## **Practicalities of 'Omics**

## 15 February 2016, Lindisfarne Room, Newcastle University

Draft Programme

10.00 - 11.00	Registration
11.00 – 11.10	Opening remarks Deb Henderson, Dean of Research & Innovation
11.10 – 12.00	Keynote: Iain Macaulay, The Genome Analysis Centre Integrated (epi)genome and transcriptome sequencing from single cells
12.00 - 12.10	Break
12.10 – 12.40	Jon Higgins, Newcastle University A microscopy-based high content RNAi screen for histone kinases in mitosis
12.40 – 13.10	Conor Lawless, Newcastle University Interpreting genome-wide Quantitative Fitness Analysis (QFA) screens
13.10 – 14.45	Lunch Sponsored by Beckman Coulter
14.45 – 15.35	Keynote: Jerry Thomas, University of York What a proteomics lab can do for you - and vice versa
15.35 – 15.45	Break
15.45 – 16.15	Nick Morrice, Sciex Newcastle's new LCMSMS System: SWATH and the TripleTOF 6600
16.15 – 16.45	Simon Cockell, Newcastle University Principles and Practicalities of Data Integration
16.45 – 16.55	Closing Remarks

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