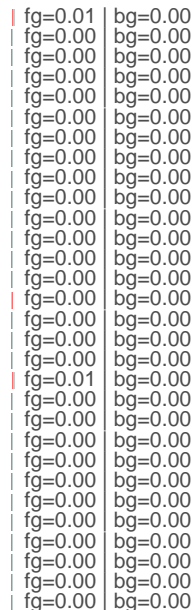


fraction genes in fg and bg



fraction

n=506/484 input genes with annotations