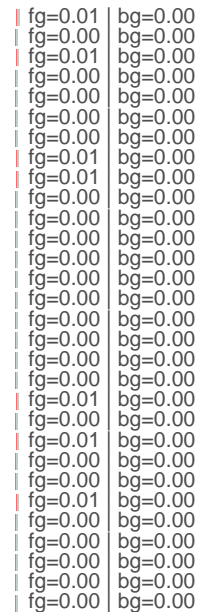


fraction genes in fg and bg



fraction

n=221/186 input genes with annotations