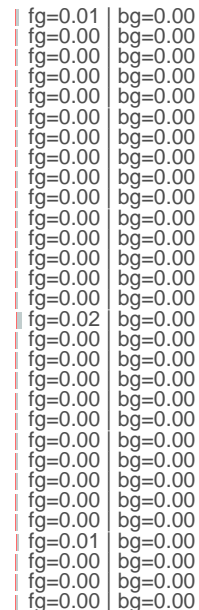


fraction genes in fg and bg



fraction

n=209/174 input genes with annotations