Choosing the right gene editing technology

February 18-19, 2016, San Diego-CA, USA

The Genome Editing & Engineering Congress brings together the key industry leaders and researchers to address the concepts, challenges and state of art methods & applications of the genome editing tools like CRISPR/Cas9, TALENs, ZNFs & AAVs etc. The case study & sessions will reveal the potential application of Genome editing tools from the modern biomedical & therapeutic application .Special emphasis on CRISPR system addressing the concept, technology, challenges like off-target effects, efficiency improvement and delivery systems etc.

## Attendees of the event will learn about:

- The cutting edge therapeutic application of Genome Editing tools: CRISPR/Cas9, TALEN, ZFN & AAVs
- How to overcome the challenges of CRISPR & other genome editing tools
- Advancements, challenges and future opportunities of CRISPR/Cas9 & other genome editing technologies
- The genome editing approaches to accelerate drug discovery, target identification, validation & screening
- ▶ The Genome Editing towards cell line engineering & disease model development
- Genome Editing application towards research animal models or transgenic animal
- The Regulatory challanges, Ethics and Intellectual property rights of Genome Editing technology

## **Conference Highlights**

- 15+ case studies on Genome editing application
- Keynote by renowned experts
- Presentations from the pharma industry
- Open forum to discuss the best tool for your research

## **Confirmed Speakers:**



For more information please contact Ajay at ajay.nimbalkar@mnmconferences.com | +91 20 6704 6819



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# Day 1, Thursday, February 18<sup>th</sup>

08:00	<b>Registration &amp; Refreshment</b>

## 08:50 Opening remarks from the Chair

**09:00** Keynote Presentation

## An Historical Perspective on the Development and Principles of Gene Editing Technologies

- Historical context, Comparision between Gene Therapy & genome Engineering, nomenclature, different mechanisms, methods and applications
- Classes of Edits or Modifications: Knockout, Knockin (gene or gene segments), Point Edits, Large Deletions

## Prof. Eric B. Kmiec

Director, Gene Editing Institute Center for Translational Cancer Research Helen F. Graham Cancer Center & Research Institute, USA

## 9:25 Understanding the role of CRISPR/ Cas 9 in Genome Editing from technology overview to the future prospective

- What are the advantages of CRISPR over other methods
- How to overcome the off-target Mutations
- What is the future of CRISPR/Cas9 system

## Methods of Genome Editing and Engineering: Concept, Technology & Challenges

### 9:50 CRISPR-revolution led paradigm-shifts in animal genome editing approaches

- Introduction to the long-used traditional animal transgenic technologies, using mouse as a model organism.
- Paradigm shifts in animal transgenic technologies caused by the CRISPR/Cas9 system
- Latest advances in CRISPR/Cas9 genome editing platforms that have completely relieved the bottlenecks of long-used transgenic technologies

## Dr. C. B. Gurumurthy

Director, Mouse Genome Engineering Core Facility, University of Nebraska Medical Center, USA

## 10:15 Solution Provider Presentation

Contact Steve Hambrook at steve.h@mnmconferences.com

## 10:45 Morning Refreshments & Poster Presentations

## 11:25 Intro to Genome Editing and Engineering

- RNA editing mechanisms and methods
- Applications of RNA editing
- New editing approaches in the context of prior editing literature
- Comparing therapeutic editing to gene therapy

### Dr. Tod Woolf

Founder and President, ETAGEN Pharma, USA

## 11:50 RNA guided genome engineering: new expansion of Cas9 toolbox and in vivo application

- New animal models using CRISPR-Cas9
- In vitro genome editing in postmitotic neurons using SpCas9
- In vivo genome editing in the mouse brain using SpCas9
- Applications of SaCas9 for genome editing in brain and liver

## **Dr. Matthias Heidenreich**

Post-Doctoral Fellow, Feng Zhang Lab, Broad Institute, MIT, USA

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12:15	Comparison of TALEN and RNAi for Targeted Mutagenesis or Target Gene Suppression in a Complex Genome
	Vector construction
	In vitro gene delivery
	<ul> <li>Alternative screening protocols to identify events with target gene knock-out and validation by sequencing</li> </ul>
	<ul> <li>Correlation of RNAi mediated target gene suppression with phenotype</li> </ul>
	<ul> <li>Rapid phenotyping protocol for knockout and knockdown events</li> </ul>
	<ul> <li>Comparison of performance of knockout or knockdown events</li> </ul>
	Prof. Fredy Altpeter University of Florida-IFAS, USA
12:40	CRISPRscan: Designing highly efficient sgRNAs for CRISPR/Cas9 targeting in vivo
	<ul> <li>A sgRNA-scoring algorithm capturing the sequence features affecting Cas9/sgRNA activity in vivo.</li> </ul>
	<ul> <li>Designing efficient alternative sgRNAs to increase the target site repertoire in the genome.</li> </ul>
	<ul> <li>Localizing Cas9 expression in the germ cells to reduce lethality and deleterious phenotypes in somatic tissues.</li> </ul>
	Dr. Miguel A. Moreno-Mateos
	Associate Research Scientist Department of Genetics, Vale University School of Medicine, USA
13:05	Lunch Break and Poster Presentation   One-to-One Networking Meetings
14:05	Indel Detection by Amplicon Analysis (IDAA): A Novel and Improved Methodology for Genome Editing Surveillance
	<ul> <li>A novel Indel Detection by Amplicon Analysis (IDAA) method for genome editing applications.</li> </ul>
	<ul> <li>IDAA is based on a simple amplicon labelling strategy and automated Capillary Electrophoresis.</li> </ul>
	<ul> <li>IDAA is emnable to high throughput detection and characterization of indels induced by precise gene targeting.</li> </ul>
	<ul> <li>IDDA is cost effective and generates indel profiles similar to Sanger and "Deep Sequencing" with sub-percentage indel</li> </ul>
	detection sensitivity.
	IDAA is highly useful for genome editing surveillance
	Dr. Eric Paul Bennett
	Associate Professor, Copenhagen Center for Glycomics (CCG), Denmark
	Application of CRISPR & other Genome Editing Technologies: Cell Line Engineering, Therapeutic Application, Animal Model, Drug discovery & Screening
Genome	Editing for Cell Line Engineering: Application of CRISPR/Cas9 & other tools
14:30	Panel Discussion: CRISPR, ZFN, TALENS; Which technologies is the Best?
14:55	Explore the Genome editing application in mammalian cell line engineering
	<ul> <li>Application of mammalian cell line using CRISPR &amp; expression of CRISPR Cas9 nuclease</li> </ul>

- Efficient strategies for TALEN-based genome editing
- Integration of human AAVS1
- Knockout generation



15:20	Disease modeling in Zebrafish using Genome Editing Tools
	• Efficient protocols for Generation of zebrafish knockout mutants for genes involved in human genetic diseases using ZFNs,
	TALENs and CRISPR/Cas9
	<ul> <li>Comparison of ZFNS, TALENs and CRISPR/CAS9 mediated mutagenesis efficiency and types of mutations</li> </ul>
	Phenotype of knockout mutants
	<ul> <li>Status of targeted knock-in mutagenesis in zebrafish</li> </ul>
	Dr. Raman Sood,
	Director, Zebrafish Core facility, National Human Genome Research Institute, National Institutes of Health, USA
15:45	Afternoon Refreshment and Poster Presentation   One-to-One Networking Meetings
16:30	Application of Cell line engineering for Ex vivo Therapeutics
	<ul> <li>Cell line based therapeutics: Case study</li> </ul>
	<ul> <li>Cell-based therapies, protein drugs, gene therapies &amp; vaccine.</li> </ul>
16:55	Understanding the regulatory Issues of Genome Editing
	<ul> <li>Why regulation is necessary?</li> </ul>
	<ul> <li>International regulations for ZFN and TALEN application</li> </ul>
	<ul> <li>Is there any regulation of CRISPR/Cas application?</li> </ul>
	<ul> <li>Genome editing is under GMO regulation or not?</li> </ul>
	<ul> <li>Regulatory approaches for Therapeutic application of CRISPR</li> </ul>
17:20	Chairman's Closing Remarks & Conference Close
17.00	

17:30 Drinks Reception





8:00	Registration & Refreshment
8:20	Opening remarks from the Chair
8:30	Keynote Presentation:
	Title: TBA
	Dr. Matthew Porteus, Principal Investigator, Stanford School Of Medicine, USA
8:55	Solution provider's presentation: Keynote-2
Therape	utic Application of Genome Editing (CRISPR, ZFN, & TALEN)
9:25	<ul> <li>Elucidating Telomere Function in Human Tumor Biology</li> <li>Mutations in the human telomerase reverse transcriptase (TERT) promoter are the most frequent non-coding mutations in cancer.</li> <li>We used genome editing to engineer these mutations in pluripotent stem cells.</li> <li>Telomerase-expressing embryonic stem with the cancer associated TERT promoter mutations showed only a modest increase in TERT transcription with no impact on telomerase activity.</li> <li>However, upon differentiation into somatic cells, which normally silence telomerase, cells with TERT promoter mutations failed to silence TERT expression, resulting in increased telomerase activity and aberrantly long telomeres.</li> <li>We conclude from these studies that TERT promoter mutations are sufficient to overcome the proliferative barrier imposed by telomere shortening without additional tumor-selected mutations.</li> <li>Dr. Dirk Hockemeyer</li> <li>Principal Investigator, University Of California, Berkeley - USA</li> </ul>
9:50	Morning Refreshments & Poster Presentations
10:35	<ul> <li>Therapeutic in vivo delivery of CRISPR/Cas9 for next generation gene therapy</li> <li>Development and characterization of RNA therapy delivery systems</li> <li>RNA delivery in: in in vivo &amp; in vitro systems</li> <li>Discuss the successful RNA therapy in case of disease like-Viral infection, Hemophilia &amp; other genetic disease</li> <li>Dr. Hao Yin</li> <li>Research Scientist, David H. Koch Institute for Integrative Cancer Research, MIT, USA</li> </ul>
11:00	<ul> <li>Genome therapy for nucleotide repeat expansion-mediated neurodegenerative diseases</li> <li>Introduction of monogenic, neurodegenerative diseases caused by nucleotide repeat expansion: Muscular dystrophies, Spinocerebellar Ataxias, Motor neuron disease.</li> <li>Mechanism of RNA/protein gain-of-function from nucleotide repeat expansion.</li> <li>Strategies of genome manipulation for nucleotide repeat expansion diseases</li> <li>Genome therapy of Myotonic Dystrophy Type 1 iPS cells</li> <li>Prospect of in vivo genome therapy.</li> <li>Dr. Guangbin Xia</li> <li>Department of Neurology, College of Medicine, University of Florida, USA</li> </ul>





## 11:25 Exploring Genome Editing Platform for immunotherapy

- Engineering of human T Cells for therapeutic approach
- MegaTAL nucleases for stem cell and T cell therapies
- TALEN application for CAR-TCell generation
- Gene modification in T cells using editing tools

## 11:55 High-throughput gene targeting using CRISPR/Cas9 for human disease modeling in zebrafish

- High-throughput method of CRISPR/Cas9 gene editing in zebrafish
- Multiplex gene editing method
- Optimize sgRNA design for improved targeted efficiency
- Orthogonal Cas9 for expanding gene targeting coverage
- High-throughput phenotyping for studying function of human deafness genes in zebrafish.

### Dr. Gaurav K. Varshney

National Human Genome Research Institute - NIH, USA

## Genome Editing application for Drug Discovery & Screening: Use of CRISPR/Cas9 & other tools

## 12:20 Explore the use CRISPR/Cas9 for high throughput screening (HTS)

- How CRISPR/Cas9 is replacing existing technologies (e.g. RNAi)
- What are the advantages, limitations and challenges?
- What are the new types of screens that CRISPR/Cas9 enables (epigenetics, in-vivo, etc.)

#### Dr. Rob Howes

Associate Director, HTS, Antibody Discovery and Protein Engineering, MedImmune, UK

### 12:45 Lunch and Poster Presentation | One-to-One Networking Meetings

### 13:45 Application of Gene Editing Technologies in Drug Discovery

- How CRISPR fastens the early drug discovery process
- How Isogenic or gene-edited cell lines are used to identify the novel biologics

### 14:15 Functional Screening with Lentiviral shRNA and CRISPR Libraries

- Role of CRISPR/Cas9 Genome-Wide sgRNA Library Screening
- · Genome-wide loss-of-function screening to identify regulatory genes
- Therapeutic target identification

## Genome Editing application in Model organism

## 14:40 Discussing the Genome editing application towards animal models

- How to create Knockout, Transgenic, and Humanized Mice/ Rats/ Rabbits
- Understand the role of Zinc Finger Nuclease or CRISPR/Cas9 systems

## Prof. Radislav Sedláček

Director, Czech Centre for Phenogenomics (BIOCEV/IMG), Czech Republic

## 15:05 CRISPR mediated Genome editing in mice

- Discuss the CRISPR/Cas9 technology platform to build mice models
- Safety assessment of therapeutics in animal model

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15:30	Afternoon Refreshment and Poster Presentation   One-to-One Networking Meetings
16:15	<ul> <li>How Genome-Editing has Made the Generation of Genetically-Modified Animals Easier, and yet as Complicated as Ever</li> <li>Brief history of genetically modifying animals via Transgenics, Knockouts/Knockins in ES cells, Lentiviral transgenics, SCNT, spermatogonial stem cell transfer and the advantages and disadvantages of each</li> <li>Genome editing technologies: ZFN, TALENs and CRISPR evolution with GEMM models and their impact on generation of species which previously could not be modified (large animal transgenics for research and agricultural production)</li> <li>Design, validation and purification of CRISPR guides for your GEMM project</li> <li>Quality control for your CRISPR injection project and Preparation of CRISPR reagents</li> <li>Analysis of the "founder" FO CRISPR generation methods</li> </ul>
16:40	<ul> <li>Ethics &amp; Intellectual property rights (IPR) in Genome Editing</li> <li>Ethical concerns and controversies of Genome Editing technologies</li> <li>Argument towards patent of CRISPR-Cas9</li> <li>International laws towards IPR in Genome engineering patents</li> </ul>
17:05	Chair person's closing remarks & Conference close

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