



**At the end of the Conference, the participants will be able to:**

- Identify diseases or disorders discussed which may benefit from genetic diagnosis or Precision Medicine.
- List examples of targeted or advanced therapeutic options for genetic diseases discussed.
- Name diseases and disorders that may be present in the Qatari population.

**Functional Genomics 2018 Target Audience**

- Physicians
- Nurses
- Pharmacists
- Allied Health

(Involved in the field of Biology, Genomics, Genomic Medicine, Genetics, Cancer and other Chronic Diseases.)

**SPC STATEMENT**

The Scientific Planning Committee has reviewed all disclosed financial relationships of speakers, moderators, facilitators and/or authors in advance of this CPD activity and has implemented procedures to manage any potential or real conflicts of interest.

**QCHP STATEMENT**

This Activity is an Accredited Group Learning Activity (Category 1) as defined by the Qatar Council for Healthcare Practitioners-Accreditation Department and approved for a maximum of 22 hours.

## Program Schedule

Day 1: Saturday 8 December 2018 "Human Genomes and Big Data"	
07:00 – 7:40	Registration and refreshments
<b>Introduction:</b>  <b>Objective:</b> <ol style="list-style-type: none"> <li>1. Able to discuss the role of Sidra Research and Functional Genomics for healthcare in Qatar.</li> <li>2. Describe the achievements of Sidra Research.</li> </ol>	
07:40 – 07:45	<b>Welcome note</b> Peter Morris CEO, Sidra Medicine - Qatar
07:45 – 07:55	<b>Overview of Sidra Research Achievements</b> Christof Von Kalle CRO, Sidra Medicine - Qatar
07:55 – 08:15	<b>The Functional Genomics Symposium Series</b> Xavier Estivill Co- Chair of the Functional Genomics Symposium 2018
<b>Session 1: The Message of Genes</b>  <b>Moderators: Nicole Soranzo and Puthen Jithesh</b>  <b>Objective:</b> <ol style="list-style-type: none"> <li>1. Describe lessons learned from large genome projects</li> <li>2. List examples of challenges encountered in decoding the genome for clinical purposes</li> <li>3. Explain advantages of population biobanks and big data projects</li> </ol>	
8:15 – 8:50	<b>"From Genomics To Therapeutics: Uncovering And Manipulating The Genetic Circuitry of Human"</b> <b>Manolis Kellis</b> MIT Computer Science, CSAIL, Broad Institute, Massachusetts, USA
8:50 – 9:25	<b>"Tohoku Medical Megabank Project- A National Challenge to Realize Personalized Medicine"</b> <b>Masayuki Yamamoto</b> Tohoku Medical Megabank Organization, Sendai, Japan
9:25 – 10:00	<b>"The Use of Population Biobanks for Translational Research"</b> <b>Cisca Wijmenga</b> Groningen University Medical Center, Groningen, Netherlands
10:00 – 10:15	<b>"Role of allele-specific RNA expression in human traits"</b> <b>Gaurav Thareja</b> Weill Cornell Medicine, Qatar
10:15 – 10:30	<b>"Incidence, Clinical Spectrum and Molecular Mechanisms of Permanent Neonatal Diabetes Mellitus in the State of Qatar"</b> <b>Sara Al-Khawaga</b> Sidra Medicine, Qatar
10:30 – 11:00	Coffee break
<b>Session 2: Genome Sequencing in Action</b>  <b>Moderator: Manolis Kellis and Stephan Lorenz</b>  <b>Objective:</b> <ol style="list-style-type: none"> <li>1. Discuss how the activities in Genome Sequencing Define the Dissection of Human Phenotypes.</li> <li>2. Explain the advantages of long-read sequencing of genomes</li> <li>3. Describe how multi-Omics approaches can be utilized for Precision Medicine</li> </ol>	
11:00 – 11:35	<b>"Genome Sequencing and Multi-Omics Phenotyping in Precision Medicine"</b> <b>Nicole Soranzo</b> Wellcome Trust Sanger Institute, Hinxton, UK
11:35 – 12:10	<b>"Long-Read Sequencing of Complex Genomes and Improved Structural Variation Characterization" *</b> <b>Evan E. Eichler</b> University of Washington School of Medicine, Seattle, USA
12:10 – 12:45	<b>"Investigating Cellular Fate Decision in Mouse Embryos by Single-cell RNA-Sequencing"</b> <b>Antonio Scialdone</b> Helmholtz Zentrum München, Munich, Germany
12:45 – 13:00	<b>"Exomes in a Clinical Setting: The Promise and the Delivery"</b> <b>Donald Love</b> Sidra Medicine, Doha, Qatar

13:00 – 13:15	<p><b>“Sidra's Pediatric Precision Medicine Program – Opportunities for Discovery and Collaborations”</b>  <b>Khalid Fakhro</b>  Sidra Medicine, Doha, Qatar</p>
13:15 – 14:45	<b>Lunch break</b>
<p><b>Session 3: Qatar Genome and Medical Knowledge</b></p> <p><b>Moderators: Asmaa Al-Thani and Richard O’Kennedy</b></p> <p><b>Objective:</b></p> <ol style="list-style-type: none"> <li>1. Able to explain the objectives of the Qatar Genome Program (QGP)</li> <li>2. Name of diseases and disorders discussed that may be present in the Qatari population</li> <li>3. Describe lessons gleaned from the first results of the QGP Analysis</li> </ol>	
14:45 – 15:05	<p><b>“Overview- Qatar Genome Project”</b>  <b>Said Ismail</b>  Qatar Genome Proram, Doha, Qatar</p>
15:05 – 15:25	<p><b>“The Biomedical Landscape of Genetic Variation in the Population of Qatar”</b>  <b>Xavier Estivill</b>  Sidra Medicine, Doha, Qatar</p>
15:25 – 15:45	<p><b>“The Genetic Architecture of Health and Disease-related Trait in Qatari Population”</b>  <b>Omar Albagha</b>  Hamad Bin Khalifa University, Doha, Qatar</p>
15:45 – 16:05	<p><b>“Exploring Pharmacogenetic Variants in the Qatari Population”</b>  <b>Puthen Jithesh</b>  Sidra Medicine, Doha, Qatar</p>
16:05 – 16:20	<p><b>“Extended Blood Group and Platelet Phenotype Prediction from Whole Genome Sequencing and Its Impact on RBC Transfusion-Related Alloimmunization in the State of Qatar”</b>  <b>Zohreh Tatari- Calderone</b>  Sidra Medicine, Doha, Qatar</p>
16:20 – 16:35	<p><b>“Rare Diseases in Qatar: Genomics and Proteomics of Heritable Muscle Disorders”</b>  <b>Alice Abdelaleem</b>  Weill Cornell Medicine, Doha, Qatar</p>
16:35 – 16:50	<p><b>“Genetics and Epigenetics linked to T2D functional pathways in Qataris”</b>  <b>Noha A. Yousri</b>  Weill Cornell Medicine, Doha, Qatar</p>
16:50 – 17:05	<p><b>“In Vivo and in Silico Models for Qatari Specific Classical Homocystinuria as basis for Development of Novel Therapies”</b>  <b>Gheyath Khaled Nasrallah</b>  Qatar University, Doha, Qatar</p>
17:05 – 17:20	<p><b>“Interactome mapping using All-vs-All sequencing (AVA-seq) method”</b>  <b>Nayra M. Al-Thani</b>  Weill Cornell Medicine, Doha, Qatar</p>
17:20 – 17:30	<b>Summary of First Day by Jithesh Puthen</b>
<p><b>Day 2: Sunday 9 December 2018</b>  <b>“Modelling and Treating Disease”</b></p>	
07:15 – 8:15	Registration and Refreshments
<p><b>Session 4: Modelling Human Disease</b></p> <p><b>Moderators: Luis Saraiva and Cisca Wijmenga</b></p> <p><b>Objective:</b></p> <ol style="list-style-type: none"> <li>1. Explain how functional genomics has been utilized for drug discovery</li> <li>2. Name the advantages of using organoids in the study of human biology and disease</li> <li>3. Describe how animal models can be used to study human disease variants</li> </ol>	
8:15 – 8:50	<p><b>“The Use of Functional Genomics to Accelerate Drug Discovery”</b>  <b>Nicholas Katsanis</b>  Duke University, North Carolina, USA</p>
8:50 – 9:25	<p><b>“Long noncoding RNAs: A Potential Gold Mine of New Disease Genes”</b>  <b>Rory Johnson</b>  University of Bern, Bern, Switzerland</p>
9:25 – 10:00	<p><b>“Liver Organoids for the Study of Human Biology and Disease”</b>  <b>Meritxell Huch</b>  University of Cambridge, Cambridge, UK</p>
10:00 – 10:35	<b>“Organoid co-culture system to study gut brain signaling”</b>

	<b>Diego V. Bohórquez</b> Duke University, North Carolina, USA
10:35 – 10:50	<b>“Hypertrophic cardiomyopathy-linked variants of cardiac myosin binding protein C3 display altered molecular properties and actin interaction”</b> <b>Sahar Da’as</b> Sidra Medicine, Doha, Qatar
10:50 – 11:20	<b>Coffee Break</b>
<b>Session 5: Exploring the Consequences of Mutation in Model Systems</b>	
<b>Moderators: Edward Stuenkel and Anne Ferguson-Smith</b>	
<b>Objective:</b>	
<ol style="list-style-type: none"> <li>1. List the advantages of utilizing animal models for gene discovery</li> <li>2. Describe more about the impact of non-self mutations in human disease</li> <li>3. Explain what spatial transcriptomics can reveal about biology</li> </ol>	
11:20 – 11:55	<b>“RNA Toxicity in Huntington’s Disease: A Therapeutic Target in Polyglutamine Diseases”</b> <b>Eulalia Marti-Puig</b> University of Barcelona, Barcelona, Spain
11:55 – 12:30	<b>“Non-self Mutations: Neurodegenerative Diseases have Genetic Hallmarks of Autoinflammatory Disease”</b> <b>Robert Richards</b> University of Adelaide, Adelaide, Australia
12:30 – 13:05	<b>“Gene Discovery Mediated by the Drosophila Model Organism Screening Center”</b> <b>Hugo Bellen</b> Baylor College of Medicine, Texas, USA
13:05 – 14:30	<b>Lunch Break</b>
14:30 – 15:05	<b>“When exomes are not enough: Approaches to working through unresolved neurological cases”</b> <b>Elizabeth Ross</b> Weill Cornell Medicine, New York, USA
15:05 – 15:20	<b>“Spatial transcriptomics of olfactory receptors for high throughput mapping of olfactory bulb glomeruli”</b> <b>Kevin Zhu (Travel Award)</b> Duke University, North Carolina, USA
<b>Session 6: Inaugural Sidra Medicine Plenary Lecture</b>	
<b>Moderators: Donald Love</b>	
<b>Objective:</b>	
<ol style="list-style-type: none"> <li>1. Describe what is Duchenne Muscular Dystrophy (DMD) and some of the challenges of DMD treatment</li> <li>2. List a few of the current DMD therapeutic approaches</li> <li>3. Discuss insights and advice on how to excel as a female in science</li> </ol>	
15:20 – 16:05	<b>“Genetic Approaches to Therapy for Duchenne Muscular Dystrophy”*</b> <b>Dame Kay Davies</b> Oxford Neuromuscular Centre, Oxford, UK
<b>16:05 – 17:25 Women in Science Workshop</b>	
Moderator : Bernice Lo and Kholoud Al- Shafai	
<b>Objective: Able to demonstrate to all Women scientists of Qatar how they can build up a successful scientific career from women who achieved top level positions in the scientific world.</b>	
<b>Panelists:</b>	
<ul style="list-style-type: none"> <li>▪ <b>Dame Kay Davies</b> Professor of Genetics Oxford Neuromuscular Centre, Oxford, UK</li> <li>▪ <b>Elizabeth Ross</b> Professor of Neurology and Neuroscience Weill Cornell Medicine, New York, USA</li> <li>▪ <b>Elizabeth Phimister</b> Deputy Editor The New England Journal of Medicine, Boston, USA</li> <li>▪ <b>Souhaila Al Khodor</b> Investigator Sidra Medicine, Doha, Qatar</li> <li>▪ <b>Mariam Ali Al Ali Al Maadeed</b> Vice President for Research &amp; Graduate Studies</li> </ul>	

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17:25-17:30	<b>Conclusion of Second Day by Xavier Estivill</b>
<b>Day 3: Monday 10 December 2018</b> <b>"Mother and Child Health"</b>	
07:15 – 8:15	Registration and Refreshments
<b>Session 7: From Models of Cells to Disease Phenotypes</b>	
<b>Moderators: Nicholas Katsanis and Khalid Fakhro</b>	
<b>Objective:</b>	
<ol style="list-style-type: none"> <li>1. Discuss lessons in the development of embryos learned from RNA sequencing</li> <li>2. Describe genetic regulators involved in maternity/pregnancy</li> <li>3. List technologies or approaches used for prenatal/carrier screening</li> </ol>	
08:15 – 08:50	<b>"Genetic Regulators of Maternal Immune Tolerance and Pregnancy Success"</b> <b>Sarah Robertson</b> Robinson Research Institute, Adelaide, Australia
08:50 – 09:25	<b>"New Applications of Genomic Medicine in Women's Health"</b> <b>Lee Schulman</b> Feinberg School of Medicine of Northwestern University, Illinois, USA
09:25 – 09:55	<b>"Pathologic Glucocorticoid Receptor Mutations: From their Clinical Manifestations to Structural Impact on the Glucocorticoid Receptor Protein Toward Future Tailored Treatment Using Modified Glucocorticoid Ligands"</b> <b>Tomoshige Kino</b> Sidra Medicine, Doha, Qatar
09:55 – 10:20	<b>"The LINC Complex is Essential for Gametogenesis"</b> <b>Henning Horn</b> Hamad Bin Khalifa University, Doha, Qatar
10:20 – 10:35	<b>"Immune Functional Studies of a Novel Pathogenic STK4 Genotype: A Case Report"</b> <b>Andrea Guennoun</b> Sidra Medicine, Doha, Qatar
10:35 – 10:50	<b>"Characterization of the Microbial Diversity in the Milk of Mexican Women"</b> <b>Karina Corona Cervantes (Travel Award)</b> Genetics and Molecular Biology Department, Mexico City, Mexico
10:50 – 11:20	<b>Coffee Break</b>
<b>Session 8: Diagnosis of the Future Child</b>	
<b>Moderators: Evan Eichler and Meritxell Huch</b>	
<b>Objective:</b>	
<ol style="list-style-type: none"> <li>1. Describe implications of epigenetics on human biology</li> <li>2. Discuss lessons gleaned from the Undiagnosed Rare Disease Program of Catalonia</li> <li>3. Discuss the findings from the noninvasive prenatal testing study at HMC</li> </ol>	
11:20 – 11:55	<b>"Variable Silencing of the Repeat Genome - Implications for Non-Genetic Inheritance"</b> <b>Anne Ferguson-Smith</b> University of Cambridge, Cambridge, UK
11:55 – 12:30	<b>"The Renaissance of Genomic Medicine"</b> <b>Stylianos Antonarakis</b> University of Geneva Medical School, Geneva, Switzerland
12:30 – 13:05	<b>"The Results of the Undiagnosed Rare Disease Program of Catalonia (URDCat)"</b> <b>Luis Perez-Jurado</b> Pompeu Fabra University, Barcelona, Spain, and University of Adelaide, Australia
13:05 – 13: 20	<b>"A Validation Study: NIPT to Identify Pregnancies at high Risk for Aneuploidies Using NGS Platform with Minimum of 2% Fetal Fraction"</b> <b>Sarmad Ali Ghulam Shabir</b> Hamd Medical Corporation, Doha, Qatar
13:20 – 14:45	<b>Lunch Break</b>

**Session 9: Patient Genome Sequencing**

**Moderators: Stylianos Antonarakis and Bernice Lo**

**Objective:**

- 1. Describe how molecular diagnosis and basic research findings can lead to novel therapies for Precision Medicine**
- 2. Distinguish between the molecular methods/assays for different types of mutations**
- 3. Describe the importance of and role of a genetic counselor in genomic research studies**

14:45 – 15:20	<b>“Genomics Research Uncovers the Primary Role of Complement Pathway in a Familial Form of Protein Losing Enteropathy”</b> <b>Ahmet Ozen</b> Marmara University, Istanbul, Turkey
15:20 – 15:55	<b>“Returning Results from Genomic Research: NIAID’S Experience”</b> <b>Leila Jamal</b> National Institute of Health, Bethesda, Maryland, USA
15:55 – 16:30	<b>“A Nuts and Bolts Approaches to Diagnosis of SCID Patients in the Era of Newborn Screening”</b> <b>Mehdi Adeli</b> Sidra Medicine and Hamad Medical Corporation, Doha, Qatar
16:30 – 16:45	<b>Mutation in the GTPase GIMAP6 Leading to Reduced Autophagy in a Severely Immune Deficient Patient”</b> <b>Brittany Chao (Travel Award)</b> Oxford University, Oxford, UK
16:45 – 17:20	<b>“Clinical Genomic: Discovery and Building Genetics Models for Complex Traits”</b> <b>James R. Lupski</b> Baylor College of Medicine, Texas, USA
17:20 – 17:30	<b>Poster Awards by Rashid Al Ali</b>
17:30 – 17:35	<b>Closing Remarks by Donald Love</b>