15	religious concept. But intelligent design is a
16	completely scientific concept which supports itself by
17	pointing to observable, physical, empirical facts about
18	the world, about life, and makes logical inferences from
19	them.
20	${\tt Q}$ . Does intelligent design require that the
21	bacterial flagellum, for example, instantaneously appear
22	from nothing?
23	A. No, it does not.
24	Q. Why not?
25	A. Because intelligent design focuses exclusively on

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1	the deduction of design from the purposeful arrangement
2	of parts. And it says nothing directly about how the
3	design was effected, whether it was done quickly, or
4	slowly, or whatever. So it has nothing to say about
5	that.
6	Q. Could the bacterial flagellum have been designed
7	over time?
8	A. Yes, it could.
9	Q. Does intelligent design require ex-nihilo
10	creation?
11	A. No, it does not.
12	Q. Why not?
13	A. Because again, the term ex-nihilo creation is a
14	theological concept, a religious concept. And
15	intelligent design is a scientific idea that relies on
16	observable facts about nature plus logical inferences.
17	Q. Is there, again, an analogy you can make here to
18	the Big Bang theory?
19	A. Yes. Yes, there is. Again, many people,
20	including many scientists, saw in the Big Bang theory
21	something that had theological implications, maybe this,
22	this Big Bang was ex-nihilo creation by a supernatural
23	being. And many people who saw that didn't like that.
24	Nonetheless, the Big Bang theory itself is an utterly
25	scientific theory because it relies on observations,

1	physical observations, empirical observations about
2	nature, and reasons from those observations using
3	logical processes.
4	Q. Is intelligent design a religious belief?
5	A. No, it isn't.
6	Q. Why not?
7	A. Intelligent design requires no tenet of any
8	particular religion, no tenet of any general religion.
9	It does not rely on religious texts. It does not rely
10	on messages from religious leaders or any such thing.
11	The exclusive concern of intelligent design is to
12	examine the empirical and observable data of nature and
13	reason from that using logical processes.
14	${\tt Q}$ . Now some claim that intelligent design advances a
15	religious belief, that it is inherently religious and
16	not science. Do you agree?
17	A. No. Again, no more than the Big Bang theory is
18	inherently religious. Although the Big Bang theory and
19	intelligent design might be taken by some people to have
20	theological or philosophical implications, both of them
21	rely on observed evidence, empirical evidence, and
22	logical reasoning.
23	Neither the Big Bang nor intelligent design
24	relies on any religious tenet, points to any religious
25	books, or any such thing.

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1	O De greationiste in the conce that Disintiffe and
T	Q. Do creationists in the sense that Plaintills and,
2	I believe, their experts use in this case require
3	physical evidence to draw their conclusions?
4	A. No. Actually, it's interesting that one could be
5	a creationist without any physical evidence. One could
6	rely a creationist could rely for his belief in
7	creation on, say, some religious text or in some private
8	religious revelation or some other non-scientific
9	source.
10	So a creationist does not need any physical
11	evidence of the kind that, for example, Richard Dawkins
12	sees in life that leads him to think that life has the
13	strong appearance of design or the kind that David
14	DeRosier sees in the bacterial flagellum. A creationist
15	can believe in creation without any such physical
16	evidence.
17	Q. Is that different than from a proponent of
18	intelligent design?
19	A. Yes, that's vastly 180 degrees different from
20	intelligent design. Intelligent design focuses
21	exclusively on the physical evidence. It relies totally
22	on empirical observations about nature. It does not
23	rely on any religious text. It does not rely on any
24	other such religious information. It relies exclusively
25	on physical evidence about nature and logical

1 inferences.

2	Q. Are intelligent design's conclusions or
3	explanations based on any religious, theological, or
4	philosophical commitment?
5	A. No, they are not.
6	Q. Again, can you draw any comparisons between
7	intelligent design and the Big Bang theory in this
8	regard?
9	A. Yes. Again, the both the Big Bang theory and
10	intelligent design may have philosophical or theological
11	implications in the view of some people, but again, both
12	are scientific theories. Both rely on observations
13	about nature. Both make reasoned conclusions from those
14	observations about nature.
15	Q. Does intelligent design require adherence to the
16	literal reading of the Book of Genesis?
17	A. No, it does not.
18	Q. Does intelligent design require adherence to the
19	belief that the Earth is no more than 6 to 10,000 years
20	old?
21	A. No, it doesn't.
22	Q. Does intelligent design require adherence to the
23	flood geology point of view which is advanced by
24	creationists?
25	A. No, it doesn't.

Does intelligent design require the action of a 1 Q. 2 supernatural creator acting outside of the laws of 3 nature? Α. No, it doesn't. 4 5 Q. Could you explain? Α. Making an analogy again to the Big Bang 6 Yes. 7 theory, the Big Bang theory is a theory which is advanced simply to explain the observations that we have 8 9 of nature, and it does so by making observations and 10 making inferences. It does not posit any supernatural 11 act to explain the Big Bang. It leaves that event 12 unexplained. Perhaps in the future, science will find an 13 14 explanation for that event. Perhaps it won't. But 15 nonetheless, the Big Bang is a completely scientific theory. Again, intelligent design is a scientific 16 17 theory that starts from the data -- the physical, 18 observable data of nature, and makes reasoned 19 conclusions from that and concludes intelligent design. 20 Scientific information does not say what is the 21 cause of design. It may never say what is the cause of 22 design. But nonetheless, it remains the best scientific 23 explanation for the data that we have.

24 Q. Can science then identify the source of design at 25 this point?

1	A. No, not at this point.
2	O. Does intelligent design rule out a natural
3	explanation for the design found in nature?
4	A. No. it does not rule it out.
5	O. Could you explain?
6	A. Yes, Again, barkening back to the Big Bang
7	theory, the Big Bang theory was proposed, and the cause
, 8	of the Big Bang was utterly unknown It's still utterly
9	unknown But nonetheless the Big Bang theory is a
10	scientific theory
11	The Big Bang theory does not postulate that the
1 1 1	The big bang theory does not posturate that the
12	Big Bang was a supernatural act. Although, you know, it
13	simply posits no explanation whatsoever. In the same
14	sense, intelligent design is a scientific theory
15	advanced to offer advanced to explain the physical,
16	observable facts about nature.
17	It cannot explain the source of the design and
18	just leaves it as an open question.
19	Q. We've heard testimony about methodological
20	naturalism. Are you familiar with that term?
21	A. Yes, I am.
22	Q. I believe you indicated in your deposition that
23	you thought it hobbles or even constrains intelligent
24	design, is that correct?
25	A. Yes, that's right.

Q. How does it do so?

1

A. Well, any constraint on what conclusion science can come to hobbles all of science. Science should be an open, no-holds-barred struggle to obtain the truth babout nature. When you start putting constraints on science, science suffers.

Yesterday, I discussed a man named Walter Nernst who said that the timelessness of nature, the infinity of time was a necessary constraint on a scientific theory. Science had to operate within that framework. If he had prevailed, progress, real progress in science would have been severely constrained.

Another reason why methodological naturalism can be a constraint on science is because oftentimes people don't think -- don't separate neatly categories in their own minds. For example, I showed the -- I showed the quotation from John Maddox, the editor of Nature, who found the Big Bang theory philosophically unacceptable and was reluctant to embrace it because of that.

There are other scientists in the past, one named Fred Hoyle, who rejected the Big Bang theory because he did not like its non-scientific, extra-scientific implications. So to the extent that people confuse a scientific theory with extra-scientific implications that some people might draw from it, then that might --

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Ţ	that might be a constraint upon the theory.
2	Q. Despite these constraints, does intelligent
3	design still fit within the framework of methodological
4	naturalism?
5	A. Yes. Despite the constraints, it certainly does,
6	just as the Big Bang theory does.
7	Q. Now we've heard some testimony about space aliens
8	and time traveling biologists. And I believe you made
9	some similar reference to that in your book, Darwin's
10	Black Box, is that correct?
11	A. Yes.
12	Q. And why was that?
13	A. Well, this was, you know, a tongue-in-cheek
14	effort to show people that, you know, intelligent design
15	does not exclude natural explanations, although some,
16	you know, explanations we might wave our hands to think
17	up right now might strike many people as implausible,
18	they are not, you know, utterly illogical.
19	And it was kind of a placemaker to say that maybe
20	some explanation will occur to us or be found in the
21	future which will, in fact, be a completely natural one.
22	Q. Now the space alien claim in particular seems to
23	fall hard on the ear of a lay person. But has that been
24	a claim that has been advanced by a notable scientist to
25	explain the natural phenomena?

1	
T	A. Yes, that's right. Surprisingly, in the year
2	1973, a man named Francis Crick, the eminent Nobel
3	laureate who discovered the double helicle shape of DNA
4	with James Watson, he published, with a co-author named
5	Leslie Orgle, he published a paper entitled Directed
6	Panspermia, which appeared in the science journal
7	Icarus.
8	And the gist of the paper was that the problems
9	trying to think of an unintelligent origin of life on
10	Earth were so severe that perhaps we should consider the
11	possibility that space aliens in the distant past sent a
12	rocket ship to the Earth filled with spores to seed life
13	on the early Earth.
14	Q. This was a claim advanced by a Nobel laureate?
15	A. Yes, Francis Crick.
16	Q. And the article in which his arguments appear,
17	was this a peer reviewed science journal?
18	A. Yes, the journal Icarus.
19	Q. Was this just a tongue-in-cheek, so to speak,
20	explanation on behalf of Francis Crick?
21	A. No, it wasn't. He mentioned it first in that
22	1973 article, and he repeated the same claim in a book
23	he published in '88 and interviews he gave later on.
24	And from what I understand, he still thought it was a
25	reasonable idea up until his death recently.

1	Q. Sir, I'd ask you to direct your attention to the
2	exhibit binder that I have provided for you, and if you
3	could go to tab 14. There is an exhibit marked as
4	Defendants' Exhibit 203-E as echo. Is that the article
5	from Francis Crick that you've been testifying about?
6	A. Yes, this is Francis Crick's article on Directed
7	Panspermia.
8	Q. Is the search for intelligence causes a
9	scientific exploration?
10	A. Yes, it is.
11	Q. Again, do you have any examples that we could
12	point to?
13	A. Well, one good example is one that I mentioned
14	earlier, which is this project called the SETI project,
15	S-E-T-I, which stands for search for extraterrestrial
16	intelligence, where scientists use instruments to scan
17	space in the hope of finding transmissions or some
18	signals that may have been sent by extraterrestrial
19	sources.
20	And they are confident that they could be able to
21	distinguish those signals from the background noise,
22	background radiation, electromagnetic phenomena of
23	space.
24	Q. Again, that's a scientific exploration?
25	A. Yes, a number of scientists are involved in that.

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1	MR. MUISE: Your Honor, I'm just do you
2	intend to go to 12:30?
3	THE COURT: I was thinking more 12:15,
4	unless you think that this is an appropriate break
5	point. Your call.
6	MR. MUISE: I certainly have more than 15
7	minutes. This next section might be divided in that 15,
8	so my preference would be to take the lunch break and
9	come back and then complete the direct during the first
10	session after lunch.
11	THE COURT: All right. We'll return then
12	at, let's say, 1:25, this afternoon, after a suitable
13	lunch break, and we'll pick up with your next topic on
14	direct at that time. We'll be in recess.
15	(Whereupon, a lunch recess was taken at
16	12:04 p.m.)
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