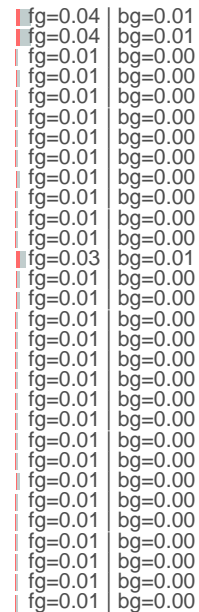


fraction genes in fg and bg



fraction

n=137/135 input genes with annotations