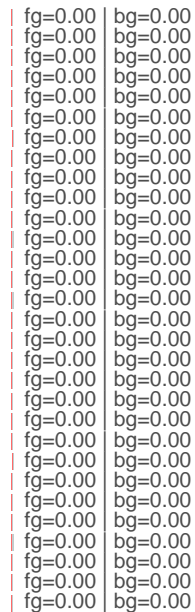


fraction genes in fg and bg



fraction

n=430/385 input genes with annotations