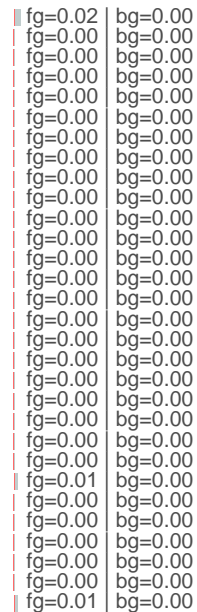


fraction genes in fg and bg



fraction

n=263/242 input genes with annotations