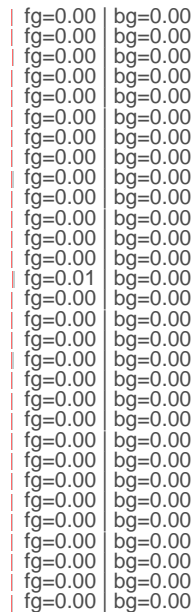


fraction genes in fg and bg



fraction

n=380/338 input genes with annotations