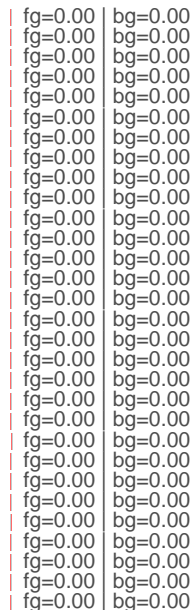


fraction genes in fg and bg



fraction

n=459/437 input genes with annotations