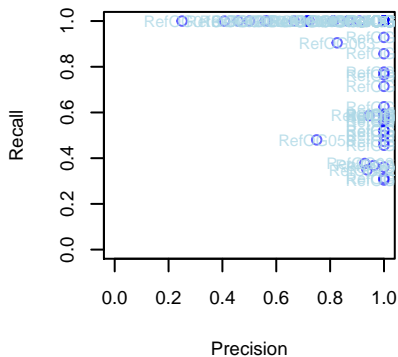
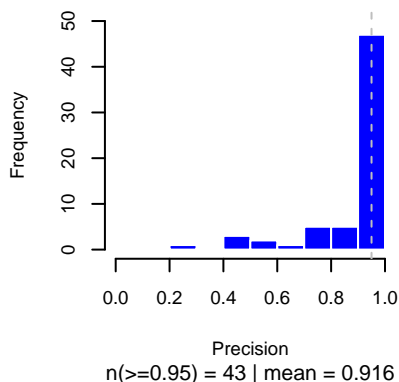


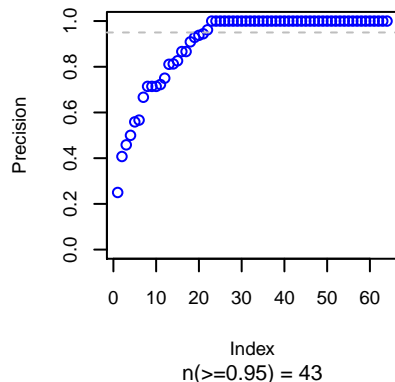
Precision & recall



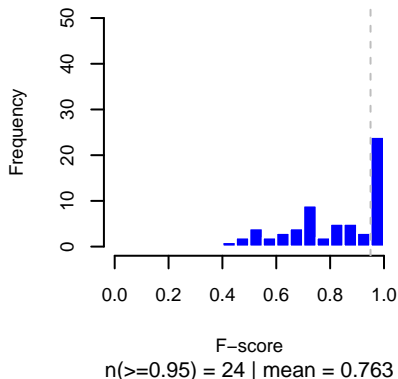
Precision



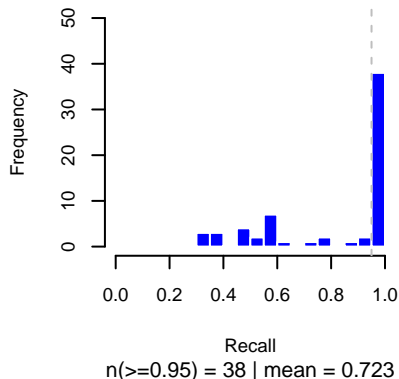
Precision



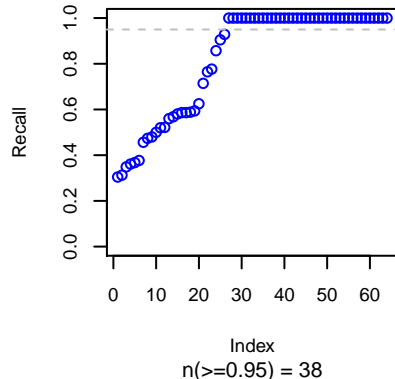
F-score



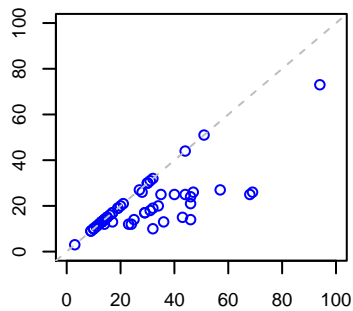
Recall



Recall

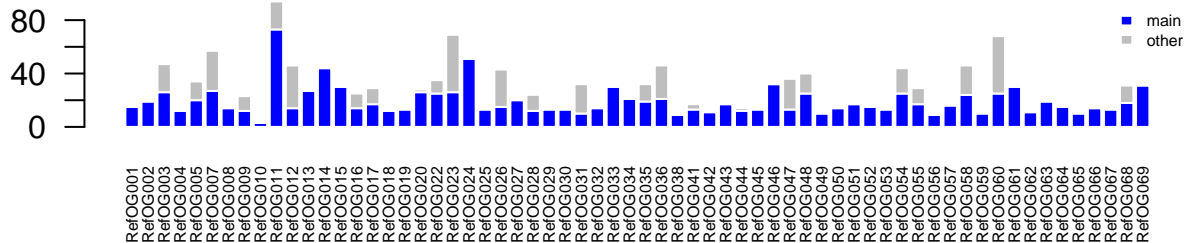


genes in best OG



n(best = all) = 39 (p=0.856 | r=0.998)
n(best < all) = 25 (p=0.981 | r=0.550)

number of reference genes in main and other OGs



num OGs per refOG

