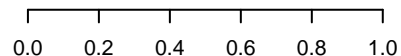
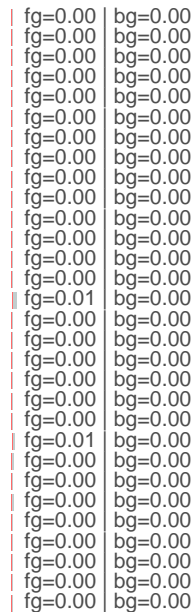


fraction genes in fg and bg



fraction

n=404/364 input genes with annotations