

# Center for Biologics Evaluation and Research



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### My comments are an informal communication and represent my own best judgment.

These comments do not bind or obligate FDA.



## Products Regulated by CBER







- Allergenics
- Blood Products
- Devices Related to Biologics
- Gene Therapies
- Human Tissues and Cellular Products
- Vaccines (preventative and therapeutic)
- Xenotransplantation Products

### CBER Regulated Products: Vaccines for Disease Prevention >150 million doses of influenza vaccine given in 2016-2017

### Annual US disease, 1900s US disease in 2013

<ul> <li>Smallpox:</li> </ul>	29,005	0	
• Diptheria:	21,053	0	Challenges:
• Pertussis:	200,752	28,639	Probiotic therapeutics Vaccines for infectious diseases of global importance/EIDs
• Polio:	16,316	1	
Measles:	530,217	187	
• Mumps:	162,344	584	
Rubella:	47,745	9	
• H. Influenzae: 20.000		31	

https://www.cdc.gov/vaccines/pubs/pinkbook/downloads/appendices/e/impact.pdf

# CBER Regulated Products: Keeping the Blood Supply Safe



Need for continued vigilance against emerging threats

Successes\*:

- HIV: 1 per 2,435,000 units
- HBV: 1 per 1,565,000 units
- HCV: 1 per 2,680,000 units
- Pathogen reduction technology –approved for platelets

#### **Ongoing Challenges:**

Continued low rates of bacterial contamination Pathogen reduction technology – apply to rbc/wb Agents of TSE EIDs, especially arboviruses (Dengue, Chikungunya, Zika) Malaria-causing species; *Babesia microti* 

# **CBER Regulated Products:**

# Advanced Therapies at the Leading Edge

Ex vivo or In vivo gene editing to treat various conditions

**FDA News Release** 

#### FDA approval brings first gene therapy to the United States

CAR T-cell therapy approved to treat certain children and young adults with B-cell acute lymphoblastic leukemia

#### FDA approves CAR-T cell therapy to treat adults with certain types of large Bcell lymphoma

Yescarta is the second gene therapy product approved in the U.S.

#### For Immediate Release

October 18, 2017

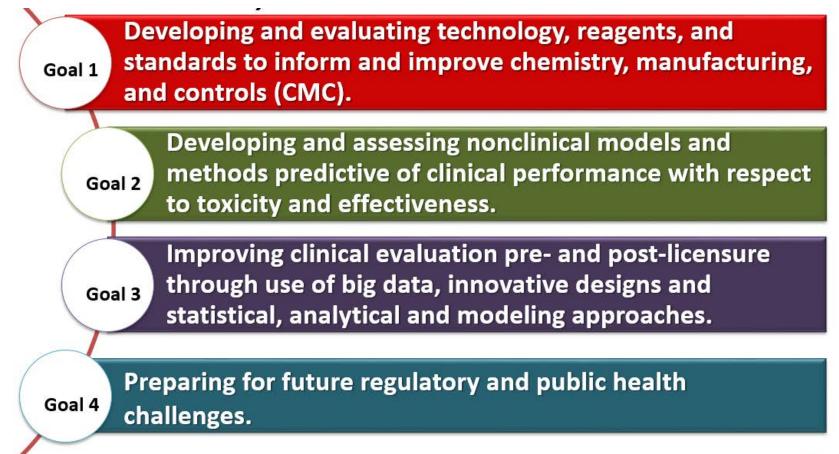
#### Challenges:

Preparing for new and evolving technologies, such as gene editing Manufacturing scale-up – applying continuous manufacturing? Stem cell-derived products Cells combined with devices (i.e. 3D-printed scaffolds) **D** 

## **CBER Research Goals**



# Advancing the scientific basis for regulation of biologics, human tissues and blood by:



# NCTR Collaborations Supporting CBER Research Goals



### • Goal 1: CMC

- Improved methods/development of standards to characterize products
- Goal 2: nonclinical
  - Develop novel approach to evaluate impact of nonsynonymous mutations
  - Bioreactor model to simulate C. difficile-host interactions
  - Detect off-target mutations of gene editing
  - PK and biodistribution of novel adjuvants (n=2)
  - Support development of animal model to support norovirus vaccine development



## **CBER Collaborations Supporting NCTR**

- Mechanisms of norovirus-*Salmonella* coinfection
  - CBER developing enteroid and animal model
- Functional screening of candidate molecules identified by NCTR SDAR for anti-trypanosomal activity



# Selected Updates and New NCTR Collaborations Supporting CBER Research Goals

**FMT-related projects (2)** 

Assays to evaluate genome editing

Use of NGS and ribosome profiling to evaluate codon optimization impact on therapeutic proteins

## Fecal Microbiota Transplantation (FMT)

Donor Screening/sample testing

What should we test for?

How good are the tests used for pathogen screening?

#### Does the manufacturing process alter efficacy?

- Oxygen exposure
- Lyophilization
- Freeze/Thaw



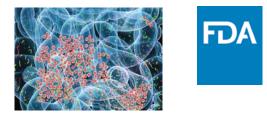


#### Product characterization:

- Potency?
- Efficacy?
- Manufacturing consistency?

- Disease/conditions associated with dysbiosis:
  - Bacterial infection/colonization
  - Inflammatory Bowel Diseases
  - Obesity
  - Metabolic Disorders
  - Autism
  - Parkinson's Disease

### CBER Goal 1: *Update* Pathogen Detection in FMT



- <u>Need</u>: Evaluate assays typically used to screen clinical samples for relative sensitivity when used to screen FMT products.
- <u>Why NCTR</u>: Experience with bioreactors that produce "stool cultures" by inoculating a base medium with human stool under controlled growth conditions.
- <u>Approach</u>: Bioreactors with active stool cultures will be inoculated with various levels of pathogen alone or pathogen plus FMT, and plating and NGS methods used to determine if pathogen can colonize. In vitro results will be expanded in mouse model of infection.
- <u>STATUS</u>: Research protocol approved; testing initiated
- <u>Potential Impact</u>: Improved understanding of assay applicability and identifying assays that need improving to ensure the safety of FMT products.

CBER Goal 2:Update Clostridium difficilehost interactions



- <u>Need</u>: Improved understanding of how commensal bacteria impact host dendritic cell responses to *Clostridium difficile*
- <u>Results</u>: Co-culture system (developed by CBER) with epithelial cells has been implemented at NCTR. Expanding to work on dendritic cells.
- <u>Potential Impact</u>: Improve our understanding of how microbiota impacts the immune response against *C. difficile*. May provide insights into how to regulate FMT, vaccine development for treating *C difficile*.

### CBER Goal 2: New



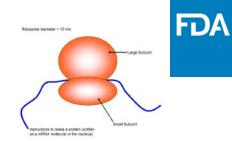
Off-target mutations in genome editing technologies

- <u>Need</u>: Address safety evaluation of advanced therapies involving CRISPR-mediated genome editing. In particular, to develop new methods and evaluate existing methods for offtarget mutation detection.
- <u>Why NCTR</u>: Well-equipped with NGS facility and extensive experience.
- <u>Approach</u>: Whole genome sequencing will be applied to enhance sensitivity of detection. Genomes of single-cellderived clones treated with CRISPR will be compared. Performance compared to other published approaches to detect DNA breaks.
- <u>Potential Impact</u>: Additional methods for detecting potential off-target mutations caused by genome editing. Valuable information on advantages and limitations of existing off-target detection methods.

### CBER Goal 2, New

Bioinformatic workflows for NGS data generated form ribosome profiling

- <u>Need</u>: Codon-optimization is employed more commonly. We know that it may impact protein structure-function in unexpected ways. Need better tools to assess.
- <u>Why NCTR</u>: NCTR scientists have experience with ribosome profiling data and analysis.
- <u>Approach</u>: Ribosome profiling will be used to identify codons that induce changes in ribosome movement across transcript; may identify how synonymous codon changes impact translation kinetics and protein conformation.
- <u>Status:</u> on-going (CBER does experiments; NCTR analysis)
- <u>Potential Impact</u>: This method may prove useful to assess impact of codon sequence changes on protein conformation as a consequence of codon optimization in regulated therapeutic proteins.





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