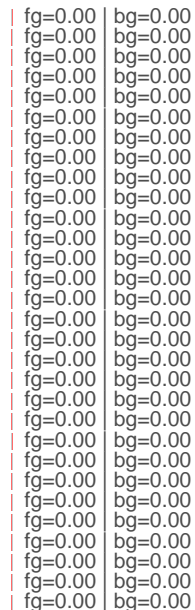


fraction genes in fg and bg



fraction

n=484/436 input genes with annotations