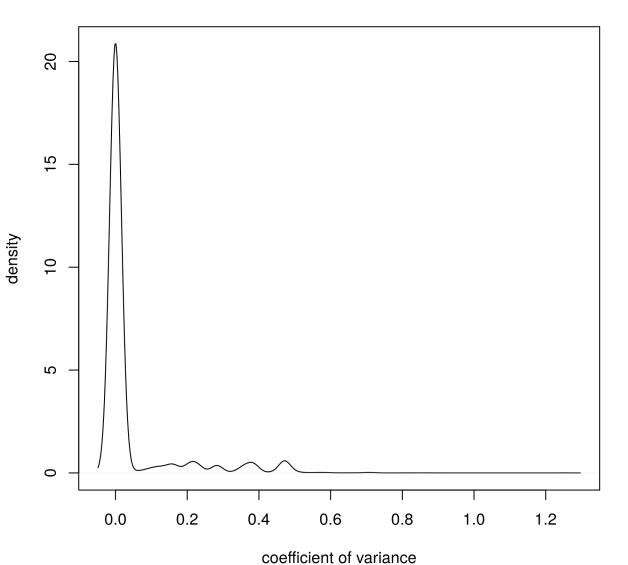
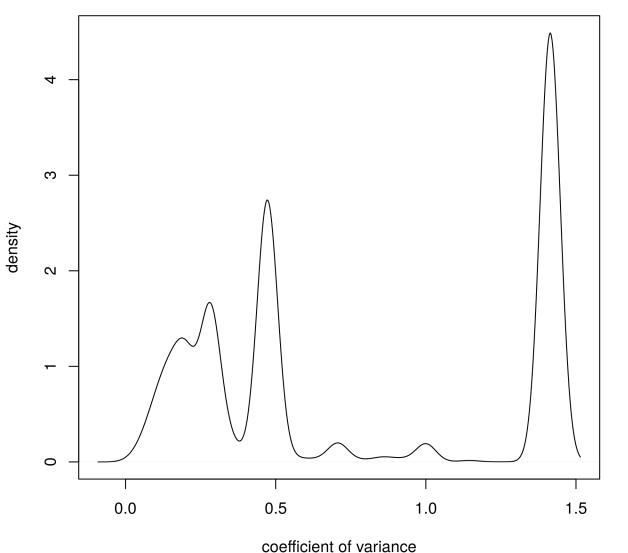
CV of counts of identical token sets (all proteins)



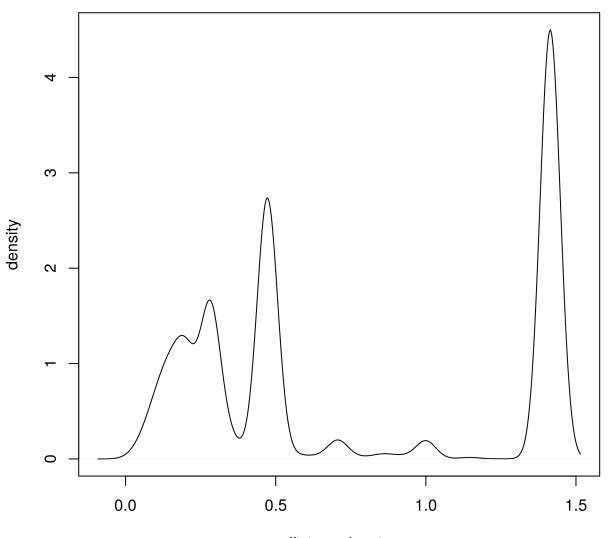
mean: 0.041, CV = sd / (mean + 1)

CV of counts of identical token sets (neighbouring tokens in same protein)



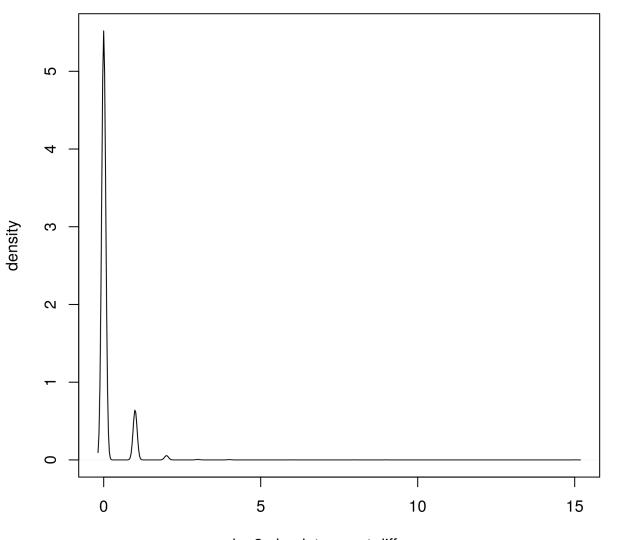
mean: 0.757

CV of counts of identical token sets



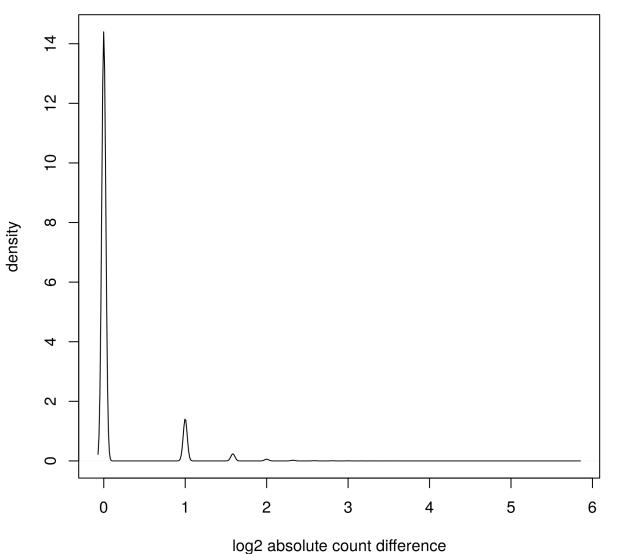
coefficient of variance mean: 0.758

count differences between identical token sets (all proteins)



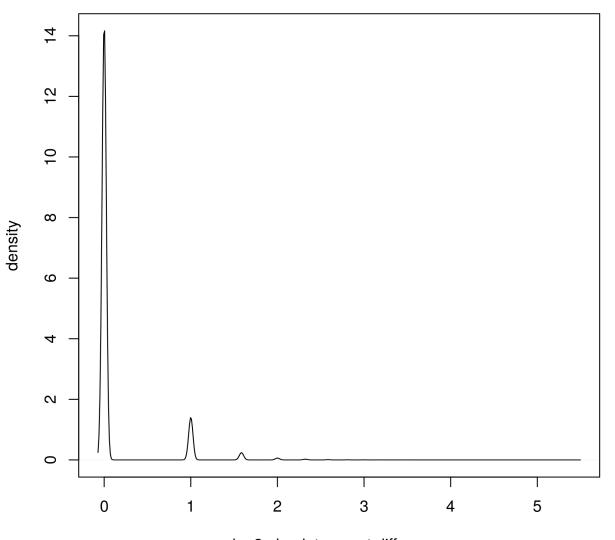
log2 absolute count difference mean: 0.129 (4.503 counts)

count differences between identical token sets (neighbouring tokens in same protein)



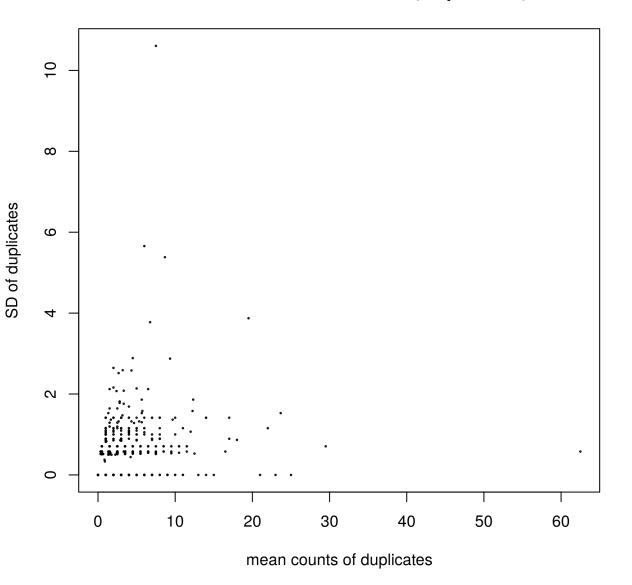
mean: 0.124 (1.141 counts)

count differences between identical token sets

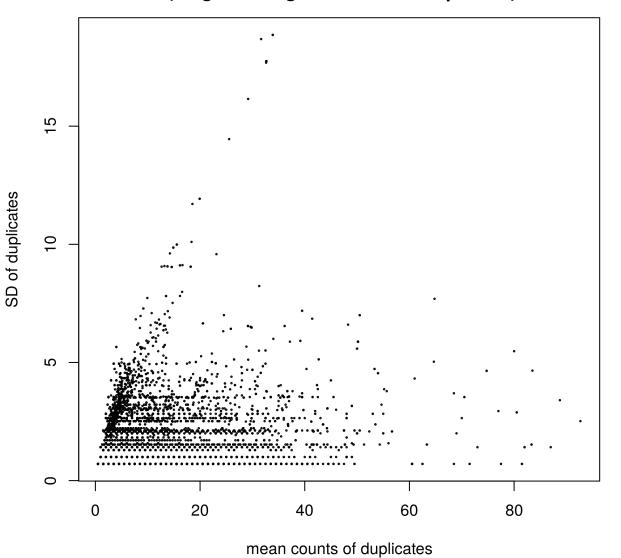


log2 absolute count difference mean: 0.125 (1.143 counts)

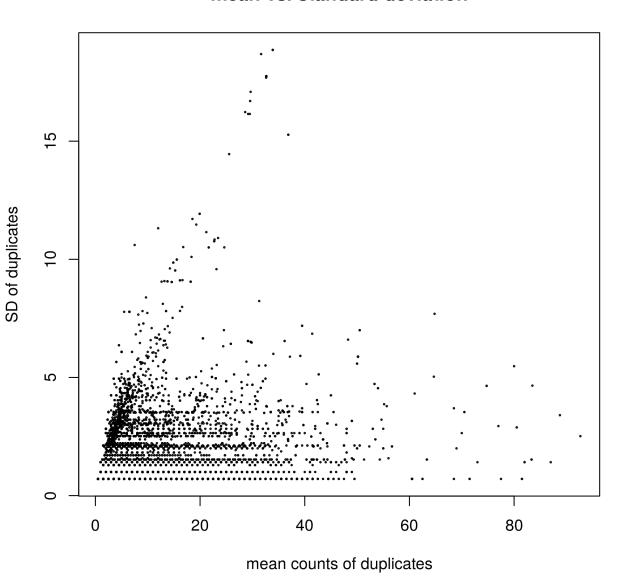
mean vs. standard deviation (all proteins)



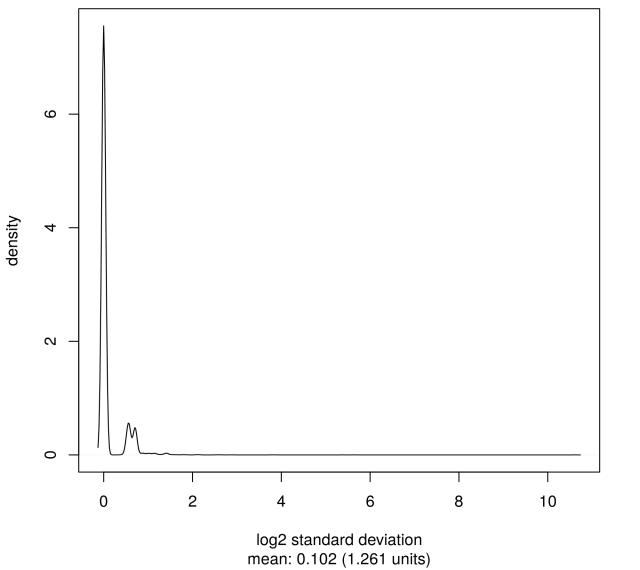
mean vs. standard deviation (neighbouring tokens in same protein)



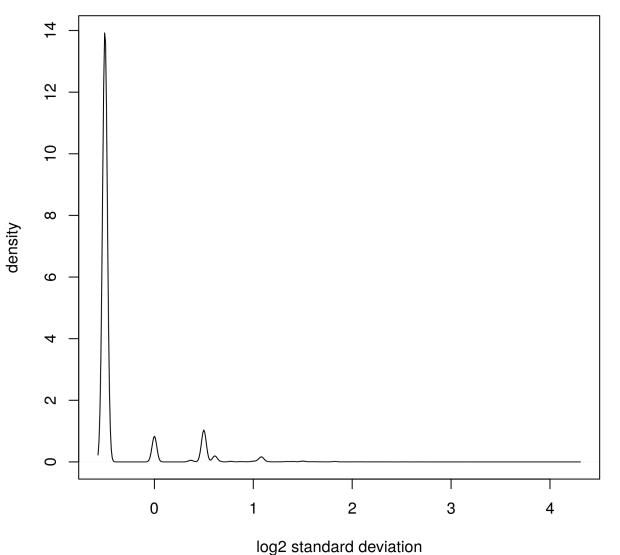
mean vs. standard deviation



standard deviation of counts of identical token sets (all proteins)

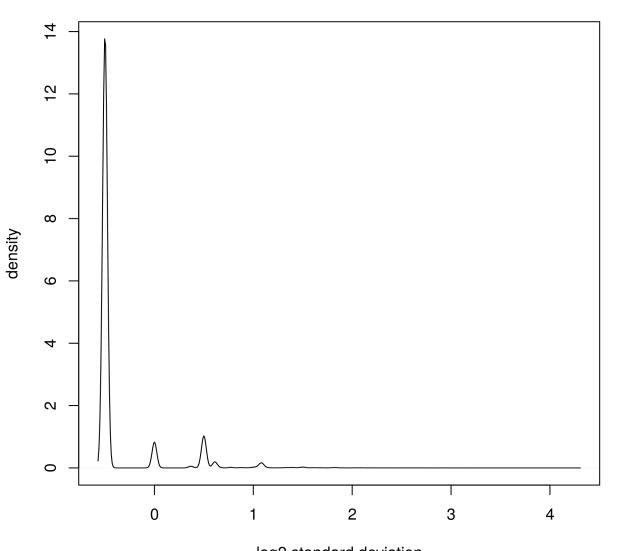


standard deviation of counts of identical token sets (neighbouring tokens in same protein)



mean: -0.359 (0.815 units)

standard deviation of counts of identical token sets



log2 standard deviation mean: -0.358 (0.816 units)