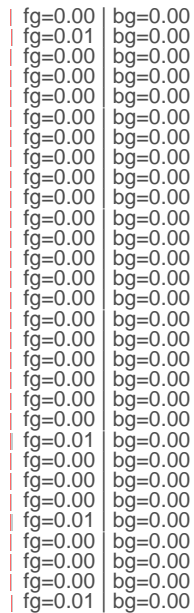


fraction genes in fg and bg



fraction

n=307/283 input genes with annotations