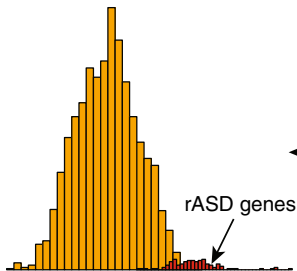


A

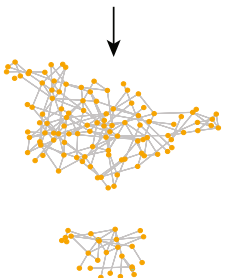
Co-expression connects
nodes (genes) in a network

**B**

Assign TADA scores to nodes
(Strong score : orange)

**D**

Identify risk (rASD) genes
among nASD genes using
FDR on TADA scores

C

Clusters of nodes with strong
scores identify network
(nASD) genes