

次世代定序資料分析軟體 Partek 工作坊



TechComm
科技共同空間

時間：8/19 (二) 9:30-17:00 PM (英文授課，提供筆記型電腦實習)

地點：生科館六樓 628 會議室

講者：Erwin Tantoso (Partek specialist)

名額有限，請先上網報名 <http://ppt.cc/xw7~>



時間	議程內容
9:30-10:30	<ol style="list-style-type: none">1. Alignment Basics with Partek Flow2. Interpreting QAQC Results Before and After Alignment; Optimizing Alignment Parameters; Interpreting Quantification Results
10:45-12:00	<ol style="list-style-type: none">1. RNA-Seq Analysis with Partek Genomics Suite2. Finding Differentially Expressed Lists of Genes or Transcripts, Alternative Splicing, Gene Ontology; Creating Plots and Graphs for Publication
13:00-14:00	<ol style="list-style-type: none">1. RNA-Seq Analysis with Partek Pathway2. Finding and Visualizing Enriched Pathways from RNA-Seq Data
14:00-15:30	<ol style="list-style-type: none">1. Exome-Sequencing with Partek Genomics Suite2. Finding and Annotating Variants from Exome-Seq Data
15:45-17:00	<ol style="list-style-type: none">1. ChIP-Seq with Partek Genomics Suite2. Detecting and Visualizing Regions of Transcription Factor Binding Sites from ChIP-Seq Data and Integration with Gene Expression Data



Software for All Your
NGS & Microarray Data



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系統生物學研究中心
Center for Systems Biology, National Taiwan University

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