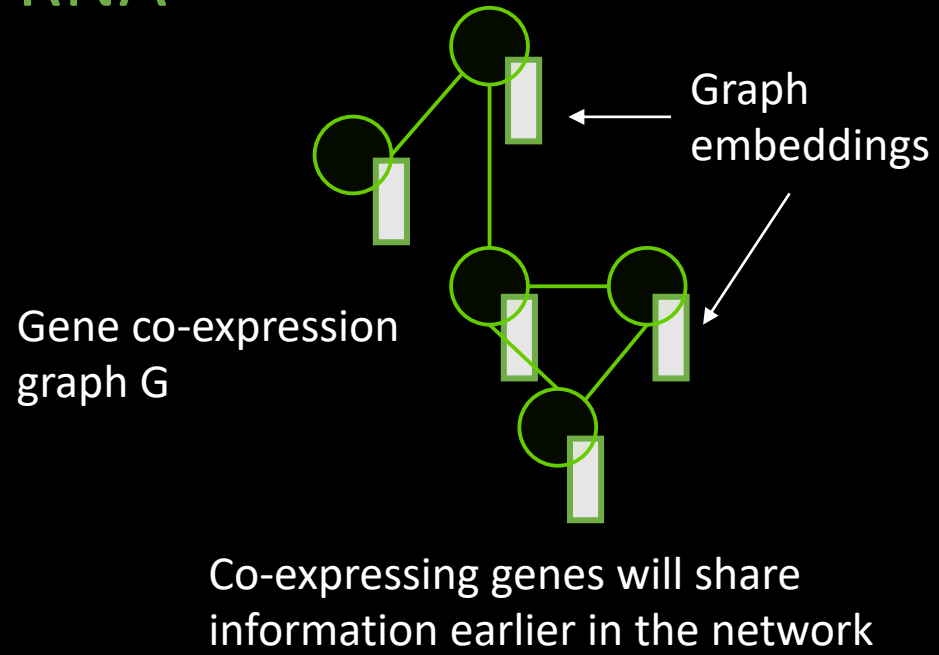
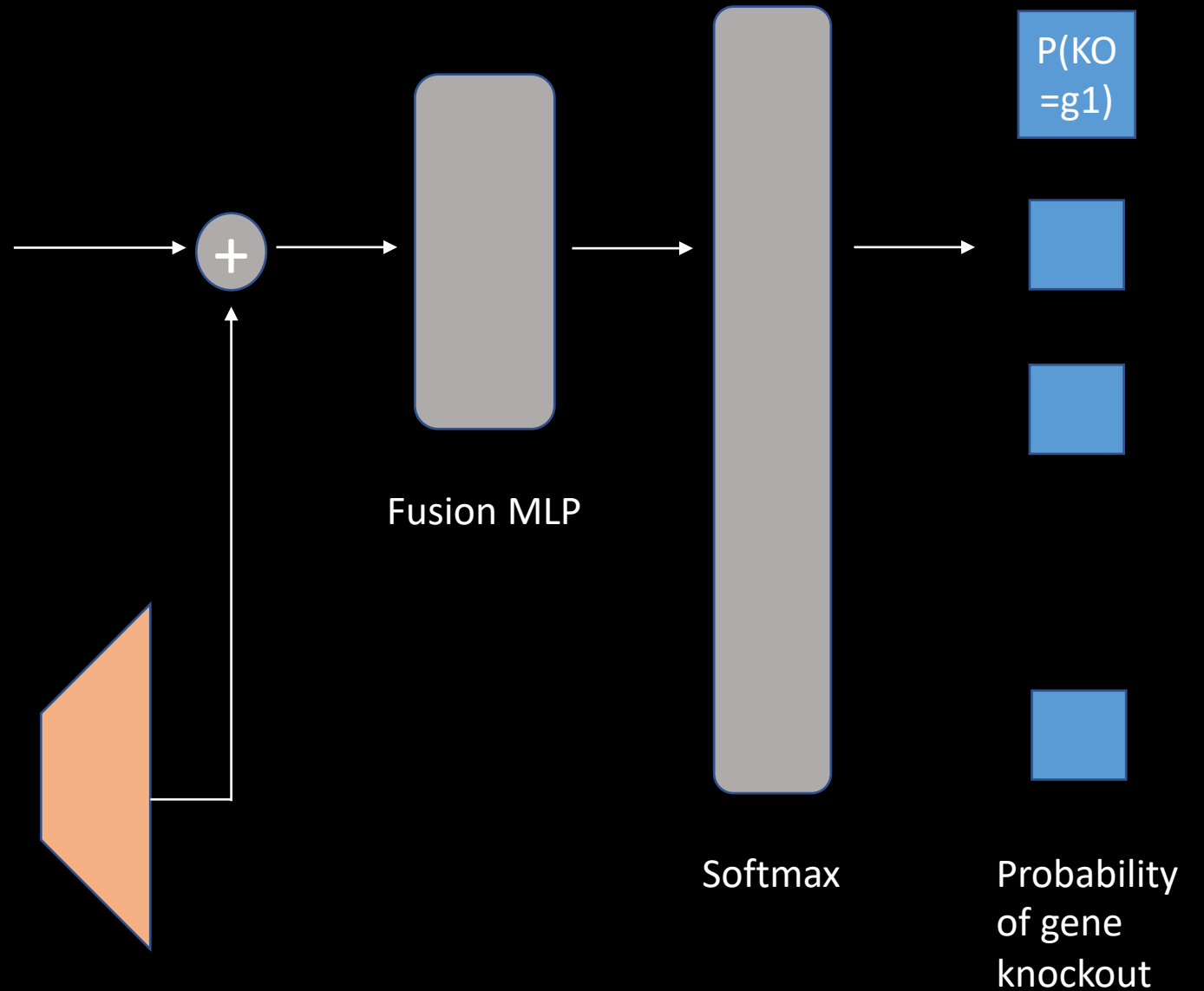


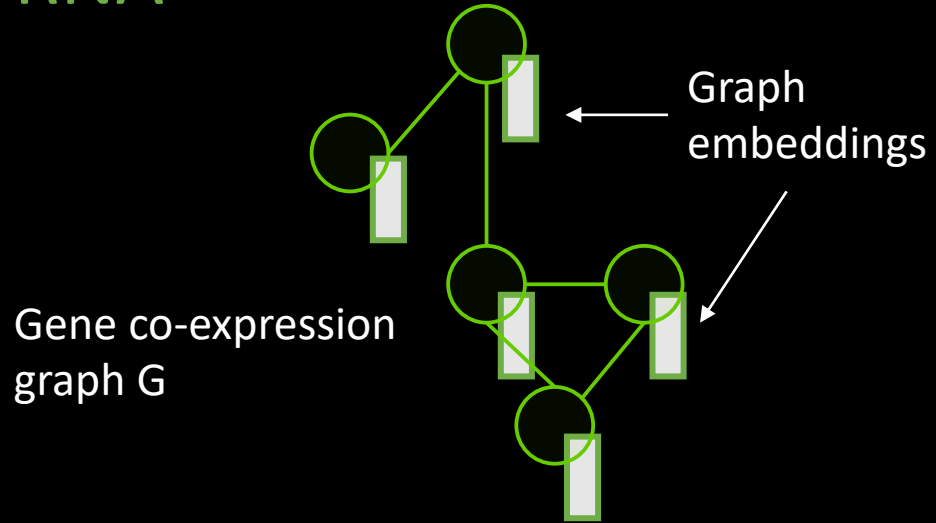
RNA



DNA



RNA



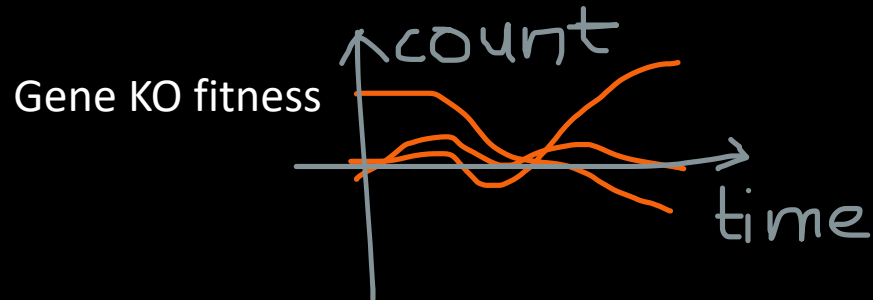
Co-expressing genes will share information earlier in the network

The graph G should be the WGCN

What if the features of the graph were the gene KO fitness data?

How do we incorporate multiple conditions? Should we have a distinct graph per experimental condition.

DNA



This is the winner of the two