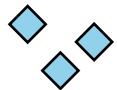
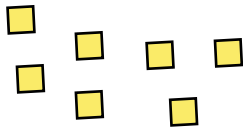


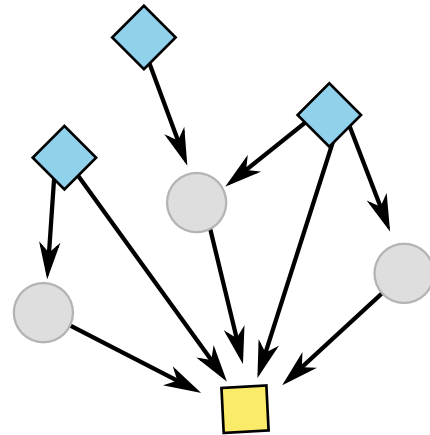
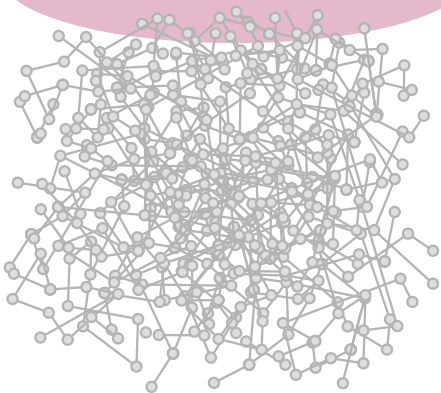
S_0 :
Initial
Proteins



S_1 :
Query
Proteins



Interactome



For each query protein t , compute
50 S_0 -to- t highest-scoring paths