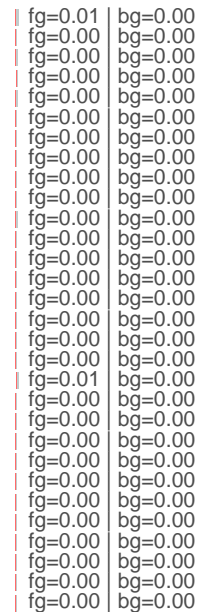


fraction genes in fg and bg



fraction

n=343/271 input genes with annotations