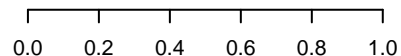
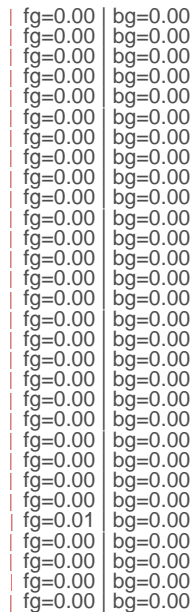


fraction genes in fg and bg



fraction

n=546/520 input genes with annotations