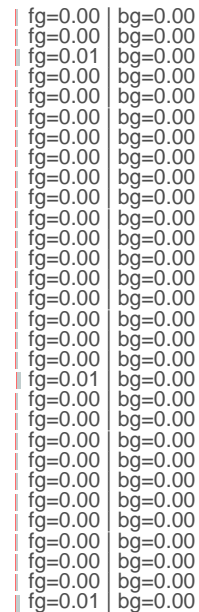


fraction genes in fg and bg



fraction

n=184/144 input genes with annotations