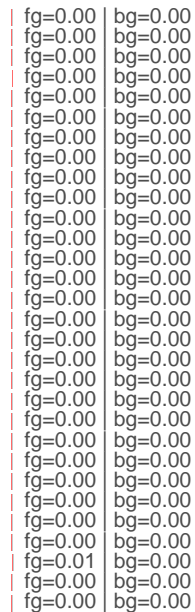


fraction genes in fg and bg



fraction

n=682/677 input genes with annotations