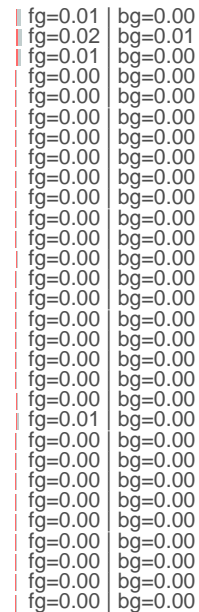


fraction genes in fg and bg



fraction

n=567/591 input genes with annotations