

**Single-Cell RNASeq
GSE57872**

Single-Cell
(TPM)

Partition by tumor

MGH26
(TPM)

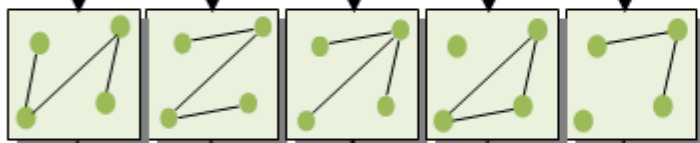
MGH28
(TPM)

MGH29
(TPM)

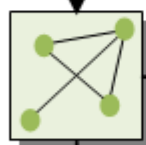
MGH30
(TPM)

MGH31
(TPM)

Construct co-expression networks



Construct summary network



equal densities

Module mining

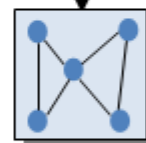


**Population RNASeq
GSE48865**

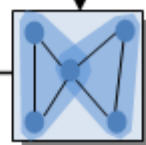
Pre-processing

Population
(TPM)

Construct co-expression network



Module mining



Compare