Bioinformatics Support

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Bioinformatics Support Unit

- Data analysis for life science
 - Collaboration
 - Analysis as a service
 - Training
- 7 members of staff
 - Mostly experienced, post-doctoral level bioinformaticians
- >50 papers published in last 5 years
- Collaborate on grants worth over £50m

The Bioinformatics Support Unit



Dr Simon Cockell Director



Dr Sirintra Nakjang NICR ESO



Robert Stones Core Facility ESO



Andrew Skelton MRG ESO



John Casement Core Facility Technician



Dr Graham Smith ICM ESO



Dr Rachel Queen NUSCU ESO

Academic Lead – Prof Anil Wipat

Bioinformatics Analysis

- Sequencing
 - Whole genome sequencing
 - Whole exome sequencing
 - RNA-Seq
 - ChIP-Seq
 - and many more...
- Microarrays
 - Expression
 - Methylation
 - Genotyping
- Proteomics
- Flow/Mass Cytometry



Bioinformatics Training

- Short courses run regularly
- Introduction to Practical Bioinformatics (2 days)
 - Linux command line
 - Scripting with BASH
 - R and Bioconductor
- RNA-Seq Analysis (1 day)
 - Alignment-free Quantification
 - Analysis using R
- Other courses being planned (DNA variation analysis, ChIP-Seq analysis, etc)
- <u>http://bsu.ncl.ac.uk/support/courses</u>



Elucidating Pathways of Steatohepatitis

- 4 year H2020 award
- Multi-omics data from core patient cohort
- RNA-Seq
- Methylation (Illumina EPIC arrays)
- Data integration and multi-omics analysis
 - eQTL
 - mQTL





Psoriasis Stratification to Optimise Relevant Therapy

- MRC Stratified Medicine award
- Large cohort RNA-Seq
- Investigating drug response
 - Range of biologic treatments of psoriasis
- Systems biology modelling of drug response





Metatranscriptomics on oral biofilms

- Industrially-funded oral microbiome investigation
- Meta RNA-Seq from 18 samples
 - Effect of treatment on oral biofilms
- Analysis of DEGs within individual species
- Leading to cross-species functional analysis
- Also annotating uncharacterised reads

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