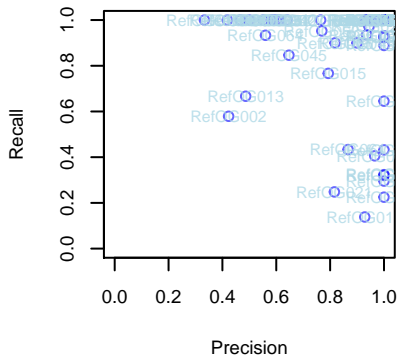
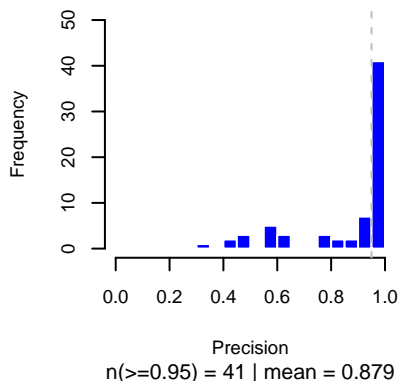


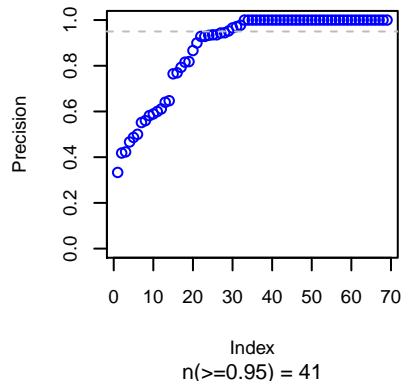
Precision & recall



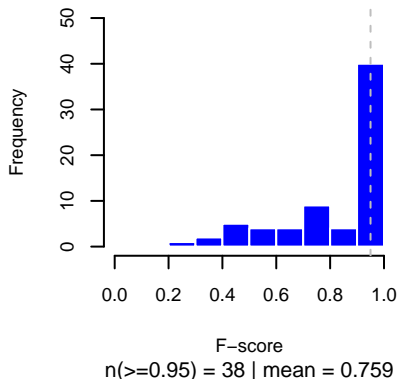
Precision



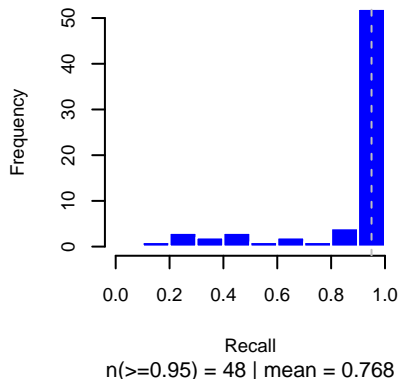
Precision



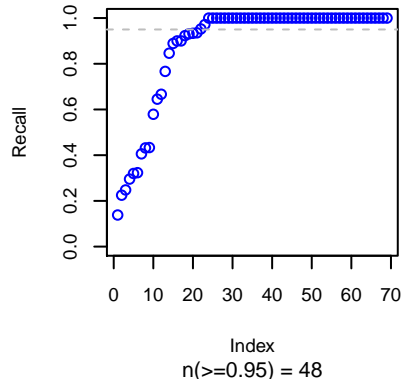
F-score



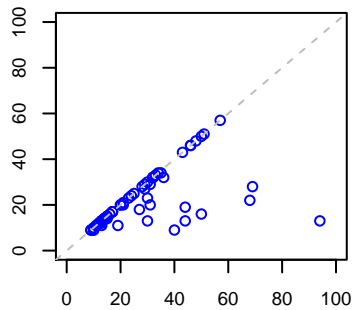
Recall



Recall

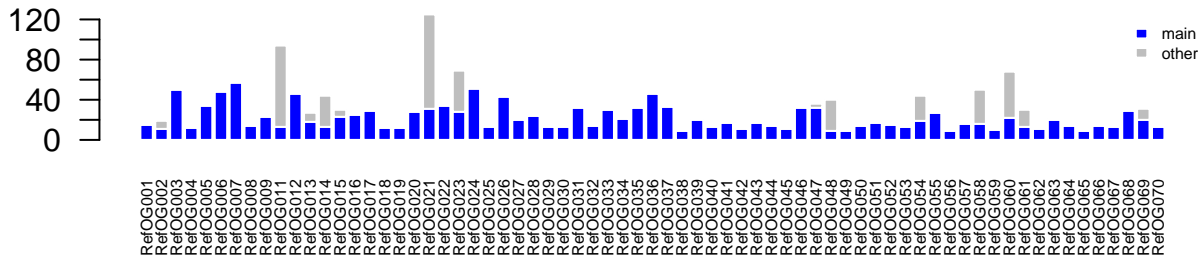


genes in best OG



n(best = all) = 50 (p=0.990 | r=0.990)
n(best = all) = 46 (p=0.885 | r=1.000)
n(best < all) = 23 (p=0.863 | r=0.637)

number of reference genes in main and other OGs



num OGs per refOG

