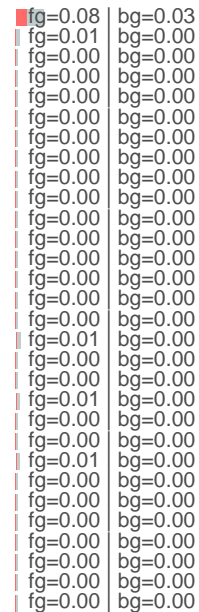


fraction genes in fg and bg



fraction

n=184/178 input genes with annotations