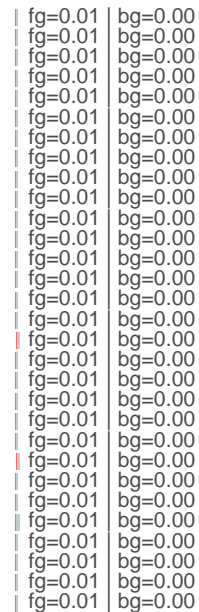


fraction genes in fg and bg



fraction

n=169/139 input genes with annotations