# Ning Zhang

## Attributed Graph Alignment

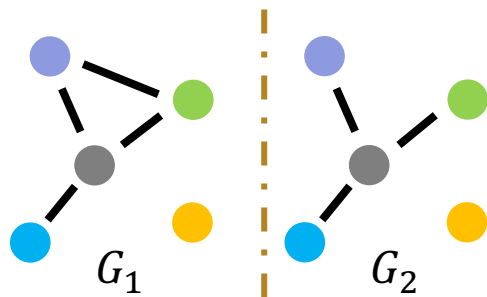Program: MSc in Information Theory

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Fig1. Protein-protein interaction networks (PPI networks)

**Existing model: graph alignment**
- Given $G_1 = (V_1, E_1)$ , $G_2 = (V_2, E_2)$
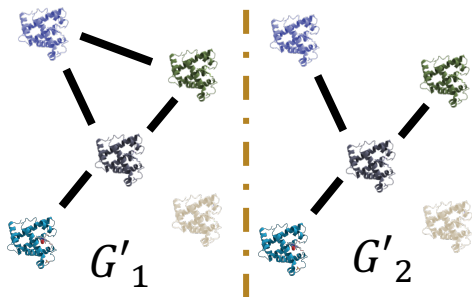- Find bijection between $V_1$ and $V_2$



Fig2. PPI networks with amino acid sequence of each protein

**Proposed model: attributed graph alignment**
- Given $G'_1 = (V_1, E_1, A_1)$ , $G'_2 = (V_2, E_2, A_2)$
- Find bijection between $V_1$ and $V_2$

**Goal:** quantify the benefit brought by attributes