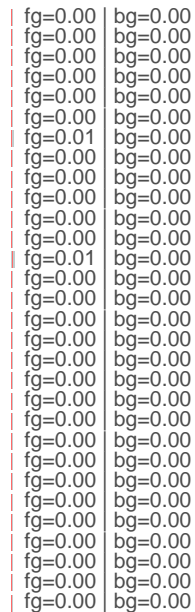


fraction genes in fg and bg



fraction

n=284/247 input genes with annotations