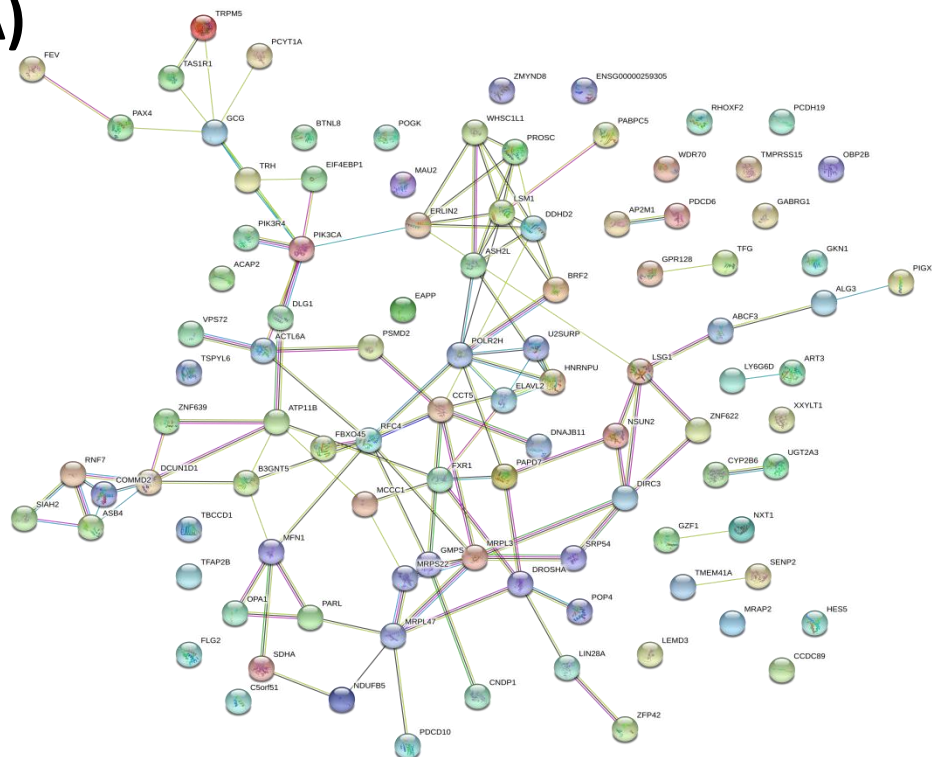


(A)

eQTL: gene-network of top-100 genes

Number of nodes:100

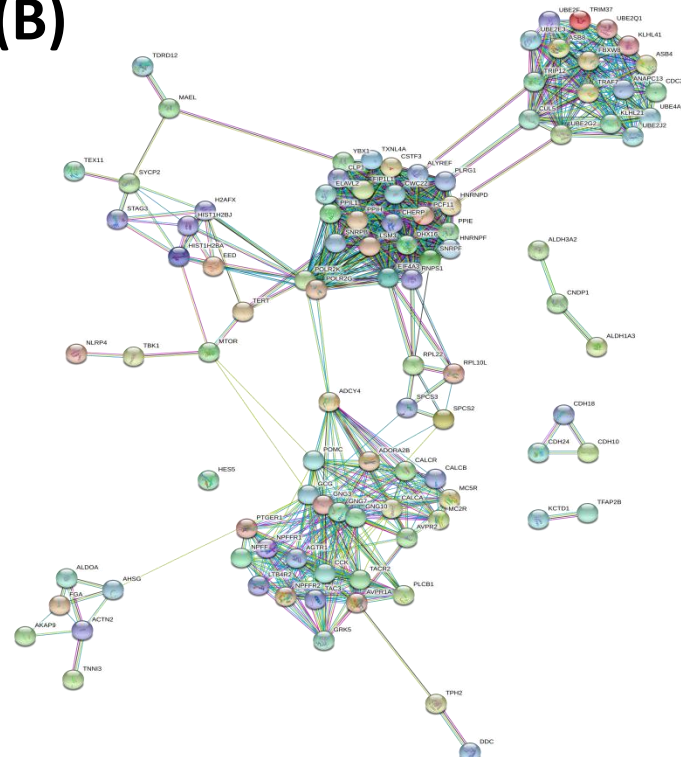
Number of edges:118

Average node degree:2.36

Avg. local clustering coefficient:0.461

Expected number of edges:62

PPI enrichment p-value:1.45e-10

(B)

nQTL: gene-network top-100 genes

Number of nodes:100

Number of edges:687

Average node degree:13.7

Avg. local clustering coefficient:0.829

Expected number of edges:218

PPI enrichment p-value:< 1.0e-16