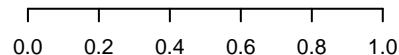
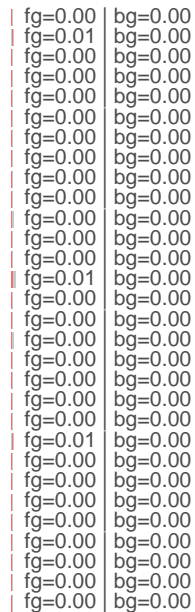


fraction genes in fg and bg



fraction

n=406/365 input genes with annotations