## WEHI Proteomics Facility

We employ the latest mass spectrometry innovations at the interface between basic and clinical research from experimental design, sample preparation, data acquisition, through advanced biostatistics.



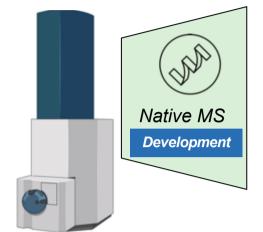
Scan the QR code to see more about WEHI proteomics facility



Global Proteome Profiling
Differential protein expression
Subcellular fractions (e.g secretome)



XL-MS (Crosslinking MS)
Structural proteomics
Protein dynamics & complexes



Impact II Q-TOF

timsTOF Pro x2 (Bruker)



DIA

Clinical Global

Proteomics proteomics

DIA

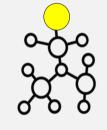
DIA = \( \mathred{\psi} \)
TurboID IP, F

(Bruker)

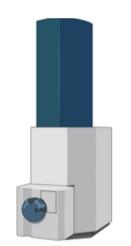
FLAG-tag



Clinical biofluid proteomics
Plasma, EVs, patient biofluids
Paired MS + Olink analyses\*



PTM & Phosphoproteomics
PTM analysis (e.g. GlyGly)
Quantitative Phosphoproteomics

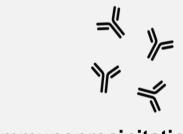


Intact MS 'top down'

Orbitrap Eclipse (Thermo Fisher Scientific)



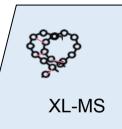
Cell surface labelling
Cell surface markers
Cell surface interactions

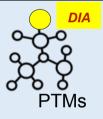


Immunoprecipitation-MS\*

Protein-protein interaction networks

\*Includes HA-tagged, FLAG-tagged











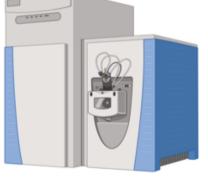
Intact 'Top Down' MS

Protein dynamics, Protein complexes

Protein isoforms (PTMs), Protein IDs



Proximity Biotin Labelling
In situ interactomics (TurbolD)
Protein-protein interaction networks



maXis II Q-TOF

(Bruker)

Q-Exactive Classic (Thermo Fisher Scientific)



Low-Input LCM Proteomics

Development

DIA

DIA

DIA

DIA

DEVELOPMENT

Development

Small Molecule

Development

Orbitrap Astral (Thermo Fisher Scientific)



Antibody de novo sequencing
Antibody characterisation
Epitope mapping



Low-input/LCM Proteomics
Low-input, high-sensitivity
Laser-capture microscopy samples

## **Contact WEHI Proteomics**



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