FOOD AND DRUG ADMINISTRATION

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CENTER FOR BIOLOGICS EVALUATION AND RESEARCH

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VACCINES AND RELATED BIOLOGICAL PRODUCTS

ADVISORY COMMITTEE

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OPEN SESSION

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TELECONFERENCE MEETING

ON CBER LABORATORY SITE VISIT

LABORATORY OF ENTERIC AND SEXUALLY

TRANSMITTED DISEASES

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CONFERENCE ROOM C
BUILDING 29B
NATIONAL INSTITUTES OF HEALTH
BETHESDA, MARYLAND

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MONDAY, NOVEMBER 18, 2002

+ + + + +

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1	PROCEEDINGS
2	(1:05 p.m.)
3	DR. DAUM: I'd like to begin by
4	calling the meeting to order. Jody, would you
5	like to go through another roll call first, or
6	are you satisfied with that?
7	DR. SACHS: No, I'm happy. There's
8	maybe two other people three other people
9	we'll just notify everybody.
10	DR. DAUM: Everyone is on except Drs.
11	Griffin, Palese
12	DR. PALESE: I'm here.
13	DR. DAUM: Hello, Dr. Palese.
14	Everyone is on except Drs. Griffin and Whitley at
15	this point. Dr. Diaz we know wasn't coming, and
16	Dr. Gellin is still a question mark.
17	ARBITRATOR HOCKENBERRY: Estuardo
18	Aguilar-Cordova is about to join us on the call.
19	DR. AGUILAR-CORDOVA: I'm here.
20	DR. SACHS: Great.
21	DR. DAUM: Okay. Thank you. You're
22	not on my voting list, D. Aguilar-Cordova, so I
23	made a mistake. I'm sorry.
24	So, with that attendance information,
25	Jody, why don't we turn the floor over to you for

announcements.

DR. SACHS: Okay. I just have a little bit of a meeting statement to read. Jody I'm Sachs, the Secretary for today's meeting of the Vaccines and Related Biological Products Advisory Committee. like to welcome all of you to the meeting of the Advisory Committee. There is a speaker phone for public participation located here in Conference Room C of Building 29B, on the NIH Campus.

This afternoon's session will consist of a presentation and committee discussion that are open to the public. We will then go into a closed session until the meeting is adjourned, as described in the Federal Register Notice of October 30, 2002.

Should a committee member get dropped from the teleconference, simply call back at the 888-324-0789 number and give the pass code as VRBPAC. The operator is under instructions only to connect the committee members to this line. Again, you have the number so, if you get disconnected, please call back. If you have any problems while we're on the call, you can press "*0" and the operator will help you.

We ask that you do not place us on

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many clinical centers hold because have background music, and it can be very distracting to those remaining on the teleconference line. However, I strongly urge everyone to use "*6" or a mute button -- it is the same as a mute button -- since there are many lines connected, background sound will be decreased quality of sound for everybody would improve if you just use mute.

I just want to let everybody know who is in the room wiht me, in front of me. Karen Midthun is here. She's the Director of the Office of Vaccines Research and Review; Dr. Neil Goldman, Associate Director for Research, Office of the Center Director in CBER is here. Richard Walker, Director of Division οf Bacterial, Parasitic & Allergenic Products. And we have two of the people in the Laboratory of Enteric and Sexually Transmitted Disease Labs, Dr. Dennis Kopecko, who is the Laboratory Chief, and Dr. Philip Boucher. Also on the call is Dr. Sam Katz, who is the Site Visit Team Chair.

I would now like to read a public statement for the record. The following announcement addresses conflict of interest issues associated with today's meeting of the

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1	Vaccine and Related Biological Products Advisory
2	Committee on November 18th, related to the review
3	and discussion of intramural research programs of
4	the Laboratory of Enteric and Sexually
5	Transmitted Diseases.
6	Based on the agenda made available, it
7	has been determined that the committee
8	discussions present no potential for conflict of
9	interest.
10	At this point, I'd like to turn the
11	meeting over to you, Dr. Daum.
12	DR. DAUM: Thank you very much. We're
13	now going to go to the open session portion of
14	this afternoon's meeting. The meeting, of
15	course, as Jody mentioned, is devoted entirely to
16	the findings of the CBER laboratory site visit of
17	the Laboratory of Enteric and Sexually
18	Transmitted Diseases.
19	So, we are first going to hear from
20	Dr. Walker regarding the overview of the Division
21	of Bacterial, Parasitic and Allergenic Products.
22	Dr. Walker.
23	DR. WALKER: Thank you. Good
24	afternoon. I'd like to take just a few minutes
25	to, as Dr. Daum said, give you a quick overview
26	of what goes on in the Division of Bacterial,

Parasitic and Allergenic Products.

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Basically, the division is focused on assuring safe and effective products for immunological control of bacterial, parasitic and allergenic agents affecting human health. It's important to keep in mind that the scientists in this division perform both research and review functions. In some cases for our investigators, this review function could be 30 to 50 percent of their time.

Another point I'd like to make focuses in on the review functions that these people do. I'd like to emphasize that their review work is cradle-to-grave type operation it because begins with pre-IND meetings with sponsors help provide guidance, and then through the IND process, continuing meetings and review, license actions, and then it's very important to keep in mind that even post-licensure of a product that the task of this division goes on because there's review of biological deviation reports, there's inspections, lot release issues, and so forth. So, these people have a lot of work in the review are as well as in the scientific area.

I'd like to make another point about - that focuses in on the scientific area that

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these people in LESTD and other parts of division deal with is in there are quite variety of products. When you consider the products that are existing now or that are possible in the foreseeable future, I think of respiratory from Streptococcus Moraxella, pathogens, to sexually transmitted diseases, pathogens might be encountered by penetrating inoculation, like Malaria and Borrelia, and so forth; special pathogens which has become a big issue in recent Bacillus anthracis, Clostridium, like years, Franciscella and Yersinia; diarrhea-causing pathogens which will be discussed a bit today, and so forth, allergenic products whether it's antigens latex or cockroach and skin antigens. So there's quite a variety of things that the people in this division have to cover.

And so the division, to do this, is made up of about 100 people organized into eight laboratories. There's the Laboratory of Methods Development Quality Control, which seeks to develop and standardize and evaluate quality control methods for bacterial vaccines, and also develop, evaluate and apply serological methods to measure immune responses in vaccine trials. And also this group more recently is working to

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coordinate quality assurance activities within the division and provide leadership and the initiative to accredit CBER quality control testing laboratories.

Another laboratory, the Laboratory of Bacterial Polysaccharides is involved in characterization of immune responses to polysaccharide and conjugate vaccines, in standardization of methods for relevant clinical application. They also have been instrumental in developing novel physical and chemical methods improved for evaluation $\circ f$ licensed and experimental vaccines involving polysaccharides.

A third laboratory is the Laboratory of Biophysics, which use high-end instrumentation to help characterize biopolymers such as polysaccharides, DNA, proteins and macromolecular assemblies such as vaccine/adjuvant complexes. And the high-end equipment like applying NMR technology to vaccine characterization.

Another laboratory in the division is the Laboratory of Respiratory and Special Pathogens, which deals with Bordetella pertussis, as well as more recently with Bacillus anthracis and Yersinia species. And the work that these people are doing is to characterize the virulence

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factors and studies on mechanism of actions of these factors, and also the regulation of gene expression of these factors.

Another laboratory is the Laboratory of Bacterial Toxins, which deals with neurotoxins as well as iron-regulated virulence factors in Corynebacterium diphtheriae, and also in Bacillus anthracis.

other laboratories There are two before the laboratory that we're going to talk The Laboratory of Mycobacterial about today. Diseases and Cellular Immunology, which evaluates protective innate and adaptive immune responses to intracellular bacteria, and they are presently assessment of DNA involved in vaccination strategies against tuberculosis, as well as identifying new antigens that might be useful in anti-tuberculosis vaccines.

The seventh of the eight laboratories that I want to talk about today is the Laboratory οf Immunobiochemistry, which deals with allergenic products. They are concerned with function, allergen structure and immunomodulation of the allergic responses, the study of chamokines and receptors in the modulation of immune responses to allergenic

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Then the eighth laboratory, which Dr. Kopecko will be talking much more about this afternoon, is the Laboratory of Enteric and Sexually Transmitted Diseases, which of course is dealing with enteric pathogens, but also in more recent times is dealing with several bioterrorism agents, and I'm sure Dr. Kopecko will mention that a little bit more.

I'd like to conclude by just bringing your attention to some of the realities facing our Researcher/Reviewers in the division. Some of the things they have to deal with are, as you would expect in large organization, а bureaucratic hurdles, like personnel and other things that sometimes make work challenging, even though we've got very good people working with Another issue these Research/Reviewers have by working in the Federal Government, they have the issue of funding levels are uncertain from year to year, and dependent upon the appropriation process.

But the most unique thing, I think, about the FDA Researcher/Reviewers as opposed to people in other government agencies like CDC or NIH is that since these people are reviewers, the

1	timing of their workload is determined by the
2	sponsor submissions and not by CBER. And so they
3	have to carry out the research work and also meet
4	deadlines that are part of the review function,
5	and that is one of the challenges that our
6	investigators are meeting.
7	And, finally, when we had the actual
8	site visit, I asked the Site Visit Committee to
9	not only help us by reviewing the individuals and
10	reviewing the programs that they are directing,
11	but also comment on the current and future
12	directions of the research.
13	So, hopefully that gives you a very
14	brief overview of our Researcher/Reviewers and
15	sort of the nature of the work that goes or
16	within this division, and I'll be glad to answer
17	any questions that you might have.
18	DR. DAUM: Thank you very much, Dr.
19	Walker. The floor is open for committee question
20	or comment.
21	DR. KATZ: This is Sam Katz. I guess
22	I'll reserve mine until after we go through the
23	report.
24	DR. DAUM: That sounds fine, your
25	choice. Any others?
26	(No response.)

Then why don't we move on to the next item, which is we're going to hear from Dr. Kopecko. Welcome, Dr. Kopecko, and we will have an overview of the Laboratory of Enteric and Sexually Transmitted Diseases.

DR. KOPECKO: Thank you very much. Good afternoon to everyone. It's a great pleasure to have the opportunity to introduce the lab to you. In ten minutes, I'll try to give you a little bit of a flavor for who we are and what we do both regulatory and research-wise, and mention a few of our future directions.

If you have the Site Visit Notebook in from of you, you can see from Section 2 that the laboratory is comprised of three sections -- the Pathogenesis Section, the Gene Regulation Section, and Dr. Stibitz and Dr. Boucher will be reviewing research in that area for the review And the third is a new section, the Immune Mechanisms Section, that Dr. Walker heading up, that's just begun, and there wasn't any significant amount of research carried out during the review period, but as we move more into anthrax and Shigella vaccine development, Mechanism Section carries that Immune importance.

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The lab also is evolving from five personnel to 14 personnel, so we're undergoing some significant changes space-wise and personnel-wise. We currently have 13 people onboard, with one more hopefully to join us within the next couple of months.

If you can see from Section 3, our lab 1994, replaced started in the former was Micoplasma Lab. It was initiated to handle a large number of INDs and regulatory work enteric diseases, and also to cover sexually transmitted diseases which, in fact, were thought t.hat. there would be onslaught an οf STD applications, which has not actually come yet, but there has been a tremendous increase of enteric disease applications.

The lab, as Dr. Walker mentioned, has a bifunction, as all labs here at CBER. We are tasked with carrying out regulatory review in our area of specialty, as well as to conduct research which enhances our knowledge base, which allows us to increase both the quality and speed of the regulatory review of various products. So, our research is aimed at our area of specialization.

Within LESTD, we conduct basic research into the genetic and molecular bases of

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disease pathogenesis and the host response to infection. And the current events that have been analyzed during the last review period involve prokaryotic eukaryotic and mechanisms bacterial involved in invasion of mucosal epithelial cells -- that is, induced endocytosis and also induced exocytosis -- as well as the global regulation of expression of bacterial virulence components, which Drs. Stibitz and Boucher will address.

Our regulatory responsibilities, you can tell from the title, cover both enteric and sexually transmitted diseases, but I wanted to summarize some of the more common and some of the less common types of agents that we cover.

This includes, obviously, cholera, Typhoid Fever, Shigellosis, as you might expect, but also enterotoxigenic E.coli and pathogenic E.coli such as Campylobacter, LOBattenuated bacterial vector based vaccine development for multiantigen use. These include modified Salmonella bot.h strains, Shiqella strains and cholera strains. We also cover Helicobacter pylori, the use of live enteric bacteria such as attenuated Salmonella typhinerium given IV to treat metastatic cancer,

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various UTI pathogens, proteus, E.coli, intestinal strain replacement for probiotic therapy, the use of bacterially derived enzymes for treatment such as alosperigenase for acute leukemia, bacterial lymphocytic phase for therapeutic antimicrobial use, bovine or chicken immunoglobulin concentrates from specifically immunized animals for human use, and genetic harbored vegetables, to mention some of the products that we cover.

These encompass products that are administered by oral, intranasal, parenteral, intrarectoral, or transcutaneous routes, and they also cover evaluation of new adjutants for orally administered products.

Currently within the lab we cover about 110 different IND products. We are receiving about 12 to 14 new IND applications per year, so you can see how that list is growing. We probably drop about five to seven per year that become inactive. We also are reviewing one full biological license application on Cholera currently, and we have several BLA supplements under review.

So, that covers the general lab and our regulatory responsibilities. I'd like to

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take a couple of minutes and go back to Section 9 and just quickly summarize some of the research carried out in Molecular Pathogenesis Section.

Our main focus was to look at the cell biology of invasion of intestinal epithelial cells by Campylobacter jejuni, which I'll get to summarize in a minute, but we also carried out a number of more minor projects which all turned out to result in publications, a couple of which I want to point out because I think they have enhanced significance.

One involves Nanobacteria, which was reported by a Alevi Pejander (phonetic) in Finland, to be a new species of bacteria that were very small, filterable through .1 micronfilters, potentially important adventitious agent biological products because they are found in more than 80 percent of bovine and human serum, thought to be the main cause for human kidney stone formation, and more recently touted to be the main cause of atherosclerotic plaque, still being pushed by Dr. Pejander.

We followed up his results, which were mostly phenomenological studies with the heaviest piece of data being PCR 16Srod assembled DNA. We were able to obtain growth of these same coccoid

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like structures that had a calcified outer shell. They looked like coccoid bacteria. However, when we got into molecular evidence, we found by PCR we obtained the same sequence that Pejander obtained, but in going to gene bank, we found that he had misinterpreted the data and, in fact, this was the sequence from phylo bacterium, which is a common water-borne contaminant that's found in PCR reactions that are carried out for a high number of cycles. Also, the growth of structures could not be inhibited in azides, which all respiration-dependent organisms would be inhibited by. So, the upshot οf these studies that we published in P&AS was that there was no credible molecular data to support the existence of Nanobacteria. We feel that these structures are actually hydroxy-appetite crystals that are triggered into initiation by various macromolecules in blood and in saliva as part of a natural process. And, in fact, Pejander reported, you can destroy macromolecules by heavy gamma-irradiation. there is a way to knock them out, but we didn't feel that there was strong evidence any this support the existence of new of organisms, removing it at least for the current

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time as an important potential adventitious agent.

The second project that I want mention quickly involved Shigella sonnei. was a project that we started some 20 years ago, trying to make a modified hybrid vaccine salmonella that would protect against typhoid and Shigella, and this involved moving the LPS genes from Shigella sonnei in Salmonella typhi, Ty21a. The difficulty is this block of genes is very large, and more recently we were able to clone downsize to about 12kb the essential genes. found they stably that are expressed in Salmonella off of illegal copied plasmid an versus the typical multi-copied plasmid that most cloners use. We also found that there is adjacent insertion sequence, Is91, that causes instability of this. Once that's deleted, stabilizes the clone region for production.

And we also found that this form one polysaccharide is expressed in two forms in Shigella and in E.coli, both as a lipoaliga saccharide attached to core as normally, but also as a capsule where it's lipid-bound and stuck in the outer surface of the cell. And we were able to transfer this into Salmonella and make a

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These studies were published recently in Infection and Immunity, patented, and we had one firm who's carrying out commercially development of this into a vaccine. We are extending these studies to include Shiqella dysenteria and couple of other Shiqella а for serotypes future vaccine development, together with development of a live oral vaccine against anthrax.

Now, the last thing I want to mention is our work on Campylobacter jejuni. We carried out a number of in vitro tissue culture invasion assays into various cell lines, using a variety of different inhibitors. And to understand the cell biology of entry, these studies have been summarized in 2001 Trends in Microbiology Summary, for the most part. We found that the organism recognizes a receptor that's probably in the junctional space, the receptor is not yet characterized, and it's carried within cuveale interactions of the bacteria with this receptor think, secretion of trigger, we invasion is effectors this work Mike ___ bу Comp (phonetic), not ours -- these invasion effectors, we found, trigger an up-regulation of PI3 kinase,

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1	a release of intracellular calcium from
2	intracellular stores, activation of calmodulin
3	and protein CON-AC, which cause a structural
4	rearrangement and this is a microtubule
5	dependent up-take mechanisms as opposed to a
6	microfilament dependent that most other bacteria
7	use the cytoskeleton is reorganized, the
8	bacteria enter through a microtubular like
9	extension, they are carried in an endosomal
10	vacuole along microtubules via the molecular
11	motor dinaene to raise the lateral surface where
12	exocytosis occurs. So, that's the current state
13	of understanding for Campylobacter jejuni entry.
14	And if there aren't any questions, I'll turn
15	things over to Dr. Stibitz.
16	DR. DAUM: Dr. Kopecko's very
17	interesting presentation is open for questions
18	and discussion.
19	(No response.)
20	Well, I guess there aren't any
21	questions or discussion, Dr. Kopecko, which means
22	you were crystal clear. Thank you very much.
23	We will call on Dr. Stibitz for an
24	overview of research activities in his area.
25	DR. STIBITZ: Thank you. And thank
26	you for letting me phone in. I'm actually in

southern Illinois, in my ancestral home, and I'm willing to bet I'm the only person who literally dialed-up to the conference call.

LESTD section of The t.hat. concerned with is interested primarily in particular aspect of bacterial of one pathogenesis, and that is the regulation virulence factor expression. The model system that we use is that of Bordetella pertussis, which has a very well-developed regulon governed by the BvgAS two-component system. Now, BvqAS, like other two-component systems, contains environmental sensor protein which is localized in the inner membrane, BvgS in this case, this protein communicates with the cytoplasmic response regulator protein, which is BvgA in this These two proteins communicate by the case. kinase activity of the BvgS sensor protein, when BvgA is phosphorylated, it binds to the promoter regions of virulence.

Now, in pertussis, these include two which I'll mention, the fha which encodes the filamentous hemagglutinin, an important adhesion, and ptx which encodes the pertussis toxin responsible for some of the hallmark symptoms of whooping cough.

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in Now, know from vitro we transcription studies that BvgA phosphate is both sufficient necessary and to transcription of virulence seen. However, there are interesting and significant differences in how this occurs, and a classic experiment which demonstrates this is that if one grows pertussis under conditions where the virulence regulon is turned off and then shifts it to permissive conditions, one sees that fha transcription is initiated within minutes while that of ptx takes And it has been suggested to several hours. represent a temporal pattern of gene expression upon infection.

Now, this difference in one of the aspects of virulence gene regulation that we're most interested in, and our ultimate goal is the complete understanding at a molecular level of the interactions of BvgA with promoter DNA and with RNA polymerase in the process of transcriptional activation. And to accomplish this, we've taken a combined biochemical genetic approach.

Now, until now, our efforts have concentrated on the fha promoter. This is one of the simplest, and actually the simplest, BvgA

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regulated promoter described. And it was our hope that we could best learn the details of BvgA action in this system and then broaden our sights to include other promoters. And, in fact, that's exactly the juncture that we now find ourselves at, and it has been quite satisfying.

Now, Dr. Philip Boucher, who will be speaking in a few moments, has been the driving force behind the biochemical analysis, and he'll be telling you more about that, but I just wanted to mention that that work has recently been accepted for publication in Molecular Cell.

I have been primarily involved in the genetic analysis, and I'd just like to go over that briefly. Past work includes both forward and reverse genetics. In terms of forward genetics -- in other words, screening for mutant phenotypes and then determining the site nature of mutations involved -- we have isolated mutants which are affected in the differential regulation -- in other words, ones which can express fha but not ptx, and a reverse of that, those which are hyperactivated for ptx expression and down for fha expression.

In terms of reverse genetics -- in other words, introducing specific mutations like

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bacterialistic mutagenesis and then determining their phenotypic effect -- we have confirmed to role of the amino acid D54, we have made short Cwhich deletions of terminal BvqA had interesting phenotype in that they were lethal to the cell and were suppressed by mutations in the subunit of RNA polymerase and, alpha recently, we've made a number of 16 substitution mutants at different positions, and many of these interesting phenotypes as well either terms of differential regulation or, in one case, believe t.he mutation which affects we oligomerization of BvqA.

Now, my future genetic analysis οf involved a full-scale genetic assault BvqA and rests on some genetic screens which BvqA we've developed, which will allow us to identify mutations that specifically affect the ability of BvqA to interact with RNA polymerase or to itself interact with in the process of dimerization or oligomerization. And I won't go into those, but input into these will be BvgA in we've done either intense in which mutagenesis or, in fact, saturation mutagenesis And by alynase scanning. what we hope identify are specific messages in BvgA which are

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involved in interacting with RNA polymerase or with itself in the process of oligomerization and dimerization.

Now, recently two other people have joined the laboratory, and I'll just briefly describe their projects. Dr. Wendy Veal Carr is ERDA Fellow who came to us from Bill Schaeffer's lab at Emory University, and Wendy is embarking on a project which, if successful, will from allow out our detailed us to zoom mechanistic analyses of BvgA to an organismal or host pathogen interaction level. And she's adapting RIVET, which stands for recombinasebased in vivo expression technology. This system was developed by Andy Kanuli, who is not Tufts, and it was to examine in vivo expression oligliocollar. The system uses a DNA recombinase as a reporter gene, and the readout is the degree of recombination of a construct in the chromosome which, upon resolution, leads to a block with antibiotic resistance. And that's after appropriate strain construction one is able to infect an animal -- in our case, a mouse -and then recover the bacteria at different time points and, by affecting the degree of recombination, affect the level of expression

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that that particular gene -- in our case, we'll expand an fha and ptx -- has experienced.

And we plan to use this to determine if the differential regulation seen in vitro with fha and ptx develop in a manner accepted in vivo and, secondly, to determine the effect mutation -- for example, which knock out important adhesions such as fha the on expression of pertussis toxin.

And then, finally, Dr. Brian James has just very recently joined us, actually, since the site visit, and he joins us from Dr. Robert Bender's lab at University of Michigan. And Brian is responsible for getting our anthrax project up and running, and in the initial phase he's developing new genetic tools specifically for powerful molelic exchange vectors and enhanced efficiency of DNA transfer, and then we plan to use those tools to study genetic regulation of virulence in anthrax. And I'll stop there.

DR. DAUM: Thank you very much, Dr. Stibitz. Dr. Stibitz' interesting presentation is now open for committee discussion and comment.

(No response.)

I was wondering if you could possibly

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1 say a couple of sentences about the big-picture 2 role of fha and the pathogenesis of pertussis -why is this of particular importance? 3 vaccine antigens that we talk about and bandy 4 5 their importance of the around acellular vaccines, why this particular antigen important 6 7 to the pathogenesis? Well, I quess the most 8 DR. STIBITZ: sort of convincing evidence of that is that the 9 10 mutants of pertussis which lack expression of fha are decreased -- they have decreased ability to 11 adhere to host cells, and in some models have 12 decreased virulence. 13 DR. KATZ: If you ask John Robbins, he 14 says it doesn't do anything. 15 16 DR. STIBITZ: Fair enough. I think it is safe to say that the animal models that we use 17 18 may not be the best animal models for assessing 19 the effect of fha, but at least, you know, one aspect of -- I think whether or not fha 20 21 absolutely important in infection in a mouse model, we can still assess the degree of in vivo 2.2 23 regulation using the system that we're going to 24 use. 25 DR. KATZ: I guess what I would like

you to comment on -- I was one of the visitors,

so I heard this, but I don't know if the committee members did -- what is the relevance of Bordetella research to a laboratory of enteric and sexually transmitted diseases?

DR. STIBITZ: Yes, I get this question every time.

DR. DAUM: I imagine you do.

There are two answers to DR. STIBITZ: that. One is sort of the historical story of how I came to be where I am, and I don't think that's particularly relevant. I think what's relevant is the fact that the type of approach that we take, until recently, we were probably the only group within the division taking a very That set of knowledge has genetic approach. turned out to be very useful for the type of products which we particularly find in Enterics Lab. For example, rationally attenuated live vaccine which use the exchange methods that we use often in pertussis and actually developed in pertussis, and then also use issues generally relating to mutation rate, population genetics, and so forth, that impact upon genetic stability, and these also impact upon live preparation, vaccine as well as issues that relate to products see, such the we as

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Т	replacement flouro-type of approach that Dennis
2	mentioned, as well as bacteria therapy, which is
3	also something that's coming to the fore again.
4	So, I think pertussis itself is not
5	relevant. I think the system that we study, the
6	regulatory system, is widely relevant to
7	bacterial pathogenesis as a whole, including
8	enterics, and then our specific method technology
9	base help us greatly in the reviews that we do.
LO	DR. DAUM: Thank you.
11	MS. FISHER: Dr. Daum?
12	DR. DAUM: Yes.
L 3	MS. FISHER: Barbara Fisher. May I
L 4	ask a question?
15	DR. DAUM: Please.
L 6	MS. FISHER: Is any of your work going
L 7	to lead us to genetic screening techniques for
18	those who would be at high risk of potentially
L 9	reacting, say, to pertussis vaccine?
20	DR. STIBITZ: I don't believe so. In
21	other words, that would be primarily a host
22	characteristic, and our techniques are not
23	designed to assess that aspect of the
24	host/parasite relationship.
25	DR. DAUM: Thank you, Dr. Stibitz.
26	Are there other committee questions or comments?

1	DR. PALESE: Yes. Peter Palese,
2	please. Can I ask a question about the anthrax?
3	DR. DAUM: Certainly.
4	DR. PALESE: A new person who is
5	coming, we work on real characteristics of the
6	anthraces. Now, are there facilities available
7	at your institution, or how are you going to test
8	for A-virulence factors?
9	DR. STIBITZ: Well, initially, we're
10	trying to develop new genetic tools, and that may
11	take some time. When we do get to the point
12	where we're working directly with anthrax, much
13	of our work can be done with the A-virulent
14	strain. At this point, we're seeing a need for
15	doing actual work with fully virulent anthrax,
16	although, if that happens, we do have a petri
17	facility. In fact, we're actually going one
18	better and we're doing much of the initial work
19	in Bacillicereous, which is a very close relative
20	of anthrax, but much, much less virulent. Does
21	that answer your question?
22	DR. PALESE: Yes. Thank you very
23	much.
24	DR. AGUILAR-CORDOVA: This is Estuardo
25	Aguilar. I wonder, Dr. Stibitz, if you could
26	comment I was also on the review team but

if you could comment on the independence of Dr. Boucher within the work that you are doing and 2 the collaboration that involved. 3 4 DR. STIBITZ: The word you use is the 5 word I would use to describe it. Basically, Phil set of knowledge and tools 6 a to 7 problems that I don't possess, and he's developed all the biochemistry. He's done that essentially 8 I work in some of the mutant 9 on his own. construct, and so on and so forth, but 10 Boucher has done all that biochemistry from the 11 ground up, and it's a project that he initiated 12 13 and it's really a collaboration. I did the 14 genetics and he does the biochemistry. Ιf have more specific questions, I could elaborate, 15 16 but the degree of independence is very high. 17 DR. DAUM: Dr. Palese, a followup? 18 (No response.) 19 All right. Thank you. Okay. 20 committee questions, comments? 2.1 DR. SACHS: I just have a very quick This is Jody Sachs. I just wanted to 2.2 remind everybody that a Transcriber is recording 23 this telecon, and it's easier for the Transcriber 24 25 if everybody repeats their name before speaking 26 when they comment. Thank you.

1	DR. DAUM: Thank you.
2	DR. AGUILAR-CORDOVA: This is Estuardo
3	Aguilar one more time.
4	DR. DAUM: Please.
5	DR. AGUILAR-CORDOVA: Just another
6	quick question for either Dr. Stibitz or Dr.
7	Kopecko. Both of you have mentioned an increased
8	number of people. Is there also an increase in
9	space that's being considered?
10	DR. STIBITZ: Do you want to field
11	that one, Dennis?
12	DR. KOPECKO: Yes, I will. This is
13	Dennis Kopecko. Yes, we do have an increase in
14	space. We still would like to get more space
15	because even with renovations that are ongoing,
16	we are still going to be fairly cramped. But
17	within the limitations of the division and the
18	Center, we can't ask for anymore, really,
19	currently. But, yes, there is some additional
20	space, and certainly some additional funding for
21	per capita research.
22	DR. DAUM: Thank you.
23	DR. KATZ: Refreshing refrain. This
24	is Dr. Katz. You said you were going from five
25	to 14 people, which is what you said in your
26	introductory remarks. Those additional nine

Are these doctoral, post-doctoral, what 2 3 people are these? DR. KOPECKO: Good question. 4 If we go back to Section 2, to the 5 see here. organizational chart, you'll see that in 6 7 Molecular Pathogenesis Section, there are two Post-Doctoral level individuals joining. 8 Dr. 9 DeQi Xu, who just recently joined within the past 10 week, who is going to be working on Shigella vaccines, and a second Post-Doctoral Fellow to 11 work on Campylobacter pathogenesis. Dr. Stibitz 12 13 mentioned two additional individuals, Dr. Wendy Veal Carr and Dr. Brian James, who joined at the 14 Post-Doctoral level. Dr. Manuel Osorio is a 15 Fellow Ph.D. Cellular Immunologist, who 16 Staff joined us back in July, in the Immune Mechanisms 17 18 Section, and then we have an additional beginning research technician, and that sort of fills out 19 20 the field. So, mostly professional level, on one technical level. 2.1 2.2 DR. KATZ: Thank you. 23 DR. DAUM: Other comments, questions? 24 DR. FAGGETT: This is Dr. Faggett. 25 One question relative to constant research versus regulatory. What is the breakout in terms of how 26

people, at what level are they scientifically?

doing primarily the research -- but in terms of 2 how much time just on a regulatory? 3 DR. KOPECKO: This is Dr. Kopecko. 4 5 That's a very good question. In order to carry out competitive research, you obviously have to 6 7 have some people in the lab that spend the majority of time there. And what Scott and I 8 have sort of evolved into is a situation where he 9 10 and I take the majority workload for regulatory, we're splitting up around 40 to 45 11 applications each, and then the remaining 20 to 12 13 30 are parceled out with other members of the 14 laboratory, that way trying to keep most of the other individuals in the lab most of the time, 15 16 but they are all involved to some extent with regulatory, other than a couple of the Post-17 Doctoral Fellows who are not FDA personnel, but 18 19 they are actually individual contractors. 20 DR. KATZ: This is Dr. Katz. T think when we made our review, if I remember correctly, 21 Dr. Kopecko, you told me 50 percent of your time 2.2 is spent on regulatory, is that correct? 23 DR. KOPECKO: That's correct. 24 DR. KATZ: So, I think that gives you 25 26 a little idea, Walter.

much time -- sounds like the new Fellows will be

1 DR. FAGGETT: Thanks, Sam. 2 DR. DAUM: Other input? (No response.) 3 thank you, everybody, for your 4 5 And let's move on to hear from Dr. comments. who will give overview of his 6 Boucher, an 7 research activity. Dr. Boucher, welcome. 8 DR. BOUCHER: Thank you. I thought I 9 would very briefly summarize the work I presented 10 to the Site Visit Team a couple of months ago, then discuss a current project, and then finish 11 12 off with what is planned for the future. 13 interest in this lab since arrival here in 1994 has been on a biochemical 14 approach to an understanding of how a bacterial 15 16 response regulator effects control of expression of virulence factors. This work has evolved from 17 18 early investigations describing how the 19 phosphorylated BvgA regulator of Bordetella 20 pertussis binds to simple promoters to 2.1 recent experiments that have global implications 2.2 for molecular genetics in bacteria. 23 Initial studies employing DNA binding and both in vivo and in vitro transcription 24 assays formed the basis of our understanding as 25 26 to how BvgA activated transcription at the fha and promoters that have already ptx been discussed by Scott. For example, it discovered that activation of both promoters involves multiple dimers organized in such a way as to suggest an interaction with the C-terminal domain of the alpha subunit, one οf polymerase subunits.

Again, as Scott mentioned, the most recent work presented a couple of months ago will be coming out in the January issue of Molecular Cell, so I'll be going through it very briefly here. Using a technique termed affinity cleavage of DNA, we have developed a structural model of how both the BvgA regulator and RNA polymerase interact at a virulence factor promoter, specifically the fha promoter.

The technique is based on the ability of a chemical nuclease to cleave DNA primarily at sites dictated by the binding specificity of the DNA-binding protein to which it is bound. Analysis of cleavage patterns generated by such modified proteins bound to promoter DNA allows the development of structural models capable of addressing certain critical questions such as how many molecules of a given regulator bind a particular promoter and how are the different

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proteins in a transcription initiation complex oriented relative to each other.

Using this technology, we described a novel architecture in which multiple dimers of BvgA bind along one fact of fha promoter DNA while a part of the polymerase complex, the alpha C-terminal domain, binds to the opposing face along the same linear stretch of DNA.

In addition, modeling studies revealed that the orientation of BvgA monomers within the dimers is the same as that recently demonstrated by x-ray crystallographic methods for the C-terminal domain of the related response regulator called NarL bound to its DNA-binding site.

Since these experiments conducted, we have extended our analysis transcription initiation complexes to other virulence factor promoters of Bordetella pertussis. Previous studies using relatively low resolution analyses made certain predictions whose validity, in light of our current understanding, were now being brought into the question. In the pertussis toxin promoter, for example, stoichiometry of BvgA binding and novel orientations for how BvgA maneuvers itself onto a very unusual binding site had previously been

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Using our affinity cleavage assay, we recently determined that the previous studies had underestimated the number of BvgA dimers bound to the toxin promoter and that the unusual BvgA binding site within this promoter bound 2 dimers of BvgA in a very usual and unconstrained manner.

another promoter, the bipA Αt have others promoter, and shown that we requires activation at this promoter concentration is intermediate of BvgA that activate required to the fha between that promoter and that required to activate the toxin promoter. Indeed, at the relatively high vitro concentrations needed for toxin activation, the bipA promoter appears to be repressed. in vitro affinity cleavage assay has aided us in distinguishing sites that are likely involved in activation from sites that are likely involved in repression. Studies of other critical virulence factor promoters are currently ongoing.

Conclusions from the work iust described suggested other also interactions between BvqA the RNA polymerase. and For example, our data suggests that BvgA may interact both of the following: with one or The C-

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terminal domain of the sigma subunit of RNA polymerase and the N-terminal domain for the alpha subunit of RNA polymerase.

Experiments utilizing both an in vitro biochemical approach and an in vivo 2-hybrid protein-protein interaction assay are underway to address these possibilities. As there are only two copies of the alpha subunit per polymerase complex, it is also important to understand whether both copies are required for full BvgAdependent promoter activity. In vitro transcription experiments are also underway to address this question.

And, lastly, another project is in the proof of concept phase of investigation. How exactly phosphorylation of BvgA engenders an active complex is not fully understood. Based on structural data of unphosphorylated forms of homologous proteins, it has been suggested that phosphorylation results in the unmasking of the DNA Binding Domain.

Perhaps due to the instability of the phosphorylated form of response regulators, it has thus far not been possible to crystalize this form. In an attempt to circumvent these limitations, we are opting for the use of a

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physical method that would structurally compare the two forms of BvgA. This technology termed FRET, for florescence resonance energy transfer, is based on the ability of an activated flourophore to transfer its resonance energy to a neighboring flourophore.

As the probability of this transfer is dependent on the distance separating the flourophores, molecular distances be can calculated if energy transfer is monitored for a series of flourophore pairs distributed within a single biomolecule. If measurements are made for each pair in both the phosphorylated and unphosphorylated forms of BvqA, a model can be derived that describes the motion of one domain -- for example, the C-terminal DNA binding domain -- relative to another -- for example, the Nterminal masking domain. The construction of preliminary BvgA mutants that will eventually harbor flourophore pairs is currently underway.

That about concludes my brief summary of my lab activities, and I can entertain any questions.

DR. DAUM: Dr. Boucher's presentation is open for committee comment and questions.

(No response.)

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1	I would like to ask the same question
2	I asked of Dr. Stibitz, if you wouldn't mind, and
3	that is, can you tell us a little bit about how
4	fha, which is good after some impressive
5	regulation, factors into pertussis status
6	physiology, and how important it is to the
7	conduct of the organism or the ability of the
8	organism to go about its business?
9	DR. BOUCHER: Certainly. First of
10	all, it is clear that fha, the expression of fha
11	in Bordetella plays an integral role in its
12	pathogenesis. Without the fha, it really has no
13	clear way of adhering to epithelial cells.
14	However, with the work that I am conducting, fha
15	really here is looked upon more as a readout of
16	the activity of the two-component systems that
17	we're studying, and less really as an antigen for
18	any possible vaccine.
19	DR. DAUM: I understand that, but it's
20	still an interesting issue for me, so I have to
21	ask it anyway.
22	Other committee comments? Questions?
23	(No response.)
24	Input?
25	(No response.)
26	Dr. Boucher, thank you.
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1 DR. BOUCHER: Thank you. 2 Jody, I think it's that bewitching 3 time for the open public hearing. Do we know if there's anybody interested in addressing 4 5 committee? 6 DR. SACHS: No, but I'll just read a 7 statement. As part of the FDA Advisory Committee 8 9 meeting, we are required to hold an open public hearing for those members of the public who are 10 not on the agenda, who would like to make a 11 12 statement concerning matters pending before the committee. 13 14 I have not received any requests at this time. If there is anyone who would like to 15 16 speak -- address the committee at this time, this is the opportunity. And I look around the room 17 18 where I am right now, and there's no one from the 19 public wishing to speak. So, I will close the open public hearing and ask you, Dr. Daum, to 20 continue the meeting. 2.1 2.2 That sounds fine. DR. DAUM: So, 23 we're going to move into closed session at this 24 point, and I guess that's going to ask those individuals who need to take their leave at this 25

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point, to please do so.

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