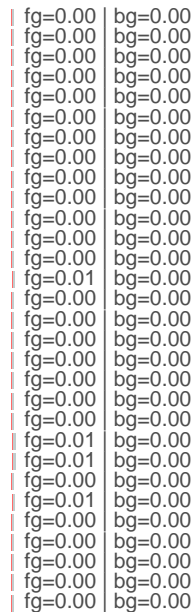


fraction genes in fg and bg



fraction

n=216/190 input genes with annotations