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 14, 2010

COEX INTERCONTINENTAL HOTEL
524 BONGEUNSARO
GANGNAMGU, SEOUL 135-975 KOREA
MEETING ROOM TBD

SEMINAR SCHEDULE		
9:30	**Registration**	
10:00	**Introduction**	**Woochul Kim, BMS**
10:10	**The Expanded Illumina Sequencing Portfolio, New Sample Prep Solutions and Workflow**	**KyungA Kim, BMS**
10:40	**A Very Small Genome Centre's Adoption of New Generation DNA Sequencing for Research and Core Service**	**Professor Si Lok, Genome Research Centre, The University of Hong Kong**
11:20	**Highly Annotated Whole-Genome Sequences of Korea Individuals**	**Professor Jongil Kim, Seoul National University, College of Medicine**
12:00	**Transcriptome Sequencing: The Switch from Arrays to Sequencing**	**Chris Streck, Illumina**
12:40	**Lunch**	
14:00	**Bioinformatics Analysis of RNA-seq Data**	**Professor Sanghyuk Lee, Korean Bioinformation Center (KOBIC)**
14:40	**Noninvasive Prenatal Diagnosis of Fetal Down Syndrome by Maternal Plasma DNA Sequencing**	**Professor Rossa Chiu, The Chinese University of Hong Kong**
15:20	**Coffee Break**	
15:40	**IlluminaCompute – An Informatics Ecosystem for Production Scale Sequencing**	**Martin Burke, Illumina**
16:20	**Closing Remarks**	**Hyeyoung Lee, Illumina**

Register now at:
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