

Programme

2:00	Renaud Gaujoux, Technion, Haifa <i>Gene expression deconvolution in practice</i>
3:00	Ruth Luthi-Carter, University of Leicester <i>Changes in cell populations during neurodegeneration: Rationale for the development of 'Population-Specific Expression Analysis (PSEA)'</i>
3:30	<i>Coffee Break</i>
4:00	Alberto Capurro, University of Leicester <i>Implementation of Population-Specific Expression Analysis and discussion of potential methodological improvements</i>
4:30	Andrew Simpkin, NUI Galway <i>Cell type heterogeneity in early life: results from a UK birth cohort</i>
5:00	Cathal Seoighe & Barbara Martinelli, NUI Galway <i>Inference of methylation haplotypes for computational deconvolution of DNA methylation data</i>
5:30	<i>Close</i>