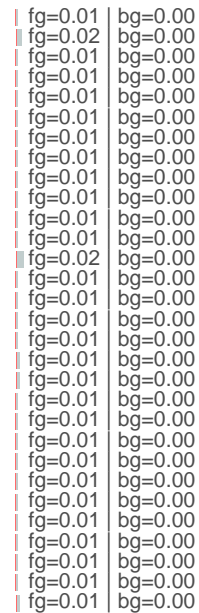


fraction genes in fg and bg



fraction

n=161/117 input genes with annotations