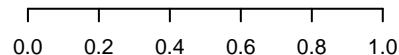
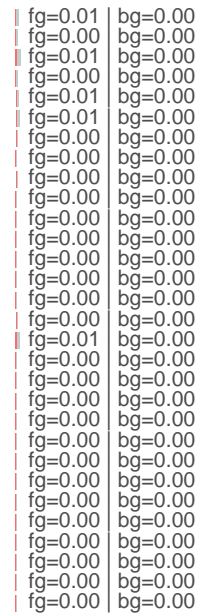


fraction genes in fg and bg



fraction

n=425/436 input genes with annotations