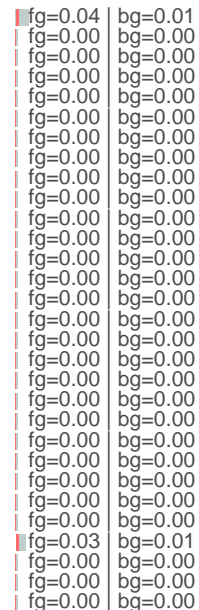


fraction genes in fg and bg



fraction

n=173/149 input genes with annotations