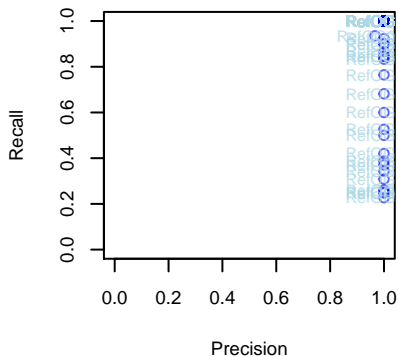
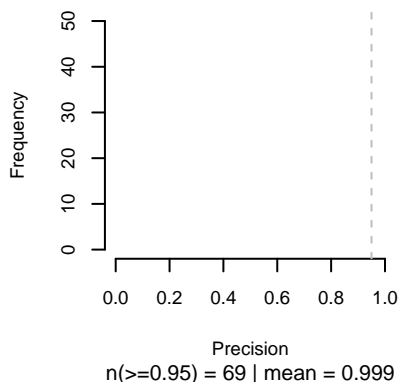


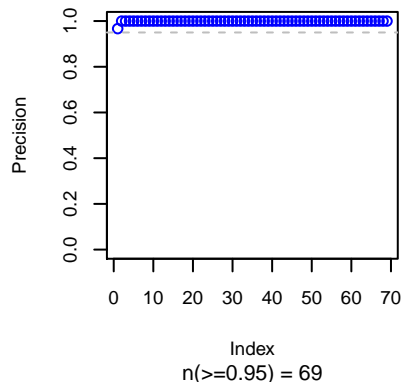
Precision & recall



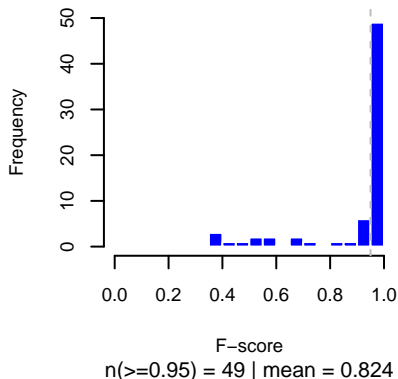
Precision



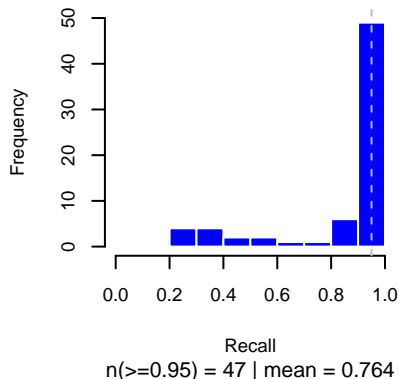
Precision



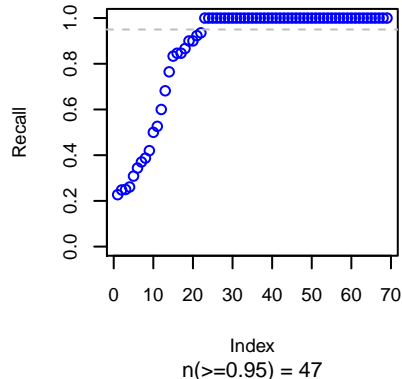
F-score



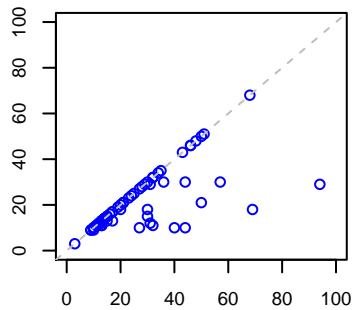
Recall



Recall

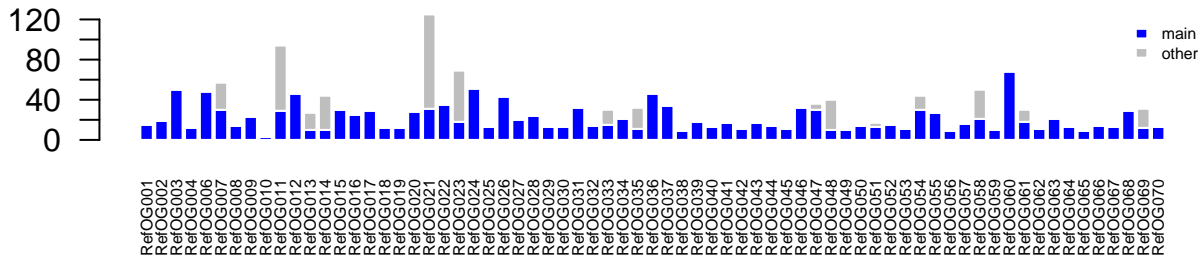


genes in best OG



n(best < all) = 51 (p=0.999 | r=0.993)
n(best = all) = 47 (p=1.000 | r=1.000)
n(best > all) = 22 (p=0.998 | r=0.588)

number of reference genes in main and other OGs



num OGs per refOG

