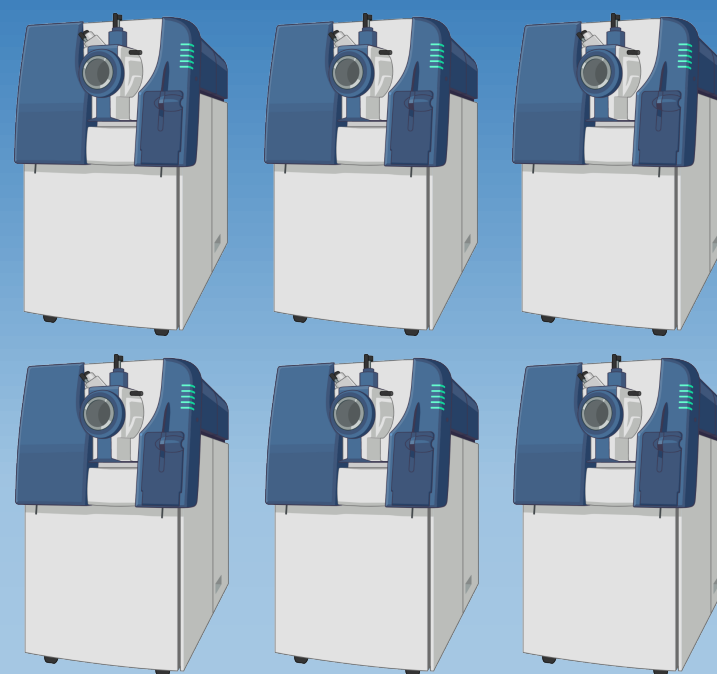
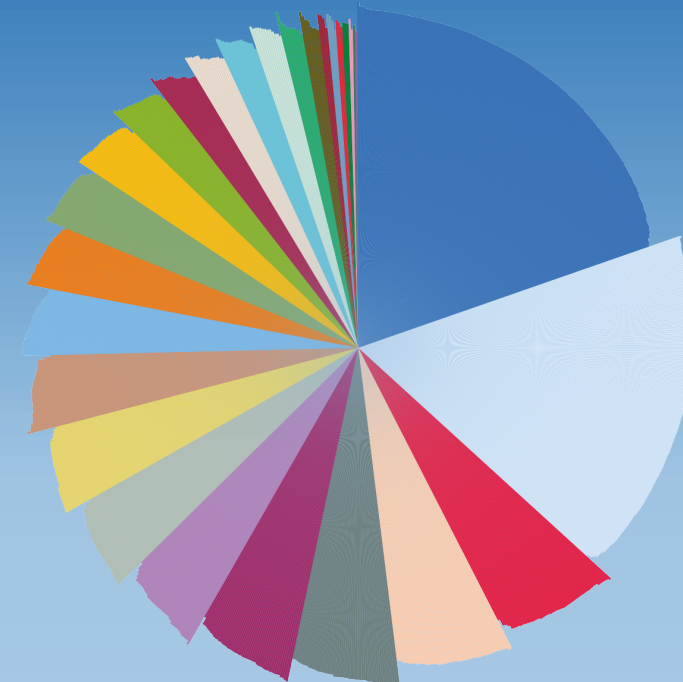


949 human
cancer cell lines

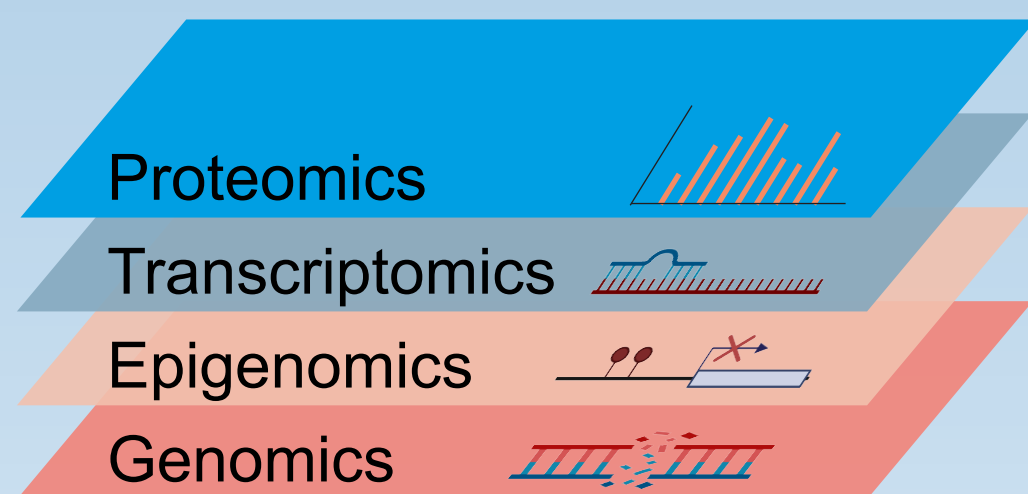


Six replicate proteomes across
six mass spectrometers



28 tissue types
> 40 cancer types
8,498 proteins

Multi-omic datasets



Cell-type
specificity

Post-transcriptional
regulation

**Biological
insights**

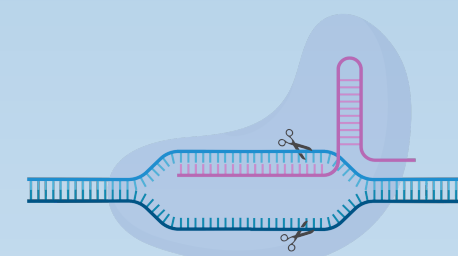
Protein-protein
interactions

Proteogenomic
integration

Cancer vulnerability screens



Sensitivity to
625 drugs

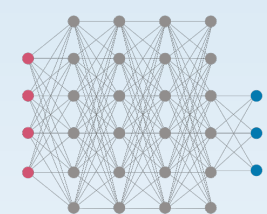


Dependency map
of 17,486 genes

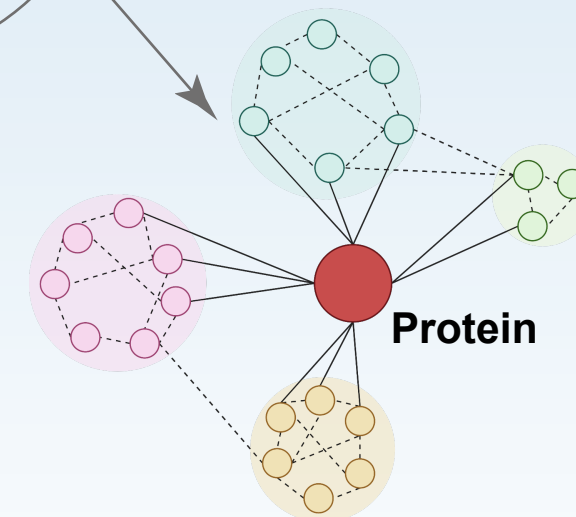


DeeProM

Deep Proteomic Marker



Protein
biomarkers



Co-regulated
protein networks