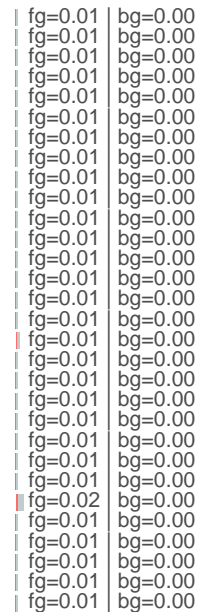


fraction genes in fg and bg



fraction

n=168/139 input genes with annotations