Let's see. Is that number one? Could you go back one just to make sure I didn't hold it down 12 times? Yeah, okay.

So what I'd like to do initially is talk about this basic premise about tumorigenicity as it relates to cell substrates. I think that's an area of concern when I read over the documentation. The potential risk statement that Dr. Lewis has provided: what are the determinants of experimental tumor formation that are being used to characterize these tissue culture cells used for vaccine production?

And then some general observations thinking about experimental tumorigenicity as the data that we understand from that relate to potential risk, and then, as I mentioned, tell you something about E1A induced sensitivity to apoptotic injury and E1A induced rejection of cells expressing these proteins, and then just a very little bit at the end about E1A in humans.

So the focus here is on the E1A as the immortalizing, enabling oncogene in the cells that are being created, and I guess theoretically its potential risk as a contaminating bit of DNA that would go across in the vaccine.

So the potential risk statement cut down

a bit says something like tumorigenic cells are more 2 non-tumorigenic cells than vaccine substrates, and that seems to be a focus of -- an 4 appropriate focus of concern, and taking that directly, I think, is an interesting thing to do, and this comes, as Dr. Lewis mentioned, from this Armed Forces Experimental Board in 1954 where there was a preference for normal cells over human tumor cells, not cells like 293 or PER.C6.

> And so I think the question at least as I think of it is the concern about the use of human tumor cells, like HeLa or something of that sort, the same thing as the concern about using human cells that form tumors. Those things may seem to be similar, but I think they're different, and that is form tumors in nude mice.

> So what I'm going to try to convince you is that I think experimental tumorigenicity as we measure it is not the same thing as primary tumor development because I think this is an important distinction. We're talking about the tumorigenicity of E1 expressing cells or E1 immortalized cells, and I don't think that's the same thing as tumor formation from a single immortalized or mutated cell in vivo that goes on to successfully form a tumor in us or in

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1 a mouse.

And it's important, I think, not to confused these things.

So the variables that can be affected, that can affect experimental tumor formation, that is, the ability to form tumor in an animal like a nude mouse with a set of tissue culture cells, the first thing is the host that you select. That makes a big difference.

And this shows you something about some experiments we've done where you can see a huge host difference in the apparent tumorigenicity of the cells.

Let me crank this up.

So these are three sets of types of mice. These are normal C57 Black 6 adult mice. These are C57 Black 6 mice that are nude and, therefore, lack of functional thymus and lack of functional T cells that have an intact natural killer cell response, a different kind of host defense.

And these are CD3-epsilon transgenic mice, and it's not interesting to know what they are, except for the fact that they lack both T cell and NK cell responses.

And then what's been done here is to take

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mouse sarcoma that is a relatively non-immunogenic mouse sarcoma, and this is collaborative study done with Jack Routes, and these cells were either tested in the three types of mice, either in their E1A negative parental state or in the state that they would be after being transfected with the E1A gene alone, no E1B, not the whole E1 region, just the E1A gene.

And the question was: did E1A expression affect the tumorigenicity, and if so, did that relate to the type of animal that you use to test tumor development?

And the answer is obvious. All of the red lines are the control cells not expressing E1A, and all of the green lines are the efficiency of tumor formation using the parameter that Dr. Lewis mentioned, the TPD-50, the efficiency of tumor formation by the cells in the different animals.

And the lower the TPD-50, the more efficient tumor formation occurred. So you can ignore all of the red lines because they're essentially all the same, and that is the non-E1A expressing sarcoma cells were very efficient at forming tumors in all three types of animals. The E1A expressing cells were essentially unable to form tumors in normal mice,

formed tumors sort of in a medium fashion reasonably well in nude mice, those that lacked a T cell response, and formed tumors with equal efficiency as the EIA negative cells in the animals that lacked both types of cellular immune defenses.

And so the type of host you pick determines a lot about the type of answer you get. You can say that this cell is, quotes, non-tumorigenic, whereas this cell is highly tumorigenic, but it really just depends on the animal chosen. So I think that's clearly an important parameter to consider.

The next thing that's important is the threshold effect. Dr. Lewis mentioned numbers of cells used to challenge animals, and so the cell dose you pick determines the result you get as well. So, again, you could create a non-tumorigenic cell by just picking too low a dose.

And here is an experiment from a study that Dr. Lewis did with these Adeno. 12 transformed mouse cells. These were Balb/c cells, and they were inoculated into adult immunocompetent animals at various doses over time. This is the number of cells injected. This is the tumor incidence of these animals.

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And what you can see is essentially no tumors up to 100,000 cells. 2 So if you happen to get unlucky and pick 100,000 or ten to the 3 fifth cells as your number, you'd call these non-4 5 tumorigenic cells. On the other hand, if you picked 100 6 7

million or 10 million cells as your dose, you'd say they're highly tumorigenic. So dose matters. matters, dose matters.

The next thing is the rout of inoculation. This is something that's not as well quantitated, but there's no question in my mind that the route that is used for a given challenge makes a difference, and in our experience intraperitoneal inoculation results in an apparently more efficient tumor positive result than subcutaneous inoculation, and there are other kinds of things you could do, like inoculate into the immunologically privileged sites.

Another result is to think about observation period. How long do you wait before you score tumor development versus non-tumor development when you're doing an experimental tumor formation assay?

This in our laboratory is a notorious experiment because we did a collaboration with another

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investigator. He had published a paper saying that
human breast cancer cells expressing the E1A gene were
non-tumorigenic and E1A had, in fact, converted these
cells into non-tumorigenic cells. It had reversed
their tumorigenicity.

Well, the thing was he looked at these cells only after about 14 to 18 days. We repeated the experiment, and so what's shown here is the percentage of animals that are tumor free over time after a challenge, with the breast cancer cell itself. These are all nude mice. So they lack T cell responses, have some NK cell defenses. So this is the E1A negative breast cancer cell. This is a ductile epithelial cell carcinoma.

There was the E1A expressing cells, and sure enough, if you look at about two to two and a half weeks, you get zero tumor formation or 100 percent tumor free animals, whereas all the animals that had been inoculated -- that's not true -- some of the animals that had been inoculated with the E1A negative cells developed tumors, and that tumor incidence continued to increase over time or the tumor free number decreased.

But if you waited a bit longer, then the E1A positive cells begin to make tumors in up to 20

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percent of the animals over time. So if you get unlucky and pick a very short observation period, you might come up with the wrong conclusion and call a cell nontumorigenic.

So the host matters. The cell dose matters, and the route of inoculation matters, and how long you wait to call the endpoint matters.

Well, then the last thing, I don't have any data to show on this, but there are several other things when you look at the literature that can make a difference in the methodology chosen to determine whether cells make tumors or don't.

One thing that I find interesting because almost all of these experiments are done by sticking a needle into a mouse or a hamster or a rat, injecting some cells into that needle track, and then asking whether they form tumors or not. That's not normal. That's not what happens after a UV irradiation or exposure to a carcinogen. This causes some kind of trauma at which site the cells now have a chance to grow.

And so I think there may be a wound effect, and we were talking earlier about some of the observations that have been made in Rous sarcoma virus where tumors form at the site of wounds.

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That is an artifact that can affect tumor formation, I think. Foreign bodies can affect tumor development as well. For example, you can attach cells to plastic plates and put them into animals and get tumors. These cells wouldn't form tumors at all through any route of inoculation like 3T3 cells if you didn't attach them onto these little disks, and so the foreign body can make a difference in your tumor assay.

So you have to be careful about how you interpret those, and the other thing that can happen is if you just provide fibroblast as a mixed culture in with your putative tumor cells, you can get tumor forming efficiency that is much greater. So there is some feeding effect or something that goes on when you can do certain kinds of mixed culture inoculations.

So my point is that tumorigenicity is not all the same. It depends on how you rig the system, how you set it up. All of these experiments can be used quite effectively to ask questions about one cell versus the other, but I don't think they are anything like at least in my mind what goes on when you get a spontaneous single transformation event and ask that cell to grow into a tumor, especially when you think about cell dose.

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There are very few spontaneous transformation events that result in the immediate generation of 100 million cells or ten million cells.

So what I'd like you to think about as a possibility is the fact that tumor formation is not the result of a single activity at least as we look at it in experimental tumorigenicity, but it's really an orchestra of events, and if all of these things aren't working right, you don't get a very good outcome or in our case you don't get a tumor.

You need to have in most cases that we know about continuous oncogene expression or continuous in our case E1A expression. That probably isn't always true, but almost always true.

That gene has to successfully cause cell cycle disregulation, and it probably has to hit to Achilles heel in the cell cycle that relates to what some people call the p53 Rb pathway where something has to occur to cause that to go out of whack. Otherwise you don't get immortalization.

The cells gave to survive this, and it's important to think about the fact that when you try to over express E1A in normal mammalian cells it usually kills them. So it is a proapoptotic event itself. So it's not easy if you're trying to transfer a lot of

E1A and some contaminating DNA to get those cells to survive because they would tend to die as a result of E1A expression itself.

Then you have to have this magical threshold cell number. I don't think single E1A positive cells probably ever form tumors. I was taught early in medicine never to say "never" or "ever," but I think it's probably true that it's probably true that it's probably true that it's very hard to get a tumor with a single E1A positive cell. You probably need millions at least.

There probably would have to also be local tissue factors. Steve Frisch from California has shown that if you plate E1A positive cells on collagen matrices as opposed to plating them on plastic, they tend to do poorly. So you could argue that maybe the reason that E1A positive cells grow poorly in nude mice compared to normal mice is that they don't like being on biometrices and that tends to be inhibitory to their growth. Most of us grow them on plastic dishes and they look fine.

And then they also have to escape from this cellular immune response that I described in this three-mouse experiment. If they don't escape from that, then the cells are likely to be killed, even if

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they make it through all of these other hurdles. So I think it's very unlikely that this events would occur of orchestrated fashion with transmitted DNA without a lot of help. I'm sorry. Would you go back one? This is just a summary of a lot of work done on trying to characterize how E1A sensitizes cells to a variety of proapoptotic injuries. We've mostly been interested in immunological injuries since they relate more to whether a cell will form a tumor or won't form a tumor in vivo, but a number of other

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So if you look at a variety of types of things, chemotherapeutic drugs, hydrogen peroxide, potassium ionophores, irradiation; this is a protein synthesis inhibitor; where the things we're more interested in like natural killer cells, cytolytic T cells, TNF alpha, or the TNF receptor apoptosis inducing ligand trail, all of these things can

selectively induce apoptosis in an E1A expressing

injuries do the same thing, and that tells us

something about the mechanism, I think.

24 So there were a lot of things going

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against E1A positive cells in trying to survive in

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cell.

vivo or trying to survive when exposed to a variety of 1 2 types of injuries. 3 This may say something about the pathway through which E1A sensitizes, but what it does say is 4 that an E1A positive cell doesn't have an easy time of 5 it trying to survive in vivo, especially when you 6 7 consider these kinds of defenses. 8 It's also important that these events, 9 this sensitization of cells to immunological mediated 10 proapoptotic outcomes are not necessarily dependent on 11 normal cell genetics. So p53 minus cells are cells 12 lack expression of the Rb gene, 13 sensitized by E1A just like normal cells are. 14 So if you wanted to postulate the worst 15 scenario, you'd say, well, maybe EIA would be 16 transmitted, and it would hit a cell that already had a mutation and that would give a chance for a bad 17 18 outcome. 19 The fact is the most common mutations in 20 human cells that lead to neoplasia are in this 21 pathway, E1A sensitized cells expressing mutations in 22 that pathway to immunological injury. So they're not 23 going to be protected by that second mutation. 24 This just shows you something about the 25 range of experiments that have been done with human

cells. This is just an experiment with natural killer cells to show you that human cells are very much like other cells. These are mouse 3T3. This is a continuous rat line that was immortalized with <u>ras</u>. This is BHK21, a spontaneously transformed hamster cell line. THE BHK21 is highly tumorigenic, and its tumorigenicity is eliminated by expressing E1A.

These other lines, all when expressing E1A become more sensitive to natural killer lymphocyte induced lysis, and this is just a human fibrosarcoma cell line, stably expressing E1A, and it does the same thing.

If you generalize this to other types of things that we know about, a variety of types cells from different tissue origins from humans are sensitized to various apoptotic injuries. Epithelial cells, notable among those 293. We've also looked at this breast cancer cell I told you about; fibroblastic cells. This is a fibrosarcoma line.

Hematopoietic cells, K562 has been shown by Stiewe in a publication in 2000 to become sensitized to reactive oxygen intermediates when expressing E1A, but not when not expressing E1A, and ACLS2, which is an interesting cell line because it doesn't express either p53 or Rb, when it expresses

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E1A becomes much more sensitive to natural killer cell injury, and I've already told you about some of the injuries. These are the ones that have been tested against human cells expressing E1A.

So this is like the atomic chart or the intermediary metabolism pathway. It's one of these things that we never remember, and I just show it for the purpose of saying that EIA probably hits multiple targets in the apoptosis cascade. This is an incomplete representation of what goes on during apoptosis, but there are preliminary data. This isn't well worked out, but there are data that EIA causes over expression of p53, which can drive apoptosis. EIA positive cells -- this was all done in 293 cells -- express some kind of what's called an oncogene associated factor that is probably like a mitochondrial-like factor that can be involved in apoptotic activity.

of Procas Base 8 to Cas Base 8 (phonetic). In certain cell types, it can increase cell surface depth receptor expression, and we've done several studies now, and I've just submitted a publication about E1A repression of the cellular NF kappa B (phonetic) defense against apoptosis.

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So probably the bottom line is that there are multiple sites in cells that are targets for E1A rendering cells sensitive to a variety of apoptotic injuries, especially immunological ones.

So the last point is E1A in humans, and I guess to try to put the idea of contaminating E1A in perspective, we should think about what goes on elsewise (phonetic), and that is there are studies. This is a study by Jimmy Hogg from Canada, from his laboratory, that shows that E1A actually may persist in normal human tissues after infection.

In these studies what they did is they took a lot of people that had lung disease, and they asked whether those that progressed to emphysema or chronic bronchitis had something unusual about them, and I think the truth be told, he had somebody who had done a post doc in Alex's laboratory or somebody's who had learned how to look for E1A, and they looked for E1A in these lung tissues, and what they found was a lot of E1A sequences in both normal people and those that progressed to emphysema based on PCR technology.

So it's not easy to do, but when people have looked, they have found E1A, at least in this one laboratory, have found E1A in normal tissues, and that may speak to the persistence of adenoviral infection.

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It may say that there is integration that goes on that
we aren't very good at detecting, but it's probably
not true that EIA is never around and these viral
oncogenes are never expressed in humans anyway, even
vaccine recipients.

And then there are two clinical trials that are going on now and fairly extensive studies in which ElA is actually being used therapeutically. It's being put into people that have cancer to ask whether it can help reject tumor cells. One of those is in a company that has the name of the virus that's called ONYX. It's really an old Arnie Levine that has a mutation that allows it to selectively kill p53 minus cells, which many human tumors are.

So this virus infection is given to people who have p53 minus tumors, and that virus preferentially replicates in those tumors and destroys the tumor tissue preferentially to normal tissue.

It's not that simply, but that's the idea.

There is also an M.D. Anderson study in which E1A is being used in gene therapy to inject in liposome vectors into people that have, for example, metastatic ovarian or peritoneal metastasis with ovarian carcinomas, and it's also being considered for breast carcinomas.

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And here the idea is that E1A represses expression of a cell surface receptor that's necessary for growth factor stimulation of the tumor cells. not only do we probably have E1A already in our If we were to look carefully and certainly can support persistent infection with adenoviruses. but E1A is being used therapeutically in humans, and far admittedly so now these aren't long-term vaccinations of babies, but so far there's no evidence from the safety testing that this has caused an adverse outcome in these patients.

So my conclusions are that experimental formation in nude mice does not predict tumorigenicity of an isolated oncogene, in this case Even if the ElA oncogene were transmitted in a contaminating DNA, it's very unlikely that it would become stably expressed in recipient cells because of transfection inefficiency. Dr. van der Eb just told you how hard it is to transfect human cells with sheared adenovirus DNA or certainly with E1A itself it's actually even harder.

Direct apoptosis that occurs during attempted establishment of these cells would probably kill many of the cells in which the DNA would try to get itself inserted.

Even if E1A did become stably expressed in 1 recipient cells, it's very unlikely that such cells 2 would provide a tumorigenic risk because 3 wouldn't be a threshold cell dose. You wouldn't have 4 the millions of cells up front that you'd need to get 5 over the hurdle of getting tumor formation initiated. 6 Now, the viability of the ElA positive 7 cells on biometrics, Steve Frisch's data, suggesting 8 that these cells do relatively poorly when cultured on 9 things like we might have in our soft tissues, and the 10 susceptibility of these cells even if they did get 11 established to immune mediated apoptosis, I think, 12 13 would rule against their survival. 14 And then also we have to think about the 15 fact that expression of E1A in humans might be normal, 16 and when therapeutically used in humans, actually be directly apoptotic and can 17 expression of critical growth factor receptors. 18 19 think E1A expression 20 contaminating force from the vaccine would probably 21 not likely be a major problem. 22 I'd be glad to answer questions. 23 ACTING CHAIRMAN DAUM: Thank you very 24 much, Dr. Cook, for an enlightening presentation. 25 We'll take a few questions. Dr. Griffin,

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then Dr. Stephens, please, and then Dr. Faggett. 1 2 DR. GRIFFIN: It's my understanding that the cells we're talking about express both E1A and 3 E1B. 4 5 DR. COOK: Right. 6 DR. GRIFFIN: And that E1B is a counter force, an anti-apoptotic factor. So how does that fit 7 in with all of your conclusions that you're making 8 9 about the safety of E1A when you're really talking 10 about giving both? DR. COOK: Right. I specifically didn't 11 12 talk about E1B. There's no evidence that E1B 13 expressed alone does anything in terms of its ability to immortalize cells. 14 15 They're looking at both. The only reason 16 that E1B is interesting in the contest of E1A is it allows E1A -- it prevents E1A from destroying cells. 17 So without E1B what you get is a tremendously 18 inefficient immortalization event, very few colonies 19 20 formed. So E1B really has to be around to prevent the 21 apoptotic response and to bind p53, the two proteins 22 that E1B makes. 23 So I don't think that E1B is a factor. 24 All I can tell you is when it gets down to what I 25 think is the key event at least from our perspective

is that if you look at the immunological injuries and 1 ask whether they still happen, from a killer cell 2 point of view, a natural killer cell, a cytolytic 3 lymphocyte or an activated macrophage still kills 4 cells expressing E1B perfectly well, even though it 5 does it in an E1A specific manner. 6 7 The only thing that E1B can do in human 8 cells when co-expressed with E1A is to repress the TNF induced apoptotic response. That's a unique thing 9 that Linda Gooding showed years ago. 10 11 So I don't think E1B really adds a whole 12 lot to the mix, other than allowing the cells to be 13 immortalized in the first place. 14 Then the other thing that I would have to think about is how do you think about transmitting 15 this DNA when these are two separate genes under the 16 control of two different promoters, and now what you 17 18 have to do is get the orchestra to work even harder. 19 You've got to put the two genes together again to get them expressed in the same cell if you're going to try 20 21 to transmit this as contaminating DNA. 2.2 So now the odds even go up higher. 23 ACTING CHAIRMAN DAUM: Thank you very much. 24 25 Dr. Stephens and then Dr. Faggett.

1 DR. STEPHENS: I was struck by one of your 2 papers talking about this family of oncogenes, and you 3 compare the HPV oncogene, which has very different properties than EA-1, and I just wondered if you could 4 5 comment the structural relationships possibility of mutation of EA-1 to produced an HPV-6 7 like oncogene. 8 DR. COOK: Well, E1A and E7 have some 9 sequence similarities. You can actually do with an 10 analogous gene SV40, you can do mix and match experiments where you can cut out a bit one, stick 11 into the other and get a competent gene. 12 13 So they have conserved sequences. I think the reason for this is that all of these genes need to 14 15 do the same thing when they get into the cell, and that is to regulate the cell cycle, and they do that 16 by binding Rb and by binding p300 and Kreb binding 17 18 proteins. 19 So there are reasons that they have co-20 evolved these. They aren't the same virus. As far as anybody know, they didn't evolve together, and they 21 22 aren't just recombinations. The data you're referring to are that EIA does all of the things I'm talking about, and so far

as we can tell, when E7 is expressed in cells with the

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exception of one or two reports, it doesn't sensitize cells to some of these other kinds of injuries like TRAIL or to killer cell injury, and so E7 is a different beast, even though it shares some of these same cell cycle regulatory bits.

The ability to convert E1A into E7, other than making a chimeric molecule, which can work -- Jack Routes has done that. He's made an E1A E7 chimera. What happens is the E1A phenotype is dominant, and so E7 sort of lacks what E1A has. It doesn't look like it has any evil force that's trying to outdo E1A.

There are no mutations so far that we have been able to make in E1A that's eliminated these activities, although if you eliminate enough of the genome, if you take enough of E1A out, you can reduce its ability to sensitize things, but it's not just a simple mutation. It's a deletion.

So I don't know that we could, other than chimera formation between E1A and E7 just making single based changes or some kind of frame shift or something in E1A create an E7-like gene product.

ACTING CHAIRMAN DAUM: Thank you.

Dr. Faggett, please.

DR. FAGGETT: Thank you very much.

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It was a very clear presentation of a very complex topic.

You did mention that you were comfortable in terms of safety issues with the A-1. What was the level of sensitivity and types of tests used to bring you to that level of comfort in terms of safety?

DR. COOK: Well, what I'm saying is that I think the -- I'm talking about odds. I suppose we could talk to statisticians, which I'm not, but the odds of getting multiple unlikely events that occur at rate of maybe one in a million to occur simultaneously to get the event that you want to see happen, which is the transfer of an E1A oncogene into a vaccine recipient at the level of DNA contamination you're talking about, and have that gene by itself now be able to immortalize the cell, have that cell survive itself, and then have it survive an immune response. I think the odds could be -- probably can't be calculated because you don't know all of the numbers, but I think they're very low.

DR. FAGGETT: It's a question of ten nanograms of DNA fragments estimated to be the smallness to get some effect. Is that a level?

DR. COOK: Ten nanograms per vaccine dose is what I think I heard earlier.

1	DR. FAGGETT: Okay. So how do you
2	DR. COOK: I've have to try to do some
.3	math. I think it would be mental gymnastics, but it
4	probably could be done.
5	DR. FAGGETT: Thank you.
6	ACTING CHAIRMAN DAUM: We will return to
·7 .	this very subject in our discussions later.
8	Dr. Diaz.
9	DR. DIAZ: You mentioned that there was a
10	lot of E1A just naturally occurring in human cells,
11	different types of human tissue. Could you expound on
12	that a little bit in terms of what's known? Is it
13	expressed? Does it perform similar immunologic
14	regulatory functions in humans?
15	And, secondly, is there any known E1B-like
16	genes in human tissue also?
17	DR. COOK: Yeah, it's important to put it
18	into perspective so that I don't overstate this. So
19	there is a laboratory in Canada in which they have
20	found evidence based on PCR analysis of E1A sequences
21	in lung tissue. It was in a lot of different lung
22	specimens that they sampled.
23	As far as I know, there aren't other
24	laboratories that have done the same thing.
25	There's a laboratory in Australia where

they've found E1A-like activity in mammalian cells, including human cells, but haven't convinced anybody 2 that it's E1A yet. 3 4 So there's a different in finding E1A, as Hogg's lab did, and finding E1A-like activity, as many 5 other people have. So there probably are E1A-like 6 7 functions in normal cells, and that's probably why E1A decided to ape this during evolution anyway, because 8 9 it helps regulate cell cycle. 10 A far as E1B expression in normal human cells outside of initial infection or persistent 11 infection, I'm not aware of any data that 12 independently has been detected in human tissues 13 1.4 unrelated to infection. 15 ACTING CHAIRMAN DAUM: There are four more 16 We'll take these four questions, and then 17 we're going to move on. Dr. Minor, then Dr. Ketner, 18 Dr. Aguilar-Cordova. 19 DR. interested MINOR: I'm in your threshold description. If you had a PD-50 of 1,000 20 21 and you give it to ten mice, five of them will go 22 down. 23 If you had 100 cells, for example, went 24 into 100 mice, are you saying that none of them will 25 qo down?

1	DR. COOK: I think that's true, actually,
2	yeah.
3	DR. MINOR: Has anybody ever done to see
4	DR. COOK: Andy, you've given a lot of
5	you know the answer.
6	DR. LEWIS: Yes, we did that very
7	experiment with the Adeno. 12 because we found that
8	the TPD-50 for the Adeno. 12 transformed Balb/c mice
9	embryo cells was a million cells, and so we asked the
10	question: what would happen if you put 100,000 cells
11	into 100 mice?
12	And we did that experiment, and we got no
13	tumors. So it looks like that threshold is real.
14	We did several other experiments to
15	address that question the same way, like cloning and
16	doing this, that, and the other, and the TPD-50 on
17	sub-clones of the population were again a million
18	cells.
19	So nothing we could do would alter that
20	threshold. I can't say that there's thing that
21	couldn't be done to alter the threshold, but we were
22	unable to do it considering that as a criticism for
23	the proposal we were making.
24	ACTING CHAIRMAN DAUM: Can I ask a follow-
25	up and this be save one of the four questions?

Is this threshold phenomenon in your belief a probability event where if you just give enough cells, one will do it, or is this a quorum sensing kind of thing where if you give enough cells they will start doing something?

DR. LEWIS: Yes. Well, I think what we ruled out by the experiments we did was the possibility that one cell out of a million cells was doing the job. All the data that we have suggested it took the collective action of million cells to overcome whatever it was that was prohibiting the cells to form a tumor mass.

Now, exactly what it's overcoming I don't know. I think the question was as to whether this was -- and these were adult Balb/c mice, by the way. They were not newborns or immunoincompetent animals. These were adult animals.

So the question I think we raised but were never able to answer was the possibility that it was something about the immune system that it took a million cells to overcome. In other words, that was the level of competence at which that animal could function immunologically with all of his anti-tumor defenses intact.

And if you challenged him with enough, he

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could no longer respond. Jim may have some other 1 ideas about that, but that's as far as we were able to 2 3 take the logic. We were never able to test that. 4 ACTING CHAIRMAN DAUM: It may be fun to return to that this afternoon as an issue. 5 to get the focus back to your presentation. 6 7 Dr. Ketner is next, and then Dr. Aquilar-8 Cordova. DR. KETNER: Yeah, this is just a comment 9 10 on the persistence of E1A sequences in human lung 11 samples. It turns out it was shown a long time ago, 12 20 years or so, that human peripheral blood contains, after an adenovirus infection has come and gone, 13 14 contains residual, intact viral genomes. 15 probably replicating slowly in peripheral blood cells, 16 and so those E1A sequences that were found, these are 17 detectable by relatively insensitive techniques like Southern blotting. 18 19 So those E1A sequences that were detected 20 by PCR may well just have been intact replicating 21 genomes in the blood in those tissues. So I think it 22 overstates it to say that there are cells in which 23 there are, you know, E1A persistent sequences. 24 ACTING CHAIRMAN DAUM: Thank you for a 25 clarifying comment.

Dr. Aguilar-Cordova.

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DR. AGUILAR-CORDOVA: Yeah, I was wondering on the cell thing you were mentioning that had E1A-like activities if these cells could complement the E1A minus vectors, and what other known genes, such as E7 or other viral genes, may have that function of complementing and then producing the replication competent activity.

DR. COOK: The clearest experiments have been done with mouse embryonal carcinoma cells, the cell line F9, and I think some of the break-away studies from Australia may be similar, but in F9 cells what happens is these are interesting cells because they can be differentiated in vitro with retinoic acid into terminally differentiated cells. Without retinoic acid, they're undifferentiated embryonal carcinoma cells.

The undifferentiated cells will complement a virus called DL312, which has no E1A and grows very inefficiently in normal cells. When these cells are differentiated, that complementation goes away.

So this is the basis upon which it has been concluded there is some E1A-like activity in normal cells. This is a mouse embryonal carcinoma cell.

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1	The question about whether other human
2	persistent viruses or like HPV where E7 is expressed
3	in an epithelial carcinoma or something like that, I
4	don't know of any evidence that E7 can be expressed in
5	the cell and complement a defective adenovirus. I
6	don't think at least as far as I know, that's not
7.	the case, but I don't know the data.
8	ACTING CHAIRMAN DAUM: Thank you very
9	much, Dr. Cook.
10	We're going to move on now to our final
11	morning, so to speak, presentation. We'll hear from
12	Dr. Peden at the FDA regarding quantitative assessment
13	of the risks of residual DNA, and after which we'll
14	take a break for lunch.
15	DR. PEDEN: Thank you.
16	ACTING CHAIRMAN DAUM: While setting up,
17	I'd like to congratulate Dr. Peden for providing us
18	with a handout with his slides on it. It makes it
19	much easier to follow the talk.
20	DR. PEDEN: Thank you.
21	I was going to say good morning, but it
22	is, in fact, good afternoon.
23	This presentation will discuss issues of
24	whether residual cell substrate DNA in vaccine
25	manufactured in neoplastic cells poses a risk to
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vaccine recipients.

At the outset, I should like to say that while these views expressed are largely my own, I would like to acknowledge the sage and helpful discussions of many colleagues, particularly Andy Lewis, Phil Krause, and Becky Sheets.

I would discuss what the perceived risks are, the proposed mechanisms whereby there could be a risk. I will review the relevant in vivo data that do exist and present some calculations and assumptions upon which they are based.

The overall conclusion I want to leave you with is that the small amounts of residual cell substrate DNA, and by this I mean ten nanograms, poses an acceptably small risk to vaccinees, and this value of ten micrograms did not come from my whim.

Oops, can I go back one more?

It has been discussed for many years, and at the WHO expert commission in biological standardization in 1998, the residual DNA from continuous cell lines, ten nanograms or less, was proposed as a reasonable level, raising it from 100 picograms, .1 nanograms or less, that was reached in 1986.

So what types of risks are there? There's

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the perceived risks associated with residual cell substrate DNA, has generally considered oncogenic risk, but there is another risk which I'm going to discuss, too, the infectivity risk.

So the proposed mechanisms whereby an oncogenic risk by cell substrate DNA comes for several One is with the introduction of a dominant activated oncogene. For example, an activated rat oncogene or in the issues we're discussing today, the dominant activated oncogenes could be E1A or E1B.

Insertional mutagenesis is another mechanism whereby DNA could be oncogenic. While this mechanism has been found in certain cancers of animals, such as the disruption of cellular tumor suppressor gene or activation of a cellular dominant oncogene, by promotion of insertion mechanisms, calculations have been done by many people, and the risk associated with a single integration event by this type of -- to disrupt the cellular tumor suppressor gene or dominant oncogene is considered by people like Reinhart Court (phonetic) to be one in ten to the minus 18 or one in ten to the 18.

So this is a very low probability event, and it comes about -- this has been seen in animals were activation has occurred from an integrating,

replicating system, such as a retrovirus, and a single insertion event from residual DNA is unlikely, very unlikely to pose this.

Another mechanism that's been pioneered by Walter Doerfler is that DNA methylation pattern changes following integration. While this occurs in system trends, the lack of a phenotypic consequence of this makes this very difficult to study.

The other risk of integrating -- of cell substrate DNA is the one I talked about, cell substrate DNA they incurred in infectious genome. The DNA virus, which is polyoma virus, Herpes virus, and papilloma virus exist as integrated into cells.

Well, in addition, of course, retroviruses have a provirus state, such as HIV-1 and HTLV and human retroviruses. So by infectious DNA what we mean is that if this DNA is introduced into the appropriate system, then infectious particles are produced, and an infection can be established.

So one other thing that influences the nature of the inoculum. For example, we concern linear versus circular DNA as not being studied greatly. The single stranded versus double stranded, where the state of the DNA is free versus the chromatin associated. In most lysed cells, of course,

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135 the DNA will not be free, but be chromatin associated. 1 2 This has not been assessed in any study except one, that chromatin was injected into animals. 3 The size distribution of the DNA is 4 5 obviously very important. If the DNA is sheared to a size that's below the size of a normal gene, such as 6

> Now, the route of inoculation can be important as you heard from Dr. Cook. Most of the studies, various animal studies were done with intramuscular, intradermal, subcutaneous, intranasal and oral route, and also by intraperitoneal route. However, an application in Phase 1 would be considered unfavorably, I would imagine, with this route.

> normal genes excluding neutrons are about 3,000 base

pairs. Then clearly this DNA will not be able to

Assumptions. Before we can go into the calculations, we need to make certain assumptions here, and I hope you can read that, but what I wanted to do is to go through these assumptions. I'll read them slowly, and I hope you have the handout.

For a given DNA, the level of response of the cell to that DNA is proportional to the amount of that DNA. That makes somewhat sense. The activity of the gene integrated in the chromosomal DNA was part of

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produce a product.

a plasmid or phage vector is equivalent or lower. Now, we have to make this assumption because we have no other way of doing this. However, it's likely that the DNA integrated into chromosomal DNA being controlled by its own promoter, requiring certain perhaps tissue specific transcription factors. may not, in fact, be active at all, whereas we put a gene on a plasmid or phage vector and we've designed it such that it expresses very well. So if anything, this is going to over estimate the risk rather than underestimating the risk, but that's the only thing we can do. So we're making this assumption. And the amount of uptake of a given gene by a cell and the expression of this gene in the cell is related to the concentration of the gene in the DNA. Now, because a single copy is represented approximately one-millionth of at the mammalian genome, the amount of DNA corresponding to a single copy gene is a million-fold less abundant for equivalent amounts of cellular DNA compared with the plasma DNA with the same gene.

Now, as I say, the genome is of nine -the haploid genome is three billion base pairs, where

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haploid

as I say, the normal gene is maybe 1,000 base pairs. 1 So there's a million-fold difference in the size of 2 3. the genomes. 4 So, for example, that is, for the uptake and expression of a gene in mammalian genomic DNA 5 equivalent to one microgram of plasma DNA, one million 6 7 micrograms or one gram of mammalian DNA is to elicit equivalent biological 8 an effect, giving the 9 assumptions on the previous slide. Conversely, the single oncogene is present 10 11 at one microgram of mammalian DNA, the equivalent amount of the same oncogene if cloned in the plasma is 12 13 one times ten to the minus six micrograms or one times ten to the minus three nanograms or picogram. 14 So there are the assumptions on which 15 16 we've made some calculations. 17 So now I want to go through some of the 18 data that has been presented in the literature about oncogenicity, and this is the work done by Hsien-jen 19 20 Kung, and colleagues, in 1983, where they're using a 21 cloned <u>src</u> gene, two micrograms induced tumors in 22 seven out of ten chickens inoculated subcutaneously in 23 the wing web. 24 If you inoculated a cloned Rous sarcoma 25 genome, the entire genome, this two micrograms induced

tumors in six and six. Presumably this comes about
because the Rous sarcoma virus DNA initiated an
infection.

So two micrograms was found to induce

tumors in about 70 percent of the animals. A study in 1990 by Halpern and colleagues got essentially the same or similar results using in this case 20 micrograms of DNA by the same route, and inoculated intravenously. Also it gave tumors.

So we can conclude that two micrograms of DNA, which is equivalent to about two and a half times ten to the 11 molecules of cloned <u>v-src</u> oncogene is oncogenic in chickens.

The study that was done by Burns and colleagues inducted the oncogenicity of <u>ras</u>, of the activated <u>ras</u> gene in mice. Activated <u>ras</u> from the T-24 gene, ten micrograms inoculated by scarification in the mouse skin.

Lymph anginose sarcomas developed in 33 out of 34 animals within 12 months, usually within 12 weeks. However, the normal <u>ras</u> gene failed to induce tumors. So at least in this route, ten micrograms, which is equivalent of about one times ten to the 12 molecules of an activated <u>ras</u> gene is oncogenic in adult mice.

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If we looked at the oncogenicity of polyoma virus DNA, polyoma virus DNA, of course, is infectious in mice, but in hamsters it is not infectious. It causes tumors by interperitoneal, subcutaneous routes, using either super clonal DNA or linear DNA. As you can see, it's tumorigenic.

This is the viral DNA isolated from the virion. If you clone the polyoma virus DNA and inject in newborn hamsters, it is also tumorigenic.

Now, the minimum amount of DNA they found activity was .2 micrograms of the linear DNA, and gave it to 22 percent of the animals. So if we use that figure, it can conclude that .2 micrograms, which is about two times ten to the ten molecules of polyoma virus DNA is oncogenic in newborn hamsters.

So then we're going to move on to what data are available on the infectivity. Now, I'm going to talk about several studies on retroviruses, and in the first ones I'm going to talk about is work done by three groups over the years, using Simian immunodeficiency virus through various routes and establishing infection in various animals, and the number of genomes, for infection.

In the bovine leukemia virus using ID route with either a facilitated Achiheinig (phonetic)

1 lipid facilitator DOTAP in the Caprine (phonetic) arthritis encephalitis virus in goats, again, and 2 feline leukemia virus and feline immunodeficiency 3 virus. 4 5 Now, the only reason for presenting this 6 slide is, although you probably can't see it, is that

number of genomes for infection with these disparate systems and different routes of inoculation is remarkably similar.

The highest number of genomes required is about two times ten to the 13, and the lowest is one times ten to the 12 in this system. So it's somewhat remarkable. Even though a dose response study was not generally done, with the possible exception of the Purcell group, it is quite remarkable the number of genomes or molecules of these proviral clones.

And yet to establish an infection, we're very similar within about a 20-fold range. The dose response study was done by Portis and colleagues with a murine retrovirus using 19, 3.8, .38, and .038 micrograms intraperitoneally and found that micrograms intraperitoneally was the last of the level to establish an infection.

Interesting, they also found that super core DNA was unable to establish an infection, whereas

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linear DNA established an infection very well.

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That result disagrees somewhat from other

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people's, but probably reflects the system as well as

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the clones.

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So the summary then of the infectivity with cloned viral genomes, viral DNA coming from the retroviruses, 15 to 500 micrograms intramuscular injection, and so that if you calculate the number of genomes, it's about one times ten to the 12 to 2.3 times ten to the 13 genomes required for infection.

Polyoma virus, in contrast, from five time ten to the minus five micrograms or one microgram variously, and the minimum estimate then is 1.3 times ten to the seven genomes. So if we can conclude from that the infectivity of different retroviral DNAs is quite similar, quite surprisingly perhaps is similar, and depending on the route of inoculation, micrograms can be infectious for retroviruses, but the infectivity of polyoma virus DNA is higher than approximately 50 picograms or .05 nanograms sufficient to establish an infection.

So it may be informative to compare the oncogenicity and infectivity where it's possible. for polyoma virus DNA, that's the only one where we can, in fact, compare directly the oncogenicity.

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Point, zero, two micrograms is the minimum amount of DNA that was tested that was oncogenic, and that's about 3.6 times ten to the ten genomes infectivity, however. The infectious dose 50 is 1.3 times ten to the minus four micrograms. So this is about 2.3 times ten to seven genomes.

So there's about three orders of magnitude difference with the infectivity of polyoma versus the oncogenicity, and SV40 is about an oncogenicity of a microgram. Retrovirus, the infectivity, again, is 15 to 30 micrograms. It's about one to two times ten to the 12 genomes. V-src was two micrograms at about two, two and a half times ten to the 11 molecules. Inactivated ras, ten micrograms, is nine times ten to the 11 molecules.

So we went in the literature to find out what evidence, what data are available for intranasal inoculation. Unfortunately there is almost none in terms of quantitative estimates. So the only one we could find was looking at this study by Timer and colleagues in Japan several years ago by using DNA vaccine administered intranasally.

So this was a clone gene of the hemagglutinin gene from influenza. It was inoculated into mice via the nasal route. Different amounts of

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DNA were administered in PBS, and after four weeks, a 1 second intranasal inoculation of 1.5 micrograms was 2 given as a good. 3 4 So down here we can zero micrograms, 0.2 micrograms, .4 micrograms, and .6 micrograms, and they 5 measured two responses, hemagglutinin inhibition assay 6 7 and mucosal IgA to the vaccine, and after the primary inoculation, nothing was found except for the six 8 9 micrograms. There was a weak response. However, after the second inoculation, 10 there was nothing. No priming inoculation was given 11 12 demonstrating that this 1.5 microgram secondary -- the 13 boost was not immunogenic. However, for all the 14 levels of DNA, there was a substantial boost. 15 So what this means to us is that even 0.2 micrograms of DNA, in other words, 20 nanograms, 16 17 administered intranasally elicits immune response since the secondary inoculation boosts this response. 18 19 Therefore, we're concluding that 20 nanograms of DNA can be biologically active when administered intranasally. 21 22 So the other route that we're considering 23 is the oral route. Now, there aren't that many 24 studies of this either, but Malcolm Martin and Mark Israel's group fed mice polyoma virus DNA between one 25

or 0.5 micrograms, and zero out of 25 became infected
with one microgram and zero out of 30 became infected
with 0.5 micrograms.

However, if you stick a gastric tube through, you can get some infectivity, but how this is doing this is through damage of the gastric tube in mice is not clear, but the important point is though that it's unlikely that small amounts of DNA will survive passage through the stomach intact.

Now, the next study that looked at oral routes is a study by Duffler (phonetic) and colleagues, and they fed a phage DNA -- it's a 7.25 kb phage DNA -- large quantity DNA and followed its fate to see whether anything did escape through the stomach, and they did find DNA in feces, blood, small intestine, and large intestine in the leukocyte population.

But the size, as you notice, is 100 base pairs to 1,700 base pairs, whereas the original genome was 7,000 base pairs. So it's clearly getting degraded during passage through the stomach.

They did find some in the blood, 194 base pairs to 976 base pairs, and perhaps interestingly, they also found about a 700 base pair fragments in blood, and they also found it integrated in 0.1

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percent of the cells.

However, these cells did not survive, and these cells came from the peyer's patches and from the spleen.

So conclusions that small amounts of DNA can, in fact, pass through the stomach, but no full length DNA was detected, and DNA can be found integrated into the mouse chromosome in a few percent of the cells. So you have to feed large quantities of DNA.

So now we can go on to try to get some estimate of risks. We'll first consider the oncogenicity.

The cloned activated oncogene, for a single dominant activated oncogene per cell, one microgram of DNA has 152,000 oncogenes, and this figure comes from the DNA in 152,000 cells that's one microgram or one microgram carries 152,000 cell equivalents. So that's where that figure comes from.

We know from our previous studies of the literature and for polyoma virus DNA about 3.6 times ten to the seven genomes or molecules are required for oncogenicity, and the activated <u>ras DNA</u>, 9.1 times ten to the 11 molecules are required for oncogenicity.

Therefore, the number of tumors expected

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from one microgram of residual DNA with a single dominate oncogene per cell is between 152,000 divided by this. It gives you a value of 4.2 times ten to the minus six for polyoma, and a value of 1.7 times ten to the minus seven for activated ras, and if you take that the other way, therefore, for ten nanograms of DNA, not one microgram, the probability of an oncogenic event is between one and two times ten the five or 200,000 for polyoma virus, and one in ten -- six times six million for the inactivated ras. So that's where the calculation comes from.

For infectivity, we've looked at polio virus DNA, and viral genomes required for infectivity is ten to the seven. So it's ten million.

The probability of an infectious event using ten nanograms of mammalian DNA with a single copy of polyoma virus DNA is one in 7,000 events, one in 7,000 events. This is rather high.

This also points to the fact that it is very important to be able to determine that an adventitious agent, such as polyoma, not just polyoma; other adventitious agents are not present in the cell substrate since infectivity is a very easy -- much easier than oncogenicity.

For the retrovirus cloned proviral DNA,

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the number of viral genomes we've determined is about four times ten to the 11, as a minimum, and the probability of an infectious event for ten nanograms of mammalian DNA with a single provirus is about one in three times ten to the eight, one in 300 million.

So it's much more difficult for this. OF course, these values all change if there are more than one copy of the oncogenic agent or of the retroviral

So we can draw some conclusions from this. Infectious risk of DNA can be more important than oncogenic risk, and as I said, therefore, it's very important to determine what level of adventitious

For the IM, intramuscular and subcutaneous routes, ten nanograms of DNA provides an estimated risk for the polyoma virus DNA, one in about 200,000 for an oncogenic event and about one in 7,000 for an

For the more likely events of cellular DNA now, cellular activated ras, there's one in six times ten to the sixth for an oncogenic event, and for proviral DNA for an infectious event, there's one in 300 million, three times ten to the eight.

For the IN route, the intranasal route,

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ten nanograms of DNA as I've shown you has an estimated safety margin of one in ten to the sixth for what I'm quoting an expression event. This is because you can detect the activity of that DNA. However, that is not an unoncogenic event. That's just what we can detect by immunological means. So that's the best one we have.

And for an oral route, one micrograms of polyoma DNA administered orally is not infectious. So, therefore, ten nanograms by arithmetic calculation with one copy of polyoma DNA, the safety margin is at least one in ten to the eighth.

However, it's not all bad for the risk. There are some mitigating factors for DNA. As you probably all know, the uptake, expression, integration are inefficient processes. Integration requires cell division, and so therefore, not all cells in the body, of course, are dividing. In intramuscular routes, the myocytes, and muscle cells have a very low division rate, and integration is required for maintenance of DNA in the absence of the replicating system.

The degradation of DNA vaccine manufacturer procedure usually requires degradation of the DNA, and of course, DNA is degraded <u>in vivo</u>. It doesn't exist for very long <u>in vivo</u>.

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So if the DNA is degraded below the size of a gene, then that increases its safety.

There's a host immune response to transfected cells, as Dr. Cook just mentioned. So even if the DNA does get in and express these cells in an immunocompetent individual, it will likely be removed.

What also Dr. Cook talked about eloquently is the multi-step nature of carcinogenesis. We now know that it's not just a single event. So even if an activated <u>ras</u> did get into that cell, that cell would not be established as a tumor right there. There are many more steps that have to proceed before carcinogenesis can occur.

And, of course, we all know that transmission of human cells in culture is much more difficult than transmission of rodent cells, and Dr. Alex van der Eb discussed some of these issues earlier on.

So what are we doing at CBER to try to help to address this? Well, Dr. Hughes mentioned that we have now initiated a study. We have finally got some funding from the FDA, and in a collaborative study with CBER and the National Cancer Institute and the Center for Drugs, we're going to study in a

systematic way to develop numbers that have been 1 2 alluding everybody for 40 years. 3 These issues come up every five to ten years, and still nothing is being done. 4 5 So we're trying to develop sensitive animal models to detect oncogenic activity of DNA. 6 The models that have been heretofore used have not 7 been systematically developed and are not very 8 sensitive. So what we're going to try to do is test 9 newborn NIH Swiss and C57 Black 6 mice, the athymic 10 11 nude mice, and K6/ODC mice, which is a constitutively promotive mouse line, and we're going to use them 12 13 making the activated ras genes. 14 our colleagues, and we're 15 grateful to have John Coffin and Don Blair and Steve Hughes from NCI and Frank Sistair (phonetic) from CDER 17 to help and advise us on these studies. The other assay we're trying to do with CDER is to get some quantitative estimate of DNA infectivity, and we want to develop quantitative in vitro assays first, of course, to affect the infectivity of proviral DNA in retroviruses. not known what level of proviral -- levels of provirus that is in mammalian DNA are infectious. Howard Tammin (phonetic) has shown that in

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If you could go into a system where some of those things are already present, if you would -- I don't know -- give us a better readout.

DR. PEDEN: Yes, you're right. I didn't mention it, but, in fact, the idea of a first hit and then you give to a vaccinee and then subsequently you get additional hits is a worrying issue.

We don't know how stable those initial hits are. If you give it to a cell, where that cell eventually gets lost, you don't retain that so-called activated state in the cell. We don't know that.

As for the models, the ODC mouse is a carotin 6 promotor that drives olefin de-Kalb oxylase (phonetic), and that's been used in carcinogenesis activities because it's constitutively what they call promoted. So if you add an initiator such as <u>ras</u>, it's a very sensitive model on the skin, of course, because --

DR. GRIFFIN: So it does provide a more sensitive --

DR. PEDEN: So it does provide that. We discussed with Steve Hughes and John Coffin and Don Blair about which mice to use. We're thinking about using the heterozygous p53 mouse, and perhaps Steve or

the others could comment on that, but we decided in the first run through to do these mice. I think the homozygous p53 knockout was generally felt, I think, by Steve just to have too much background.

And so we're trying to balance it. I'm glad you do point out the importance of doing these quantitative studies. I don't want to lessen what's done in the past, but they were never intended for that role, and of course, the infectivity in monkeys is an incredibly expensive thing. So there isn't any information on that.

So I don't know whether Steve wants to comment on the particular animal models or Don or John.

DR. BLAIR: No, I think we chose, you know, sort of a reasonable spread of different strains of mice which might show up some strain differences. The nudes as an immunosuppressed, perhaps more sensitive mechanism, and as you say, the problem with some of the p53 knockouts and things is as you increase the sensitivity, you increase the background, and you may obscure things. So we went with something where we felt the background wasn't going to be insurmountable and we should see positive.

DR. HUGHES: I also would add that I don't

will necessarily be the only things that will ever be done, and I think that based on what we would like to do is to have what we hope is a reasonable preliminary survey and see, in fact, the degree to which we get response with relatively lower amounts of DNA.

And depending on the outcome of these experiments, I think we would be better prepared to go back and actually plan and propose a second round in which, depending on what the outcome is, something like a p53 heterozygous animal might be employed.

But because they're actually, as you point out and Dr. Peden has pointed out, there's relatively little really quantitative assessment, we thought we'd make what we intend to be the first attempt to do that, and then based on what we learn from that, try and go ahead in a reasonable fashion and maybe expand this to other model systems.

ACTING CHAIRMAN DAUM: This is, again, an issue that I think we should encourage everybody to raise in this afternoon's session when we're talking about the big picture issues with respect to this approach.

I'd like to focus now on Dr. Peden's presentation though, and Dr. Aguilar-Cordova is next

1	and then Dr. Faggett and Dr. Moulton.
2	DR. AGUILAR-CORDOVA: Yes. Given the
3	numbers that you presented and there may be various
4	oncogenes in any one cell or activated genes or
5	whichever, in your interpretation would it make a
6	significant difference whether it was a spontaneous
7	tumorigenic cell line versus a designer cell line in
8	which one of the many events might be known?
9	DR. PEDEN: So now are you saying the
10	oncogenes from a spontaneously transformed cell is
11	studied? I mean, these are cloned oncogenes injected.
12	DR. AGUILAR-CORDOVA: This is like the
13	worst case scenario. What I'm saying is based on your
14	statistical analysis that you presented
15	DR. PEDEN: Well, I wouldn't grace it with
16	statistical analysis, but
17	(Laughter.)
18	DR. AGUILAR-CORDOVA: Or statistical
19	deductions. I guess the general question would be
20	would, say, in the presentation we heard earlier from
21	Dr. Lewis, do you think that based on that would A549
22	residual DNA pose any significant different risk than
23	293?
24	DR. PEDEN: I think there's not universal
25	agreement, I think, among us about that, but I think
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the general feeling is yes. The main reason, you don't know what the oncogenes were that brought about 2 the A549 phenotype. So that's one thing. 3 And also the other issue about A549 is you 4 don't know what's in there. Is there an oncogenic 5 agent that brought about that? 6 7 So I think in general terms there is a big difference between a cell that you know hits passage 8 history and you transform it by another mechanism. I 9 10 think that's why we are calling these designer cell substrates and we call them at low risk, such as A549. 11 12 So there's two reasons for that. The both 13 I've just said to you. I think for the study what we want to do 14 15 is, in fact, determine ultimately the most sensible animal model where we can look at genomic DNA, DNA 16 from A549, from PER.C6, to see where that can induce 17 an oncogenic event. So that's really where we're 18 going at, although I didn't, in fact, say that. 19 to see whether you can, in fact, detect an oncogenic 20 21 event using DNA from such things as A549 cells. 22 That's where we want to go, but we need to 23 develop these animal models, sensitive animal models 24 for that reason. 25 Thank you very ACTING CHAIRMAN DAUM:

much.

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Dr. Faggett, and then Dr. Moulton, and then we will adjourn for lunch.

DR. FAGGETT: Now, what prompted the change from the WHO requirements for residual cell substrate DNA levels from 0.1 nanograms in 1986 to the ten nanogram level in 1999? Was there some --

DR. PEDEN: I think as we get more and more information, we do these experiments. See, more information came, more experiments were done. You may think they were deficient, and we may agree with you, but we have more knowledge and more data and also more clinical experience with some of these cells, such as Vero cells.

So I think it's a combination of all things, but I wasn't at that meeting. So maybe Phil and Andrew were at the meeting. I don't know what specifically lowered except for these numbers that I just presented to you.

But, Phil, do you want to say anything?

DR. KRAUSE: Yeah, maybe I could just comment. I think an important thing to remember is that these meetings that have gone on in the past, in general, have been talking about the older kind of cell substrates, and here we're talking about newer

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And so I don't think that the idea of using an A549 cell, for instance, was specifically considered as one of the things that was going into raising that limit from 100 picograms to ten nanograms at that meeting, but instead the idea was to consider the kinds of cells with which there was much more substantial experience in vaccine and other biological production.

ACTING CHAIRMAN DAUM: Thank you very much.

Dr. Moulton.

DR. MOULTON: Yeah. This is a great start, I think, towards the full risk analysis. think a full risk analysis would have probability distributions associated with every one of assumptions there. There's a lot of assumptions we're making. For example, the potential mitigating factors going to reduce the risk. are There's other assumptions that could increase it, and a couple of people have already mentioned the problem of the low dose extrapolation, which could affect things by a couple of orders of magnitude.

And I was wondering if there's a plan even before we get the result of these future experiments

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to try and put some probability distributions on 1 there, every one of those assessments, and also a more 2 general question of how does this risk analysis fit 3 into the general risk management program at FDA. 4 5 It's one thing to estimate or calculate a 6 risk, estimate that point estimate, and if you're 7 lucky get some kind of tolerance interval around it. It's another to actually manage the risk once it's 8 been estimated. 9 Well, the way we manage the 10 DR. PEDEN: 11 DNA risk is to require that the levels are low. mean that's the simple way to do that. 12 13 As for the probability, many people have 14 15 16

tried to make estimates, probability estimates at all of these steps, and i can give you lots of references where that was done. I don't find that that's satisfactory nor useful because there isn't -- I mean, the probability of integration event, the probability of out-take of DNA, that's all those events you're talking about, and that's the evidence.

The consideration of those is all done from in vitro studies. They've been transfecting DNA in culture and measuring events, and so to me that's not very satisfactory to do that, to extrapolate from what goes on in vitro to what goes on in vivo.

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That's why I think we need the in vivo 1 data to try to get real numbers as opposed to these 2 estimates. 3 ACTING CHAIRMAN DAUM: This is a topic that I think must be also returned to this afternoon. 5 So perhaps, Dr. Moulton, you will remind us that we 6 need to address this and bring it up again. 7 I'd like to thank Dr. Peden at this point 8 very much for another fine presentation that helped us 9 10 Dare I say you provided us a great deal of 11 thought for food. 12 We will now break for lunch. It's 12:45 here in the Eastern time zone, and we will reconvene 13 1.4 at 1:45 exactly. 15 Thank you. 16 (Whereupon, at 12:45 p.m., the meeting was recessed for lunch, to reconvene at 1:45 p.m., the 17 18 same day.) 19 20 21 22 23 24 25

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# A-F-T-E-R-N-O-O-N S-E-S-S-I-O-N

with the adventitious agent issue this afternoon.

introduce us to the topic.

have three or four presentations on that, and we'll

begin with Dr. Philip Krause from FDA who will

effective public health interventions ever devised.

Vaccines, however, are very dependent on public

confidence in their safety, and that's really for

several reasons, but one of the most important is that

vaccines are generally given to healthy people, many

of whom may never be exposed to disease which one is

trying to prevent, and another reason, of course, is

the need in many cases for many vaccines to generate

high enough immunization rates to get some level of

herd immunity, without which one doesn't get maximal

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(1:49 a.m.)

We are in

Okay.

Welcome back from lunch, and we'll begin

DR. KRAUSE: Vaccines are among the most

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ACTING CHAIRMAN DAUM: afternoon session.

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And of course, since many vaccines are given to healthy children, these issues are even more important.

An important component of public

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vaccine benefit.

confidence in vaccine safety, of course, is their freedom from adventitious agents. So what is an adventitious agent?

Well, for the purposes of this discussion at least we're defining an adventitious agent as an infectious agent that is extraneous to the product, and so a vaccine strain obviously is not an adventitious agent, but something that's carried in from the outside or carried in with the cell substrate, is potentially an adventitious agent, and an obvious goal then is to insure that final products don't contain adventitious agents.

Now, there are two additional points I want to make, and just because I say final products here doesn't mean that intermediate products should contain adventitious agents. In fact, all through the production process it's important to keep adventitious agents out of vaccine manufacture, and the other thing which we've spent some time talking about already is trying to define in some quantitative way the terminology of "should not contain."

So any negative result obviously needs some quantitation associated with the sensitivity of the testing, which allows one to exclude that something is there.

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Now, in general the OVRR approach to adventitious agent issues has included identifying potential issues, including theoretical ones, discussing these issues in public, and Dr. Lewis described the two previous Advisory Committee meetings at which this came up, as well as the public international meeting on cell substrates, and of course, this meeting is another component of that, making decisions based on the best available science and, in addition to that, insuring that any potential issues are known and understood by research subjects and investigators.

So today we're going to be talking about two potential adventitious agents or types of potential adventitious agents. One is transmissible spongiform encephalopathy agents, and the other is viruses.

Directly following this brief introduction, Dr. Sue Priola will talk in some more detail about TSE agents, and then after that, I will provide some general principles of adventitious agent testing, as well as a more detailed discussion of virus as potential adventitious agents.

So to introduce Dr. Priola's talk, I'm just going to go over a few things related to our

consideration of transmissible spongiform encephalopathy agents and cell substrates. And the kinds of issues that can come up that might increase the risk that a cell substrate could harbor such an agent.

And I guess one possibility is that because some transmissible spongiform encephalopathy diseases, such as Creutzfeldt-Jakob disease and Gerstmann-Straussler syndrome are at least in part genetically determined. The question arises of whether a cell substrate which contains mutations that are associated with these types of diseases might then harbor a greater risk of either having TSE agents in it or, if exposed to TSE agents causing infection or being infected with them and thereby transmitting them to a vaccine recipient.

The second question, which is a general one which comes up and which this committee has addressed before to some degree at least, is the consequence of potential exposure to serum from countries where bovine spongiform encephalopathy or the risk of BSE exists.

And this is an issue which has come up in the context, in particular, of cells which have been in laboratories in the early to mid-1980s at a time

before this risk was as widely appreciated as it is now.

And then, of course, there are other factors that could theoretically increase the risk of TSE infection of cell substrates. One of these is PrP expression levels. It's been shown at least in some systems that cells that express higher levels of the PRNP gene that encodes PrP, that the risk of those cells being able to propagate TSE agents is higher, and then also there's the question of whether cells of neuronal or retinal origin might also have greater risk of at least propagating TSE agents if exposed or of containing them in the first place.

Now, there are a couple of neoplastic cell specific notions that are also TSE related. One of them is that because neoplastic cells often have some associated genomic instability, whether, in fact, that genomic instability might itself induce mutations in the PRNP gene, which might then have some consequence on the ability of those cells to harbor TSE agents.

And then also, it appears that at least in some cells the ability of a cell to undergo apoptosis helps to prevent infection with TSE agents in them, and because neoplastic cells have in most cases at least some apoptosis pathways abrogated, the question

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then is whether a neoplastic cell might a priori be a little bit more susceptible to infection with TSE agents.

So at this time, we're following the approach to TSE issues that's outlined on this slide, and that is where possible to determine the family and medical history of the cell donor with respect to TSE risk factors; to ask that the PRNP gene be sequenced; to ask that Western blots be performed to look for the presence of protease resistant PRP; and to insist that they be negative, of course; and to determine if exposure to fetal bovine serum from countries with BSE or with the risk of BSE could have occurred, and then if some possible exposure to questionable fetal bovine serum has occurred, to perform a risk assessment based on the dilution factor from the time of that exposure, based on the assumption then that the cells cannot support replication of a BSE agent.

There are several evolving ideas which we don't think can be applied right now, but which may be applicable in the future. One of them is if possible exposure to fetal bovine serum of unknown origin has occurred, to document that the cells cannot support replication of the bovine spongiform encephalopathy agent.

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And this time I don't believe that there are any cells in tissue culture that have been demonstrated to replicate the BSE agent.

To evaluate the level of PrP expression, which although it is generally accepted, I think that cells that express higher levels of PrP might have a greater risk of propagating TSE agents if exposed. The question then also comes down to defining what exactly would constitute a safe level and understanding in a way that can be applied quantitatively what kind of controls would be necessary to use such an assay in a regulatory setting.

To evaluate for the presence of TSE agents by animal inoculation, and as many of you know, the monkey, the primate models take a very long time to carry out, but there are promising transgenic mouse models which could potentially be used after they are appropriately validated to get more rapid answers to these kinds of questions in an animal model, and of course, the general principle that once new assays for detection of TSE agents become available, to introduce them for cell substrate testing as soon as is feasible.

So with that, I would like to stop here.

1	I'm happy to take any questions, but if there aren't
2	any, then we'll go on to Dr. Priola's talk on TSE
3	agents as an issue in the use of neoplastic cell
4	substrates, and I think she'll concentrate, in
5	particular, on issues associated with TSE agents in
6	cell culture.
7	ACTING CHAIRMAN DAUM: I think we'll go
8	right on if that's okay because we'll have a change to
9	ask questions.
10	DR. KRAUSE: You'll get another crack at
11	me.
12	ACTING CHAIRMAN DAUM: That's right.
13	Thank you.
14	Dr. Priola here? There she is.
15	DR. PRIOLA: I would just like to thank
16	can everybody hear me? to thank Dr. Krause for
17	giving such a terrific introduction. He left me with
18	a few things to talk about, but he hit really on every
L9	major point that is going to be an issue in what I go
20	over.
21	What I want to do is basically give a very
22	brief introduction, basic introduction to TSE
23	diseases, and then describe using experimental
24	examples what we know about the difficulties
25	associated with passaging TSF infectivity into tiggue

culture. 1 2 So do you have the presentation up? There 3 you go. I think as we all know that TSE diseases 4 are long term transmissible, degenerative diseases. 5 Have you frozen up there? It's coming on. 6 It's the good at explanation point. Oh, no, there it 7 8 goes. 9 (Laughter.) 10 DR. PRIOLA: I want to wait until this 11 Okay. There we go. Thank you. comes up. 12 All right. So the TSEs are slow, fadable, and transmissible brain diseases that affect a variety 13 14 of mammals, including, of course, humans. Creutzfeldt-Jakob disease in humans is the prototypic, 15 one of the prototypic TSEs, scrapie in sheep, and of 16 17 course, BSE in cattle. 18 One of the things about the TSE diseases is there's this phenomenon called the species barrier 19 20 to infection, and it's this that has caused all of the 21 concern when it was realized that SE from cattle 22 apparently has crossed over and caused a new form of 23 CJD in people in the U.K. 24 And usually the species barrier is quite 25 strong in these diseases, and so it was unexpected

that cattle BSE would necessarily pass into people. 1 And, of course, it's this concern that now 2 3 have BSE in cattle populations contamination of bovine products used in tissue 4 culture could then passage that TSE infectivity into 5 tissue cultures using those products. 6 Experimentally, the primary systems are TSE infections of mice and hamsters, and most of what 8 9 I'm going to talk about today deals with that. 10 It's important to remember that within the TSE there are different strains of TSE agents. 11. has to 12 there be some caution in interpreting 13 experiments using one strain of agent and one species 14 of animal and extrapolating those data to other 15 species and other strains. I'd like you to keep that 16 in mind. 17 Infection, of course, is believed to be 18 primarily from ingestion or inoculation. It's an extremely long disease course. 19 It can take from 20 months to decades to appear. It is always fatal. 21 There are no preclinical diagnostic tests available. 22 There's no effective post or preclinical treatments 23 available. 24 If that weren't bad enough, it turns out 25 that we're still not entirely clear what the exact

composition of the infectious agent is, but the fact
that it's unusually hard to kill and that there's no
viral or bacterial association with TSE infectivity -good Lord. Can I go back? Good luck. Okay -- led to
the hypothesis that, in fact, it was an infectious
self-replicating protein that was responsible for
these diseases.

In the early '80s, a protein was found that was very closely associated with TSE infectivity in Stan Prusiner's lab, and that protein was called prion protein, or PrP, and I think as most of us are aware here, PrP is, in fact, a normal host cellular protein. It's something that's expressed in all mammals, and it's expressed almost ubiquitously. It's very difficult to find a tissue or cell that does not express PRP.

In its normal form it's sensitive to digestion with cellular proteases, and for that reason I'm going to call it PrP-sen for protease sensitive.

As I said, it's almost ubiquitously expressed, and it's soluble.

During TSE disease, the normal form of PrP, which is shown here -- it's a cell surface glycoprotein -- is converted to an abnormal form that is now partially resistant to proteinase K, and the

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partial proteinase resistance comes about because when 1 you expose this molecule to proteases, what happens is 2 instead of digesting everything away as it would if it 3 were just PrP-sen, it basically can only clip away 4 part of the end terminus, and you see this very 5 characteristic side shift down like this, and that is 6 what we define as PrP-res or protease resistant PrP. 7 8 This form is now insoluble. It's heavily aggregated, and it is completely TSE specific. the marker that everybody looks for. If you find PrP-10 res, by definition you have a TSE disease. 11 12 It's found primarily at high levels in the 13 CNS animal models it's found 14 lymphoreticular system as well. 15 Ιt different has confirmational 16 structure than normal PrP. The normal PrP 17 primarily optihelical. When it gets converted to the 18 abnormal form, it becomes beta sheeted structure, and it's believed that it's this difference, it's this 19 20 change in confirmation that accounts for the different 21 properties of these molecules. 22 while there's still 23 within the field about the role of whether or not PrP-24 res is the infectious agent, the role of PrP in TSE 25 diseases is absolutely undeniable. You need normal

PrP for infection. Mice that don't express this gene do not get sick following exposure to TSE agents. 2 3 Mutations in this molecule can strongly influence disease susceptibility and species barriers 4 infection. 5 tò PrP-res, the abnormal form associated with toxic events in the brain and is, of 6 7 course, always associated with infectivity. So in terms of human TSE diseases and the 8 9 dangers involved, not dangers, the possibility involved with deriving neoplastic tissue culture from 10 human cells, there are three groups of human TSE 11 12 diseases. 13 There is the sporadic form known as Creutzfeldt-Jakob disease or CJD. There's no known 14 15 exposure to TSE infectivity. This form of the disease 16 is not associated with mutations in PrP. It accounts 17 for the vast majority of TSE cases. It's extremely 18 It happens annually worldwide at an incidence 19 of about one case per million people. 20 There's also familial forms of the human 21 TSEs. These include familial CJD, Gerstmann-22 Straussler-Scheinker syndrome. These forms are also

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not associated with any known exposure to infectivity.

They are, however, associated with mutations in the

PrP molecule. They are even more rare than this radic

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(phonetic) disease with about one to ten cases per 100 2 million people. There are very few families in the world that have been identified that carry these 3 4 mutations. 5 Finally, I have lumped together here under the term "infectious TSE" infections which are a 6 result of exposure, a known exposure or presumed .7 exposure to infectivity, and these include iatrogenic 8 cases that arise 9 CJD following exposure to contaminated medical instruments, and of course, 10 variant CJD in Great Britain. 11 12 Is there a problem? You can't see the 13 I'm not speaking clearly enough for you to see the slides? Sorry. I'm glad you told me. 14 15 All right. So where in the heck was I? 16 Okay. So in terms of it being an issue in cell culture, the problems are can you infect tissue 17 culture cells easily with TSE agents. 18 19 You're going to have to wait a minute. I've got to reorganize myself. 20 21 And as alluded to by Phil, there are three 22 basic issues. First of all, development of new cell lines from CNS tissue of patients who are potentially 23 24 contaminated or who are potentially infected with

Creutzfeldt Jakob disease or a derivation of cell

PrP

two

from an individual carrying familial 1 lines mutations, and I'll address both of these issues. 2 There's also what has become a concern 3 particularly in the last couple of years whether or 4 not tissue culture cells exposed to bovine derived 5 products could become TSE infected, and I'll address 6 7 this and the susceptibility factors that might be involved. 8 9 And finally, given these 10 possibilities, what is our ability to detect TSE infectivity in these cells? 11 12

All right. So the first thing I want to address is it, in fact, possible to derive persistently infected TSE infected cultures from immortalized cell lines derived from human cells, and these experiments were all done many years ago, 20 to 30, almost 40 years ago now.

these experiments essentially tissue was taken from the brain or spinal cord of CJD or Kuru patients who had died of CJD or -- this thing is extremely sensitive. I've got it. It's okay -had died from CJD or Kuru, were maintained in tissue culture for over 300 or up to 300 days, and then the presence of infectivity assayed for by bioassay in primates, and they could always detect infectivity.

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Similar experiments were done in mouse and sheep models of scrapie. The point I want to make here is that even though you could maintain infectivity if you just kept the cells alive, the minute you start passing them, you lose infectivity.

Okay. So after one or two passes in the human system, you lose infectivity, and you can't recover it upon further passage. You can, however, derive immortalized cell lines infected with TSE agents and other systems because this approach did work in both the mouse and sheep system.

So even though it didn't work for human, theoretically it's been shown to work in at least two other system. So the possibility exists.

What about familial TSE diseases? There are a wide variety, up to 20 mutations in the PrP gene that have been associated with familial TSE disease, and they're spread throughout the molecule, and we don't know exactly how they lead to disease, but the idea based on the protein only theory of the TSEs is that these mutations lead to altered biochemical properties of the normal PrP molecule. It then spontaneously converts over to the abnormal form, accumulates, and causes disease.

So what is the evidence that this actually

happens? Because, of course, this is the concern if you should immortalized either cell line derived from an individual carrying a familial mutation. Does this actually happen? And in my laboratory over the last few years, we've been working with PrP mutants that involve insertion of extra copies of an octapeptide repeated motif, and when you look at these mutations, so what we've done is basically taken a PrP molecule and just added more and more numbers of these repeats and asked about the properties. I hope you can see this. What you find is

that if you look at two of the cardinal properties of PrP-res, the abnormal form, protease K resistance and aggregation, you can see the red line here. As you increase the number of repeats, you do, increase the relative protease resistance of PrP.

To put it into perspective, the green line represents the abnormal form PrP-res. At the level of PK where all of the protein is destroyed, PrP-res hasn't even been touched. So this molecule is not PrP-res by this definition.

Similarly, when you look at the amount of it aggregated, yes, as you increase, as you have these mutations, the molecule begins to aggregate more.

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still does not do so to the same extent as PrP-res.

So while the biochemical properties of these molecules change and they begin to acquire some properties reminiscent or res, they are not PrP-res, and believe me, we've tried. We cannot get them to spontaneously form PrP-res.

All right. So this is all in tissue culture. What about <u>in vivo</u>? If you take these cells and stick them into animals, you also don't get TSE disease. Okay? So just sticking in the cells.

If you make transgenic mice over expressing these mutations, so try to mimic what happens in people, what do you get?

And what I've done here is really very briefly summarized experiments done by several labs where they basically inserted human associated familial TSE mutations in to the PrP gene, overexpress them in mice, and then assayed them for disease, and a couple of points I want to make here.

Number one, you need overexpression to get any sort of neurological disease. You don't need the mutation necessarily because without the mutation and over expression, you still get neurological disease.

Okay. So overexpression itself is sufficient to cause symptoms in the animal.

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at these mice, they did show signs of neurological dysfunction, no TSE disease, and they certainly didn't passage infectivity. There's this one GSS associated mutation where this was overexpressed in animals, had the neurological disease, and there may have been some transmission into limited -- in a limited way into other mice.

so this is the one instance where there may be some evidence that this mutation could induce disease. When you look, however, at this same mutation in a context where it is not overexpressed, which is much more analogous to the human system, all right, so this is an instance where mouse was made where the mutation was put into the PrP gene and that gene put right back into the correct spot in the chromosome at the correct copy number. You don't get neurological disease. You don't get TSE disease, and you don't get infectivity, and the laboratory that did this the last I heard had looked at 800 and some mice. So they're looking very carefully.

So there's no really strong, convincing evidence that mutations, familial TSE mutations will generate spontaneous TSE disease in tissue culture or in transgenic mouse models.

Now, what about infection of tissue culture cells, particularly in regard to exposure to bovine derived tissue culture products?

What are the susceptibility factors? How much infectivity do you need? How is easy is it to get an acute versus persistent infection?

And what I've tried to do on this next slide is summarize all of those points in a little model, and what I've shown here is each of these black boxes represents what I think is a major block to persistent TSE infection of tissue culture cells.

First of all, you need extremely high -you need high multiplicity of infection. So routinely
what we used to do is brain material, high titered
infectious brain material. This automatically calls
into question whether or not the very low amounts of
TSE infectivity that have been hypothesized to be
carried around in blood would be sufficient to
overcome this barrier where you need a whole lot of
agent to get a little bit in.

The second thing, even if some of this agent gets in that has to be considered is, first of all, does the cell express PrP-sen theoretically any cell that expresses normal PrP is susceptible. So if it expresses PrP-sen, it's got the right sequence. So

say this is bovine, BSE, PrP-res, and this is human 2 PrP-sen. Right away you've got a problem. The sequences of these genes do not match, and that will 3 be an issue in terms of the abnormal form reacting 4 with the normal form and causing propagation of the 5 abnormal form. 6 7 Okay. So let's say all of this is okay. 8 This matches that. There's enough PrP there. 9 sequence is the same. It's folded appropriately. This process occurs. 10 You've passed the second 11 barrier. 12 Then comes the third one, and this is a 13 big one, and that is that as you start to passage these cells and dilute out whatever cells might have 14 become infected, you lose infectivity. 15 16 Even if you get an infected monolayer, in 17 many cases there are as few as one percent of these cells infected. All right? And the consequence of 18 19 that is that if you look at an infected cell layer like this for PrP-res, by current techniques you're 20 2.1 probably not going to find it. Okay? 22 So while the presence of PrP-res is a very 23 good indicator that the cell is infected, the absence 24 does not mean it is uninfected, and that's what I want

to show here. This is an experiment I did years ago

where we tried to infect cell lines that express relatively low levels of PrP-sen. We assayed at various times after exposure to an infectious brain homogenate.

We looked both for PrP-res, which we could not find, and we looked for infectivity in mouse bioassay, and we easily found infectivity, and at every pass and at equivalent levels.

Okay. When we did this experiment in cells that had been maintained in tissue culture a bit longer, so that these were relatively new cells; these had been maintained a couple of months longer; all of a sudden we couldn't infect them, and this gets back to the point that Dr. Hughes made earlier this morning that cells change over time in culture, and in this instance, it changed in a good way for you guys and a bad way for me because, you know, I want infected cells.

The other point I want to make, and Dr. Krause referred to this a couple of times in his talk, is this issue that if you overexpress PrP-sen, and this again is probably not something that should be a concern with the cell lines that are being used here, if you artificially overexpress PrP-sen, now you get a tissue culture line that is more susceptible to

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infection, and the idea is with more PrP-sen, you can accumulate more PrP-res, which you can then detect more easily, and you can infect more easily.

And our experience has been this is simply not true. It's completely unpredictable, and this is a piece of data that hasn't been published yet that just shows in neuronal and non-neuronal cells exposed to four different strains in this instance of mouse scrapie. So eight combinations. In only one case is there very clear evidence of formation of the abnormal form, and that's this one right here. All the others are negative.

And I could not have predicted. I mean I could have thrown a dart at this thing and predicted it just as easily as the result I got. So even overexpression, while it could be considered a susceptibility factor, you can't use it to predict susceptibility.

And just to summarize, all the different cell types that have been looked for susceptibility to TSE infection, there have been both neuronal and non-neuronal cells described that are susceptible to mouse and this bottom one here, the kidney epithelial, to sheep scrapie if they expressed the sheep PrP gene. All of these cells, including a couple of cell lines

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that were referred to earlier this morning by Dr. 1 Lewis, they're using vaccine production. These guys 2 3 have not been shown to maintain human TSE infectivity. 4 So there are a lot of nonsusceptible cell lines, and you can see a lot of these are human 5 derived. 6 7 Okay. So finally, to briefly mention what our ability to detect TSE infectivity in the absence 8 of detectable PrP, in the absence of bioassay, there 9 10 are now three approved and marketed tests for checking for PrP-res in infected animals, and these are being 11 12 used extensively in Europe at slaughterhouse levels to 1.3 look at cattle, potentially BSE infected cattle. 14 These tests are all maximized for high titer tissue, such as brain or spinal cord, and 15 they're only sensitive to PrP-res levels that you 16 17 detect shortly before or at clinical signs. So this is where you have a lot of infectivity present in the 18 19 animal. 20 This is not like tissue culture cells 21 where you have very little infectivity present. Okay? 22 There is another test that's coming 23 through in the approval process. This is the Adelphia 24 test, which may be a bit more sensitive than these

There are a couple of future tests that have

three.

not really been -- in particular, the capillary immunoelectrophoresis have not been rigorously tested, but have been proposed to be able to detect extremely low levels of PrP-res in tissues such as blood.

So the difficulties with all of these

So the difficulties with all of these tests, the postmortem test, they're based only on res detection. They're maximized to high titer tissue, and we haven't actually tested them against our tissue culture cells.

Their sensitivity, again, is not terribly

-- not as good essentially as bioassay in either the
same species or in appropriate transgenic mice. So
bioassay remains the gold standard test for detection
of TSE infection, and as Dr. Krause referred to, the
time required for this assay often makes it
impractical, but it's very important.

So given the data that we currently have in tissue culture, it's, you know, unlikely that human neoplastic cell substrates will be TSE infected or that exposure to potentially BSE contaminated bovine tissue culture products could lead to persistent TSE infection, but of course, as we're all well aware, you cannot guarantee zero risk.

And in the absence of that, what can you do to at least assess it as carefully as possible, the

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presence of TSE infectivity in these cultures? this is the answer I came up with, and after finishing it, I realized it's basically what we do in the lab to assess whether our tissue cultures contain infection.

There are now commercially available tests that can be used to do this, and as, again, Dr. Krause mentioned, you assay for PrP-sen, its expression level, determine its sequence. If it's a familial mutation, throw the cell line out.

Assay for PrP-res, and this would best be done in multiple cell sub-clones from the passage, as well as periodically at different cell culture passes because of the potential instability of the cell.

Bioassay for infectivity if all of this is negative. If any of these are positive, you know, again, throw out the cell line. If these are all negative or if this is negative, bioassay for TSE infectivity would have to be done, I should think.

And in terms of looking for human TSE infectivity or BSE, perhaps the best bet is to use human PrP or bovine PrP expressing transgenic mice. Keep those animals for up to two years, watch for clinical signs, and at the end of the day, look in the brains for pathology and/or PrP-res, and to do this, you know, relatively routinely if you're worried.

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1	So I'll stop there pretty much on time and
· · · · 2	take any questions.
3	ACTING CHAIRMAN DAUM: Thank you very
4	much, Dr. Priola.
5	And we'll have this presentation open for
6	discussion. We'll begin with Dr. Decker.
7	DR. DECKER: Just a simple clarifying
8	question to see if I've absorbed your implicit
9	definitions correctly. When you and Dr. Krause used
10	the term "infected," you're referring to a cell line
11	that actively is producing the abnormal prion protein,
12	and if we're just talking about pouring in some
13	preformed abnormal prion protein, you're going to use
14	the word "contaminated" or something like that?
15	DR. PRIOLA: Yeah, that's correct. When
16	I use "infected," I mean persistently infected.
17	DR. DECKER: So we've got a cell there
18	that's actually
19	DR. PRIOLA: Producing infectivity.
20	DR. DECKER: producing.
21	DR. PRIOLA: Yeah. It's making PrP-res,
22	it's accumulating PrP-res, and it's making new
23	infectivity.
24	DR. DECKER: Rather than simply pouring
25	contaminated
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1	DR. PRIOLA: Exactly.
2	DR. DECKER: bovine serum in there which
3	eventually gets diluted out and doesn't reproduce.
4	DR. PRIOLA: Right, which is what's
5	suggested by those immortalization experiments, yeah.
6	ACTING CHAIRMAN DAUM: Thank you.
7	Dr. Coffin, please.
8	DR. COFFIN: It's technically feasible,
9	although probably not very pleasant, to take cell
10	lines that one imagines one might use for vaccine
.11	production purposes and actually knock out both copies
12	of PrP. From what I understand, such cells would be
13	expected to be perfectly viable and you wouldn't
14	notice the difference. The whole animals are
15	perfectly viable without
16	DR. PRIOLA: Yes.
17	DR. COFFIN: PrP. Do you think that
18	would be an advisable thing to do?
19	DR. PRIOLA: Well, if you took away the
20	PrP, then I think you'd have a cell line that could be
21	very strongly argued would be completely resistant to
22	TSE infection, and I have never made knockout cells
23	myself, except from knockout mice, for example, PrP
24	knockout mice, but if it could be done, and it was not
25	technically too difficult, then, yeah, I think it

would be -- it would take away an element. You know, 1 what little risk there is, it would take that away 2 because PrP knockout mice are resistant completely to 3 infection. 4 5 ACTING CHAIRMAN DAUM: There was one more hand here. 6 Was it Dr. Myers? 7 DR. MYERS: When you're growing, effect, cell in culture and it loses the infectivity, 8 talking to people about this, the implication for me 9 is that if you're growing the cell fast enough, the 10 cell outgrows the PrP-res. Is that not an appropriate 11 12 way of looking at it? 13 DR. PRIOLA: Well --14 DR. MYERS: Because an awful lot of your infected cell lines are very slow growing things, and 15 you mentioned this business about the TSE from the 16 infected brains and so on. Is that a misconception? 17 18 DR. PRIOLA: You know, I don't think it's a hard and fast rule. 19 The neural 2A cells, which are mouse neural blastoma cells, we have troubles with 20 21 losing infectivity in these cells if we don't use very 22 specific types of medium. 23 There's another persistently infected cell 24 line called an SMB cell that is perfectly stable. 25 is slow growing. These fibroblast cells I showed you

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1	where it appears we got infectivity in there based on
2	the res signal, we have to split those if we're not
3	careful every two to three days. They are not slow
4	growing, and they are very reproducibly infectable
5	based on PrP-res with this one strain. So I don't
6	think it's a hard and fast rule.
7	ACTING CHAIRMAN DAUM: Thank you.
8	Dr. Katz.
9	DR. KATZ: I have a naive mechanical
10	question. When you talk about passages and failure or
11	success at pass, you're talking about passing
12	suspensions of cells, not just tissue culture fluid;
13	is that correct?
14	DR. PRIOLA: That's correct. These are
15	passage of TAP cells, yeah.
16	DR. KATZ: Thank you.
17	ACTING CHAIRMAN DAUM: Dr. Aguilar-Cordova
18	and then Dr. Stephens.
19	DR. AGUILAR-CORDOVA: I was wondering.
20	When you talk about some people being contaminated
21	with hospital instruments, would that be expected to
22	be a large inoculum, and then from your experiments in
23	vitro, it seems like you really need a huge inoculum
24	to get anything going. So could that be a big
25	differential between <u>in vivo</u> and <u>in vitro</u> ?

<u> </u>	DR. PRIOLA: Yeah. I mean, <u>in vitro</u> you
2	definitely need much more infectivity to get a
3	productive infection of tissue culture cells. As to
4	how much infectivity is actually on a contaminated
5	surgical instrument that passes this, I don't know,
6	but I do know that when it happens, the people who
7.	have what essentially happens is an operation is
8	done on a subclinical CJD patient, and those
9	instruments are used in other patients before it's
10	realized, and those other patients do get CJD.
11	So it's pretty effective, and with
12	presumably, you know well, I don't know the dose.
. 13	I don't know the dose, but it may be very low.
14	ACTING CHAIRMAN DAUM: Dr. Stephens, and
,15	then Dr. Coffin.
1,6	DR. STEPHENS: Just a follow-up on the PrP
17	knockout question. What does PrP do in normal cells?
18	DR. PRIOLA: That's a really good
19	question. It's not precisely known, but there was a
20	recent really nice paper in science that suggested
21	that if you cross-link it in differentiated neurons,
22	it activates the Finnkinees (phonetic) pathway.
23	DR. STEPHENS: And as a follow-up, was
24	there any evidence, is there any evidence of an
25	inoculum effect in new variant CJD in terms of BSE

1	exposure?
2	DR. PRIOLA: You mean inoculum effect, how
3	much you need?
4	DR. STEPHENS: Yeah, how much? I mean, is
5	there a relationship in any of the epidemiological
6	studies to beef exposure?
7	DR. PRIOLA: Yeah. Not being an
8	epidemiologist, I don't know the precise answer, but
9	I do know that the current thinking right now is it's
10	inefficient because it's inefficient. You're
11	crossing a species barrier. You're going orally, both
12	of which are bad things. So you presumably need more,
13	a pretty good bolus.
14	DR. STEPHENS: I appreciate that, but is
15	there data that supports that?
16	DR. PRIOLA: Not that I'm clear on.
17	ACTING CHAIRMAN DAUM: Michael Decker
18	might have data about that, an answer for that.
19	DR. DECKER: I don't know if there's the
20	type of hard data we'd like, but an illustrative
21	anecdote is the town in England where the butcher shop
22	that served the local community was using centuries
23	old techniques that heavily contaminated the resulting
24	meat with neural tissue, and there was a very high
25	attack rate, a cluster of human variant CJD in that

1	town.
2	So they found both an unusual
. 3	preponderance of disease and an unusual practice that
4	would give them a high load.
5	DR. PRIOLA: Yeah, I'm not sure what
6	the I know there was a cluster of five patients,
7	but I'm not clear what the total number of people were
8	that were exposed, and so what the exact attack rate
9	was. Do you?
10	DR. DECKER: No, but five out of the 90 in
11	England is a pretty high proportion.
12	DR. PRIOLA: Oh, for sure. Oh, yeah.
13	DR. DECKER: The proportion of that town's
14	population to the total English population.
15	DR. PRIOLA: Definitely, yeah, yeah.
16	ACTING CHAIRMAN DAUM: Dr. Coffin and then
17	I'd like to move on.
18	DR. COFFIN: So what do we know about
19	infectivity in serum of affected animals and their
20	offspring?
21	DR. PRIOLA: In terms of cattle?
22	DR. COFFIN: Yeah.
23	DR. PRIOLA: In cattle?
24	DR. COFFIN: In cattle, yeah.
25	DR. PRIOLA: There are only two reports

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1	that I know of that have suggested there might be very
2	low levels of infectivity in blood of cattle, one of
3	which is in an experiment where they experimentally
4	inoculated cattle and assayed tissue over the entire
5	course of disease. There was a point at which they
6	detected infectivity in bone marrow cells from a pool
. 7 	of three samples.
8	And there was a recent report last year
9	where it's a bit more indirect, where sheep infected
10	with BSE a bunch of sheep were infected with BSE,
11	and at various time points blood was transfused from
12.	those sheep into naive sheep, and one of those sheep
13	got sick, the implication being that infectivity got
14	out into the blood.
15	DR. COFFIN: But have there been serious
16	attempts made to look into
17	DR. PRIOLA: There have.
18	DR. COFFIN: serum products and other
19	blood products?
20	DR. PRIOLA: Commercial serum products?
21	Not that I
22	DR. COFFIN: Or in serum that's obtained,
23	you know, large amounts of serum from large numbers of
24	significant
25	DR. PRIOLA: Yeah, and in terms of

1	hamsters, yes, there was. Bob Rohwer did a really
2	heroic effort to do that, and pure concentrated
3	hundreds of mLs of hamster blood and was able to
4	detect using lots of animals that there's less than 1
5	LD-50 out of all that material.
6	DR. COFFIN: So it seems
7	DR. PRIOLA: It's extremely low.
8	DR. COFFIN: at the moment, based on
9	our current thinking, it seems improbable
10	DR. PRIOLA: Extremely.
11	DR. COFFIN: that there were these
12	significant risks from serum, to begin with.
13	DR. PRIOLA: Based on our current
14	knowledge.
15	DR. COFFIN: Based on our current
16	knowledge.
17	DR. PRIOLA: Absolutely.
18	ACTING CHAIRMAN DAUM: A burning final
1.9	comment.
20	DR. MINOR: I mean, while the bone marrow
21	result is out there, okay, I think it was regarded as
22	slightly questionable, and I think also that the
23	efforts that people have made to find infectivity in
24	cow blood as opposed to bone marrow have uniformly
25	been negative, and that's even going cow to cow, as I

understand it. 1 2 DR. PRIOLA: Yeah. 3 DR. MINOR: And I think there's also an 4 issue of the species that you're looking at will give you a different level of infectivity in the blood 5 perhaps. 6 7 DR. PRIOLA: Sure, absolutely. 8 ACTING CHAIRMAN DAUM: Thank you for that 9 clarification. 10 I'd like to move on to Dr. Krause, who promised we would have our opportunity to have at him 11 12 again, and here it is. Dr. Krause will talk about adventitious agent testing of neoplastic cell 13 14 substrates. 15 DR. KRAUSE: I'm just going to dive right 16 in and view this as a continuation of where I left 17 off. This slide shows a number of different episodes 18 throughout the history of the world and the U.S. of contamination of biological products with adventitious 19 20 agents, and I don't want to go into this in very much 21 detail, but I do want to focus on the contamination of 22 polio and adenovirus vaccines in the late 1950s and 23 early 1960s with SV40 as an example of something we

And as many of you know, millions of

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really want to avoid repeating.

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people received SV40 contaminated polio and adventitious vaccines into the late 1950s and early 1960s, and these vaccines were produced in primary Rhesus monkey kidney cells.

SV40 was identified as a virus; it was discovered after this, and the basis for discovering SV40 was the fact that supernatants of primary Rhesus monkey kidney cells cause tumors in laboratory animals and also were shown to cause cytopathic effect in primary Circa Pithacus (phonetic) monkey kidney cells, or different species of monkey, which then led to the discovery by Dr. Hillaman (phonetic) at Merck of SV40.

The vaccine seeds were treated with antiSV40 neutralizing antibodies in the early 1960s to rid
them of SV40. Epidemiological studies suggest luckily
that there was no adverse sequelae to the vaccinated
children, and however, recently SV40 DNA has been
detected in some human malignancies by PCR, and this
highlights a significant problem that one has with
adventitious agents, which is that sometimes the
concerns and the potential effect might not become
apparent for a substantial period of time, until a
substantial period of time after the contamination
event has occurred.

So a couple of lessons learned here are

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obviously the value of insuring that the products are free of adventitious agents, and also the particular importance of insuring the freedom of products from oncogenic agents or potentially oncogenic agents, especially for vaccines that are given to children.

So I've listed here a few basic principles of adventitious agent testing. We've already talked about how this kind of testing should consider quantitative issues and how important it is to understand the sensitivity of one's assays and what one precisely is tested for and what the meaning of a negative result actually is.

It's important to consider issues that are specific to the material in question, and I would argue that also where possible one should use tests that have the potential to detect unsuspected or even undiscovered agents.

Now, to talk about adventitious agents, I think it's useful to think a little bit about how viral vaccines are produced. The center of viral vaccine production is the cell substrate, and of course, for live viral vaccines or inactivated viral vaccines, some cell substrate is required to grow the virus.

You add to that a vaccine seed and various

other raw materials, which can include serum, medium, stabilizers or things like that, which then together in the production process after usually some kind of purification or inactivation lead to the final product.

This entire thing takes place in a facility which hopefully is designed in such a way as to minimize the opportunity for contamination at any of these steps, although there is the risk, of course, that personnel who perform these steps might contaminate a vaccine.

But a basic principle of producing these vaccines is that all of the materials that go into the vaccine, the vaccine seed, the raw materials and the cell substrate, need to be shown to be free of adventitious agents as they enter the process.

Now, purification steps could potentially reduce adventitious agent burden if some were able to get into the production process, and there is a model for thinking about that that has been used in evaluating therapeutic products at CBER in which some cell substrates might contain viral particles, and the ability of a purification scheme then to remove those particles is then determined, and that's called investigation of viral clearance.

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And a general principle there as well then is that the purification scheme which is used needs to be able to remove no only the amount of material that you think could be present in the cell substrate, but also some additional amount which represents a safety factor.

It's also worth noting that for live vaccines a purification process could actually potentially concentrate adventitious agents.

Okay. So as we look at adventitious testing then, I've talked about the importance of quantitation, but I just wanted to go through four individual points here because in policy making not only do we need to understand how sensitive an assay is, but it really does need to estimate the pretest probability of a problem because that will influence the kinds of assays that one thinks needs to be done.

One needs to consider the number of doses or dose equivalents that can be tested. One needs to understand the sensitivity of the assays, and a very important component of that then is in performing assays to include the appropriate controls that allow one to define how sensitive any given assay was, and the need to consider safety margins in deciding what kind of assays need to be performed.