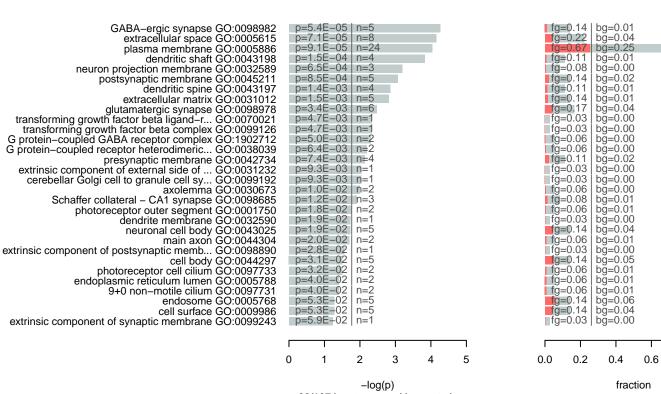


GO:CC gc026 fraction genes in fg and expected value

0.8

1.0



n=36/107 input genes with annotations