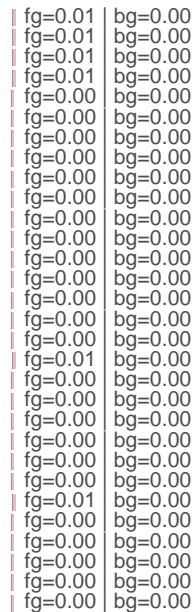


fraction genes in fg and bg



fraction

n=167/158 input genes with annotations