

# CRISPR and Beyond: Perturbations at Scale to Understand Genomes

22-25 September 2020

## Virtual Conference Agenda

Tuesday, 22 September				
13:00		13:05		Welcome
<i>Scientific Programme Committee:</i> <a href="#">John Doench, Broad Institute, USA</a> <a href="#">Leopold Parts, Wellcome Sanger Institute, UK</a> <a href="#">Lea Starita, University of Washington, USA</a> <a href="#">Jolanda van Leeuwen, University of Lausanne, Switzerland</a>				
13:05		14:35		Session 1: Links to Disease
			Live	Introduction to the session <i>Chair: Leopold Parts, Wellcome Sanger Institute, UK</i>
13:05	13:25	20 mins	pre-recorded	Genome-wide characterisation of BRCA 1 functioning in DNA damage repair <a href="#">Sylvie Noordermeer, Leiden University Medical Centre, Netherlands</a>
13:25	13:45	20 mins	pre-recorded	Using Stem Cells to Explore the Genetics Underlying Complex Genetic Disease <a href="#">Kristen Brennand, Icahn School of Medicine at Mount Sinai, USA</a>
13:45	13:55	10 mins	pre-recorded	Whole genome CRISPR screen for entry of extracellular tau in human neurons <i>Lewis Evans, UCL Great Ormond Street, ICH, UK</i>
13:55	14:05	10 mins	pre-recorded	Transforming diagnostic confidence in developmental disorders using CRISPR-based saturation genome editing <i>Hong Kee Tan, Wellcome Sanger Institute, UK</i>
14:05	14:35	30 mins	Live	Q&A <i>Chair: Leopold Parts, Wellcome Sanger Institute, UK</i> <i>Moderator: Shondra Miller, St Jude Children's Research Hospital, USA</i>
14:35	14:55	20 mins	Break	
14:55		16:25		Session 2: Coding Variation
			Live	Introduction to the session <i>Chair: Leopold Parts, Wellcome Sanger Institute, UK</i>
14:55	15:15	20 mins	Pre-recorded	High throughput functional annotation of pharmacogene variants using yeast <a href="#">Maitreya Dunham, University of Washington, USA</a>
15:15	15:35	20 mins	Pre-recorded	Understanding the functional effects of coding variation, at scale <a href="#">Lea Starita, University of Washington, USA</a>
15:35	15:45	10 mins	Pre-recorded	Massively parallel assessment of human variants with base editor screens <i>Ruth Hanna, Broad Institute, USA</i>
15:45	15:55	10 mins	Pre-recorded	Large-scale phenotypic characterization of genetic variants of the DNA damage response using CRISPR-dependent base editing <i>Alberto Ciccia, Columbia University, USA</i>
15:55	16:25	30 mins	Live	Q&A <i>Chair: Leopold Parts, Wellcome Sanger Institute, UK</i> <i>Moderator: Shondra Miller, St Jude Children's Research Hospital, USA</i>
16:25	16:45	20 mins	Break	
16:45	17:00		Pre-recorded	Poster Session 1 lightning talks
17:00	18:00			Poster Session 1

# CRISPR and Beyond: Perturbations at Scale to Understand Genomes

22-25 September 2020

Wednesday, 23 September

13:00		14:30		Session 3: Genetic Screens	
				Live	Introduction to the session <i>Chair: Sylvie Noordermeer, Leiden University, Netherlands</i>
13:00	13:20	20 mins	Pre-recorded	CRISPR screening for cancer drug discovery <a href="#">Mathew Garnett, Wellcome Sanger Institute, UK</a>	
13:20	13:40	20 mins	Pre-recorded	Optimisation and meta-analysis of genome-scale signals <a href="#">Kenneth Baillie, University of Edinburgh, UK</a>	
13:40	13:50	10 mins	Pre-recorded	Major functional bias in genome-wide CRISPR screens for mitochondrial complexes <i>Maximilian Billmann, University of Minnesota, USA</i>	
13:50	14:00	10 mins	Pre-recorded	Temporal analysis of time-series CRISPR screen data using mixed linear models. <i>Lorena Sofia Lopez Zepeda, Berlin Institute of Medical Systems Biology, Germany</i>	
14:00	14:30	30 mins	Live	Q&A <i>Chair: Jolanda van Leeuwen, University of Lausaane, Switzerland</i> <i>Moderator: Sylvie Noordermeer, Leiden University, Netherlands</i>	
14:30	14:50	20 mins	Break		
14:50		16:20		Session 4: Rewriting DNA	
				Live	Introduction to the session <i>Chair: Tom Ellis, Imperial College London, UK</i>
14:50	15:10	20 mins	Pre-recorded	Genome-wide interrogation of gene functions through base editor screens empowered by barcoded sgRNAs <a href="#">Wensheng Wei, Peking University, China</a>	
15:10	15:30	20 mins	Pre-recorded	Recording cellular and molecular events in DNA <a href="#">Nozomu Yachie, University of British Columbia, Canada</a>	
15:30	15:40	10 mins	Pre-recorded	Pooled protein tagging, cellular imaging and in situ sequencing for monitoring drug action in real time <i>Andreas Reicher, CelMM Research Center for Molecular Medicine, Austria</i>	
15:40	15:50	10 mins	Pre-recorded	Characterization of synthetic single guide RNAs for CRISPR/Cas9 genome editing - An extensive evaluation of gRNA formats, purity, and delivery methods <i>Garrett Rettig, Integrated DNA Technologies, USA</i>	
15:50	16:20	30 mins	Live	Q&A <i>Chair: Lea Starita, University of Washington, USA</i> <i>Moderator: Tom Ellis, Imperial College London, UK</i>	
16:20	16:40	20 mins	Break		
16:40	17:15		Pre-recorded	Poster Session 2 lightning talks	
17:15	18:15			Poster Session2	

# CRISPR and Beyond: Perturbations at Scale to Understand Genomes

22-25 September 2020

## Thursday, 24 September

13:00		14:10		Session 5: Genetic Interactions	
		Live		Introduction to the session <i>Chair: Jolanda van Leeuwen, University of Lausanne, Switzerland</i>	
13:00	13:20	20 mins	Pre-recorded	Crosstalk between cellular metabolism and DNA repair <a href="#">Joanna Loizou, Research Centre for Molecular Medicine, Austria</a>	
13:20	13:30	10 mins	Pre-recorded	Comprehensive prediction of synthetic lethality between paralog pairs in cancer cell lines Barbara De Kegel, University College Dublin, Ireland	
13:30	13:40	10 mins	Pre-recorded	Understanding Chemical-Genetic Interactions <i>Loan Vulliard, CeMM Research Center for Molecular Biology, Austria</i>	
13:40	14:10	30 mins	Live	Q&A <i>Chair: Jolanda van Leeuwen, University of Lausanne, Switzerland</i> <i>Moderator: John Doench, Broad Institute, USA</i>	
14:10	14:30	20 mins	Break		
14:30		16:00		Session 6: Single Cell + Perturbation	
		Live		Introduction to the session <i>Chair: Shondra Pruett-Miller, St Jude Children's Research Hospital, USA</i>	
14:30	14:50	20 mins	Pre-recorded	Biochemistry at single-cell resolution: a new approach to understand functional heterogeneity <a href="#">Jay Hesselberth, University of Colorado School of Medicine, USA</a>	
14:50	15:10	20 mins	Pre-recorded	Human tissues & Cellular Phenotypes <a href="#">Sarah Teichmann, Wellcome Sanger Institute, UK</a>	
15:10	15:20	10 mins	Pre-recorded	Relating thousands of variants to function with massively parallel pooled genetic screens <i>Kathryn Geiger Schuller, Broad Institute of MIT and Harvard, USA</i>	
15:20	15:30	10 mins	Pre-recorded	Towards genome-scale CROP-seq screens <i>Tilmann Buerckstuegger, Aelian Biotechnology GmbH, Austria</i>	
15:30	16:00	30 mins	Live	Q&A <i>Chair: Lea Starita, University of Washington, USA</i> <i>Moderator: John Doench, Broad Institute, USA</i>	
16:00	16:20	20 mins	Break		
16:20	17:00		Pre-recorded	Poster Session 3 lightning talks	
17:00	18:00			Poster Session 3	

# CRISPR and Beyond: Perturbations at Scale to Understand Genomes

22-25 September 2020

Friday, 25 September

13:00		14:40		Session 7: Technology	
		Live		Introduction to the session <i>Chair: Mathew Garnett, Wellcome Sanger Institute, UK</i>	
13:00	13:20	20 mins	Pre-recorded	Tricks of the Trade: Functionalizing Genome Editing for a Broad Range of Cellular Targets <a href="#">Shondra Pruett-Miller, St Jude Children's Research Hospital, USA</a>	
13:20	13:40	20 mins	Pre-recorded	Exploring the flexibility of a eukaryotic genome using synthetic yeast <a href="#">Tom Ellis, Imperial College London, UK</a>	
13:40	13:50	10 mins	Pre-recorded	Massively parallel kinetic profiling of natural and engineered CRISPR nucleases <i>Stephen Jones, University of Texas at Austin, USA</i>	
13:50	14:00	10 mins	Pre-recorded	A benchtop platform for generating highly multiplexed variant libraries <i>Deanna Church, Inscripta, Inc. USA</i>	
14:00	14:10	10 mins	Pre-recorded	Discovery and characterization of transcriptional effectors with high-throughput protein domain screens in human cells <i>Josh Tycko, Bassik Lab, USA</i>	
14:10	14:40	30 mins	Live	Q&A <i>Chair: John Doench, Broad Institute, USA</i> <i>Moderator: Mathew Garnett, Wellcome Sanger Institute, UK</i>	
14:40	15:00	20 mins	Break		
15:00		16:10		Session 8: Non-coding Variation	
		Live		Introduction to the session <i>Chair: Lea Starita, University of Washington, USA</i>	
15:00	15:20	20 mins	Pre-recorded	CRISPR, cancer and long noncoding RNAs <a href="#">Rory Johnson, University College Dublin, Ireland</a>	
15:20	15:30	10 mins	Pre-recorded	Systematic characterisation of functional circular RNAs <i>Matt Neve, Monash University, Australia</i>	
15:30	15:40	10 mins	Pre-recorded	High-throughput screening of engineered microRNA target sites enables precise control of gene expression patterns in mammalian cells <i>Yale Michaels, University of British Columbia, Canada</i>	
15:40	16:10	30 mins	Live	Q&A <i>Chair: Lea Starita, University of Washington, USA</i> <i>Moderator: Jolanda van Leeuwen, University of Lausanne, Switzerland</i>	
16:10		16:20		Closing remarks	
<p><i>Scientific Programme Committee:</i>  <a href="#">John Doench, Broad Institute, USA</a>  <a href="#">Leopold Parts, Wellcome Sanger Institute, UK</a>  <a href="#">Lea Starita, University of Washington, USA</a>  <a href="#">Jolanda van Leeuwen, University of Lausanne, Switzerland</a></p>					