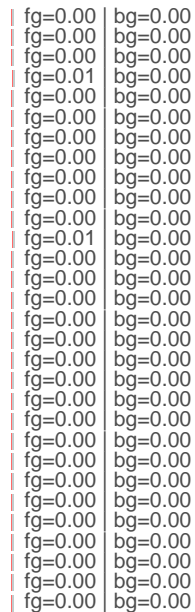


fraction genes in fg and bg



fraction

n=206/181 input genes with annotations