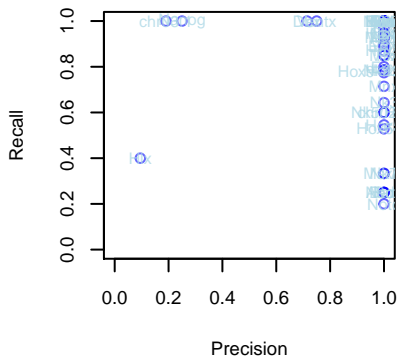
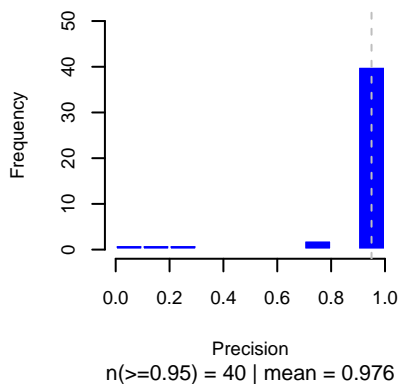


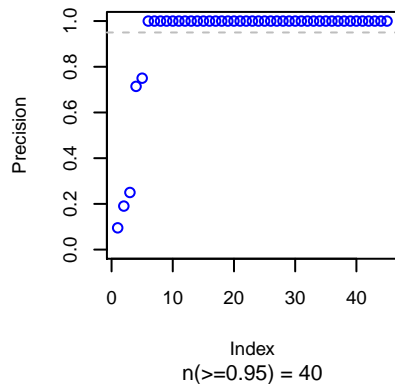
Precision & recall



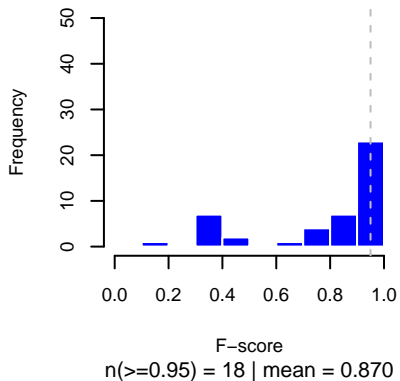
Precision



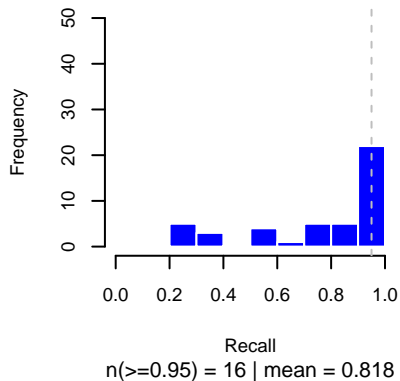
Precision



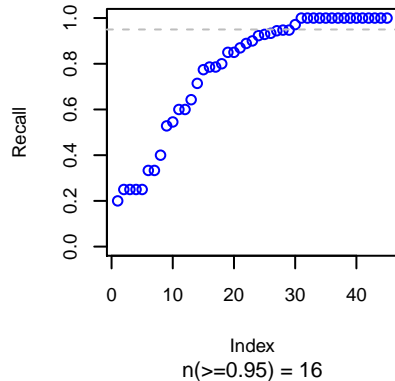
F-score



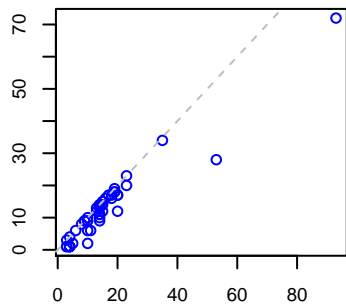
Recall



Recall

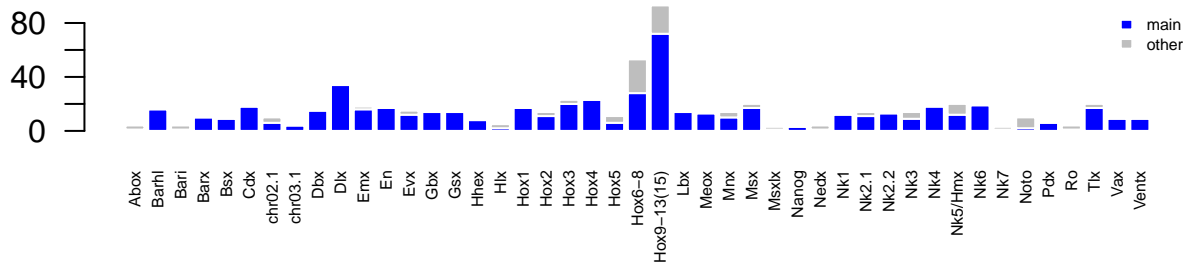


genes in best OG



n(best = all) = 23 (p=0.900 | r=0.978)
n(best = all) = 15 (p=0.860 | r=1.000)
n(best < all) = 30 (p=0.970 | r=0.667)

number of reference genes in main and other OGs



num OGs per refOG

