

Sequencing Seminar Series

Sequencing Simplified

Illumina sequencing is fuelling groundbreaking discoveries across a wide range of research areas. By enabling the broadest range of applications including whole-genome resequencing, de novo sequencing, targeted resequencing, transcriptome and epigenetic analysis, researchers are publishing at amazing rates.

This seminar series will showcase presentations from scientists using Illumina technology to power their next-generation sequencing studies. We invite you to join us, learn about their research, and receive an update on new sequencing products from Illumina.

TUESDAY, SEPTEMBER 7, 2010

**CROWNE PLAZA MANCHESTER CITY CENTRE
70 SHUDEHILL
MANCHESTER, M4 4AF ENGLAND
SMITHFIELD SUITE**

SEMINAR SCHEDULE		
10:00	Registration	
10:30	Introduction	Illumina
10:35	The Expanded Illumina Sequencing Portfolio, New Sample Prep Solutions and Workflow	Marcus Hausch, Ph.D., Illumina
11:00	The Compendium of Genomic DNA Sequencing Applications	Stephanie Brooking, Illumina
11:30	Sequencing Cancer Genomes	Phil Stephens, Ph.D., Cancer Genome Project, Wellcome Trust Sanger Institute
12:00	Medip-Seq, Profiling the Cancer Methylome	Andrew Feber, Ph.D., University College London
12:30	LUNCH	
1:30	Transcriptome Sequencing: The Switch from Arrays to Sequencing	Shawn Baker, Ph.D., Illumina
2:00	Diagnostic Applications of Clonal Sequencing Using the Illumina GAI Platform	Graham Taylor, Ph.D., Leeds Institute of Molecular Medicine
2:30	Application of Illumina Sequencing to the Genetics of Human Cancers	Gordon Blackshields, Ph.D., Source BioScience
3:00	Illumina NGS Sequencing Analysis Process: Turning Data Into Biology	Lukas Smink, Illumina
3:30	Closing Remarks	Illumina

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