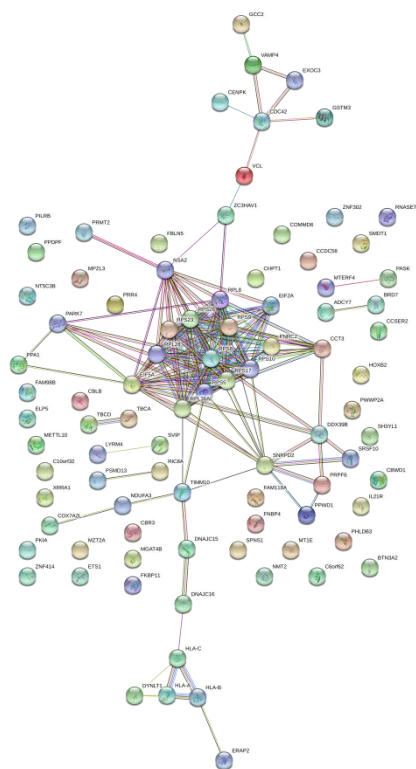


(A)

eQTL: gene-network of top-100 genes

Number of nodes:90

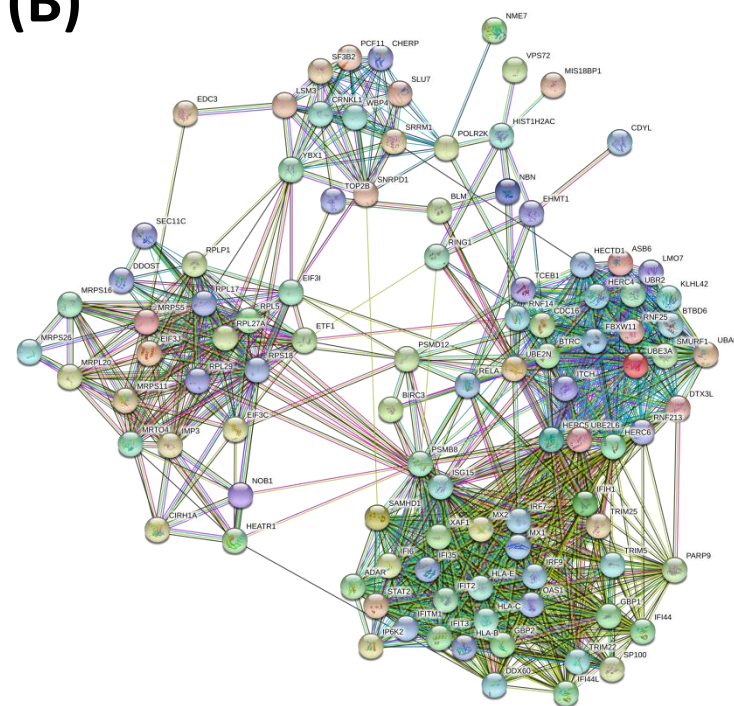
Number of edges:147

Average node degree:3.27

Avg. local clustering coefficient:0.427

Expected number of edges:65

PPI enrichment p-value:<1.0e-16

(B)

nQTL: gene-network top-100 genes

Number of nodes:103

Number of edges:1024

Average node degree:19.9

Avg. local clustering coefficient:0.746

Expected number of edges:220

PPI enrichment p-value:< 1.0e-16