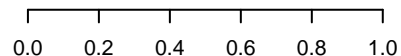
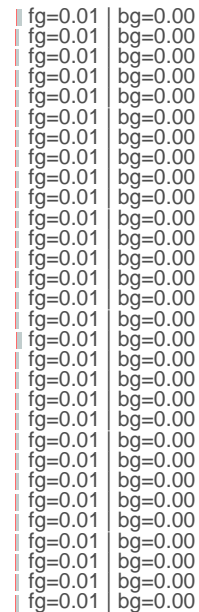


fraction genes in fg and bg



fraction

n=123/97 input genes with annotations