Lunch and Learn: 3D Epigenetics Tools for Multi-Omics Research

Hosted by Arima Genomics and Illumina

Summary:
Learn how epigenetics techniques such as Arima-HiC and HiChIP can be part of a multi-omics research approach to enable new discoveries. Hi-C is an exciting technique which uses Illumina Next-Generation Sequencing to investigate chromatin conformation state.

Please attend for epigenetics-focused presentations and discussions lead by Arima Genomics and Illumina. Lunch will be provided.

Time:
March 27th, 12:00PM-1:00PM

Location:
CIEMAS Building Room 2240

Agenda:

Arima-HiC for 3D Chromatin Conformation Research

Jay Clark, Epigenetics Project Consultant, Arima Genomics

Utilizing Epigenetic Tools in a Multi-Omics Sequencing Approach

Matthew Angel, Sequencing Specialist, Illumina

Please RSVP to jay@arimagenomics.com to be marked down for lunch.