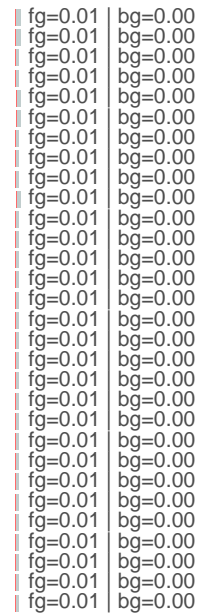


fraction genes in fg and bg



fraction

n=130/113 input genes with annotations