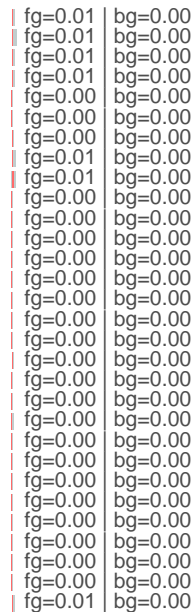


fraction genes in fg and bg



fraction

n=319/314 input genes with annotations