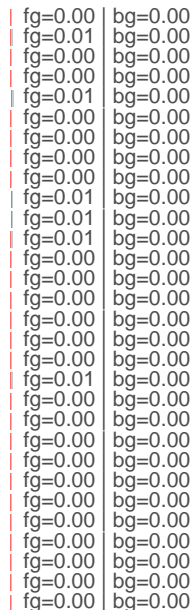


fraction genes in fg and bg



fraction

n=305/243 input genes with annotations