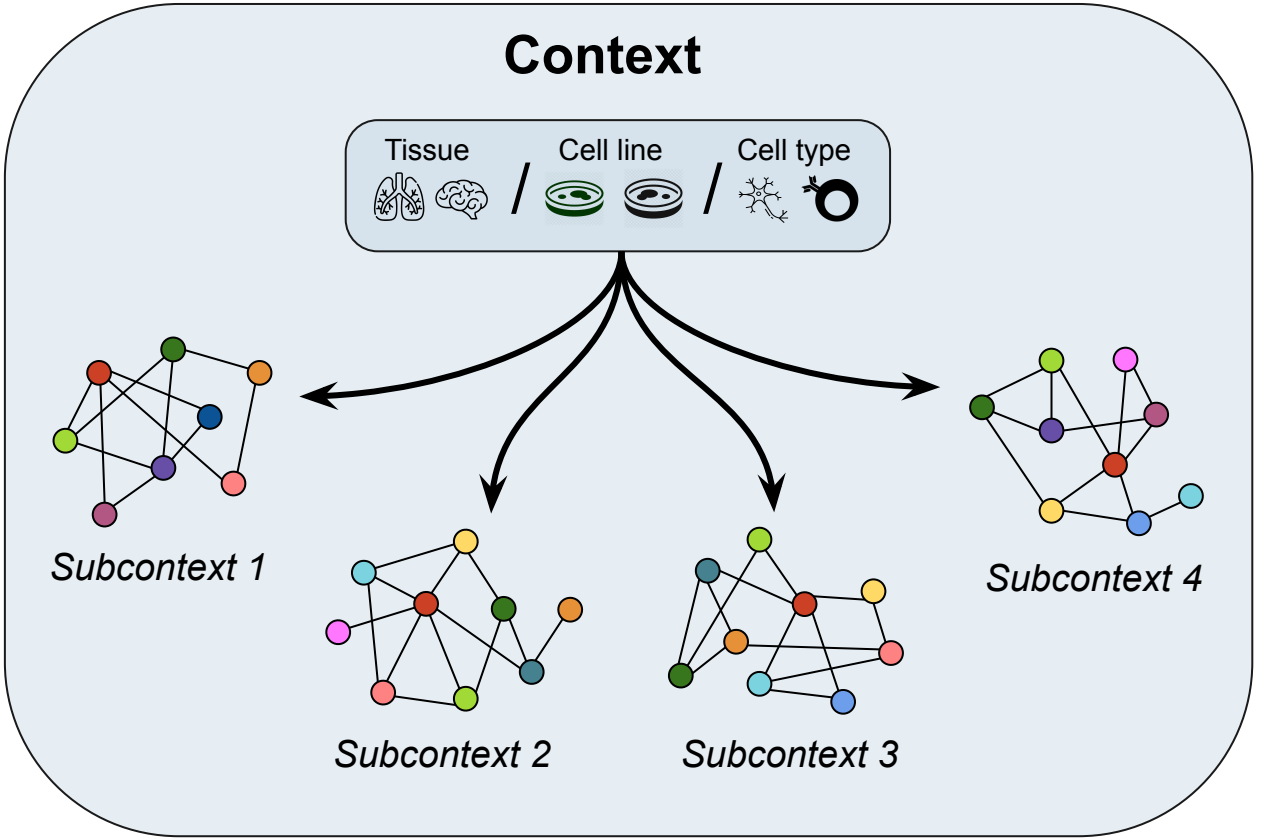

Analyses

Protein-level analysis

- Most common nodes →
- Least common nodes →
- Node degree
  - Subcontext 1 → :3, :4, :2, …

Network-level analysis

- Unique edges →
- Most common edges →
- Node neighbors
  - Subcontext 1 → :(, ,), …
- Permuted networks
  - Subcontext 1 → , , …
- Edge-based similarity →

Pathway-level analysis

- Pathway-based similarity →