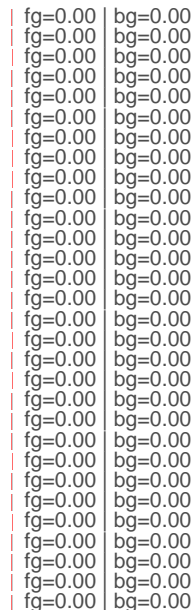


fraction genes in fg and bg



fraction

n=832/866 input genes with annotations