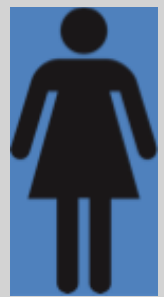

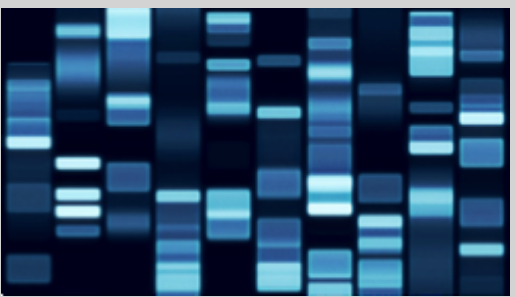
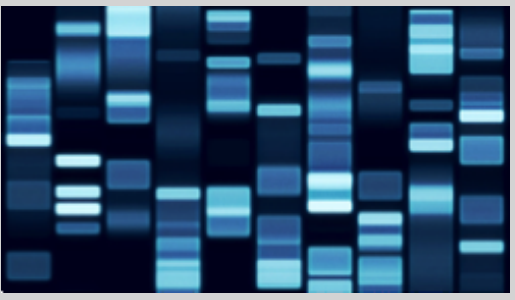


Coherent sub-networks in genome-scale metabolic models

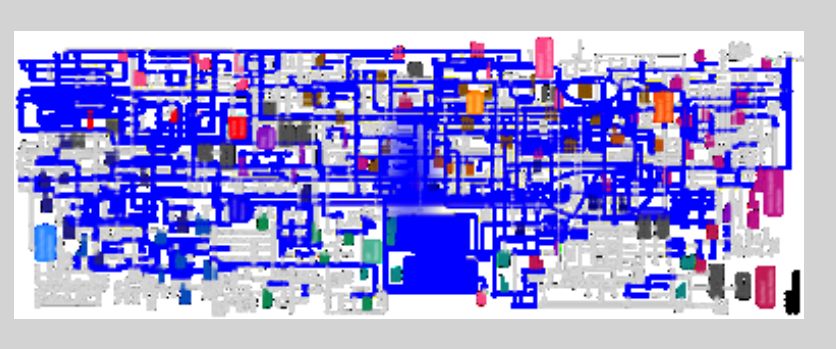
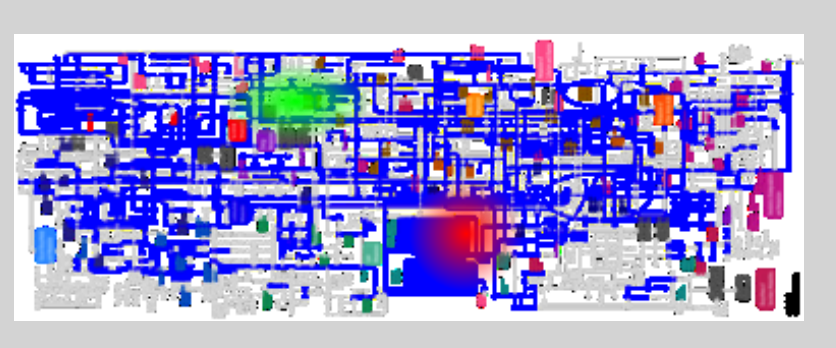
patient

mutations

metabolic model

http://www.metabolicatlas.org/

- 917 patients
- $|V| \sim 4000$
- $|E| \sim 15000$

**Aim** Find **dense common subgraphs** in patients with specific markers e.g.  $mutbrca = 1$  and  $mutp53 = 1$

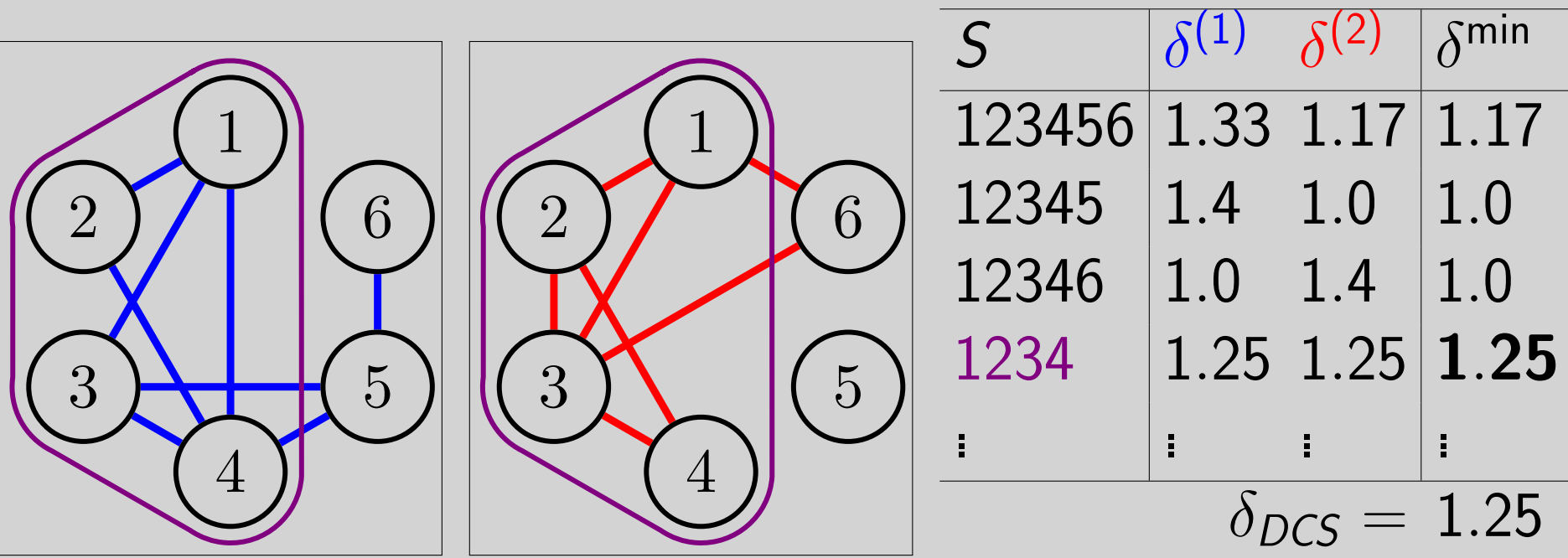
Existing methods do not scale [jiang2009mining; li2011integrative ]

Dense Common Subgraph (DCS) problem

**DCS** Given relational graph set  $G^{(1)} = (V, E^{(1)})$ ,  $G^{(2)} = (V, E^{(2)})$ , ...,

$$\delta_{DCS} = \max_{S \subseteq V} \min_{G^{(m)}} \frac{\#\{\text{edges induced by } S \text{ in } G^{(m)}\}}{|S|}$$

**Example**



References

- GoldbergG84
- charikar2000greedy
- jiang2009mining
- li2011integrative