

# The Otago Genomics Facility Microarray & Technologies Meeting Friday February 8<sup>th</sup>, 2008



University of Otago, Department of Biochemistry, Room 228, 710 Cumberland Street, Dunedin, NZ



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Ian Campbell, Peter McCallum Institute, Melbourne  
Cancer gene discovery using ultra-high density SNP arrays

Paul O'Sullivan, Pacific Edge Biotechnology  
Bladder Cancer Diagnostics: from Microarrays to Clinical Trials

Matt Anaka, The Ludwig Institute, Australia  
Wilms tumor histogenesis is determined by the precursor cells,  
rather than constitutive epigenetic errors

Jo Stanton, University of Otago  
High Throughput Sequencing at Otago University using the GS FLX

Lesley Collins, Massey University  
Solexa Analysis Pipeline and Downstream Genomics: Bioinformatics  
for Solexa Sequencing

Mik Black, University of Otago  
High throughput sequencing data analysis

Allen Rodrigo, University of Auckland  
A new client for GRID computing

Caroline Beck, University of Otago  
Identification of regeneration specific genes in amphibian hind  
limb regeneration

Julia Horsfield, University of Otago  
When development comes unstuck: investigating chromosome  
cohesion syndromes in zebrafish

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