

## Protein-Protein Interactions

### AP-MS Workflows

- Affinity purification-mass spec
- Pull-down interacting partners
- Identify protein complexes

### Proximity Labeling

- BiOID, APEX, TurboID
- Spatial proteomics
- In vivo labeling

### Cross-linking MS

- Chemical cross-linkers
- Distance constraints
- Protein structure information

### Network Construction

- Interaction databases
- Scoring significance
- Pathway analysis