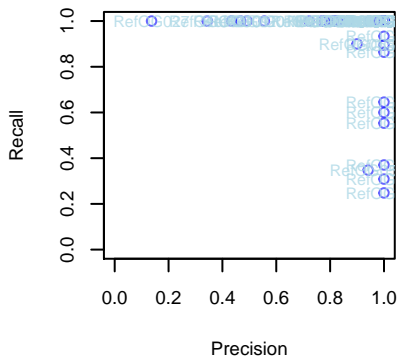
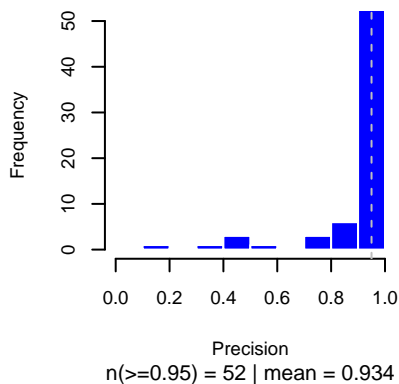


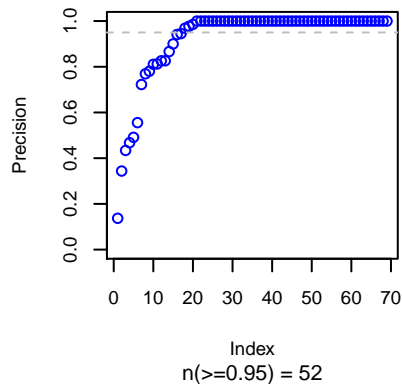
Precision & recall



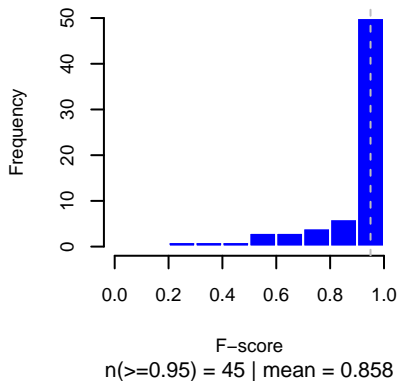
Precision



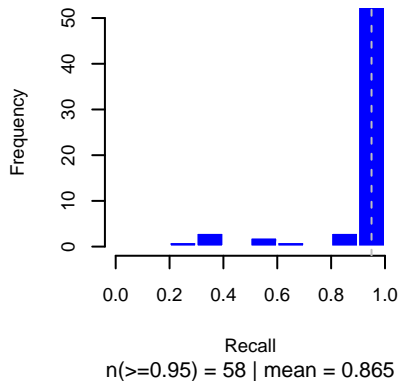
Precision



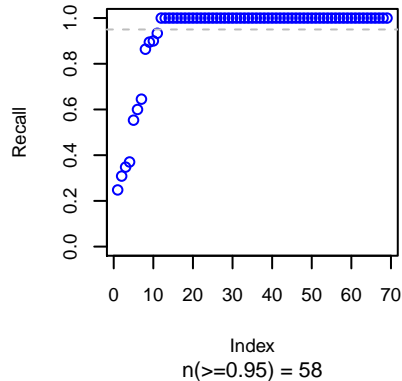
F-score



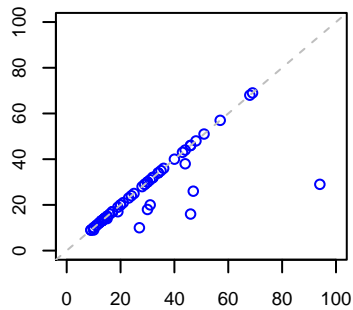
Recall



Recall

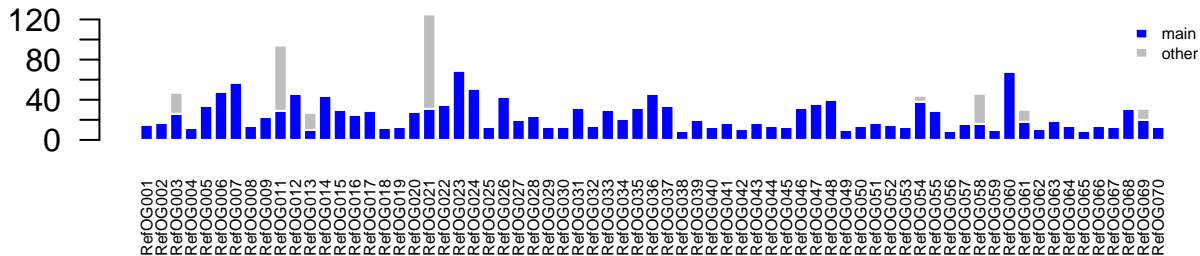


genes in best OG



n(best = all) = 66 (p=0.997 | r=0.997)
n(best = all) = 58 (p=0.909 | r=1.000)
n(best < all) = 11 (p=0.986 | r=0.606)

number of reference genes in main and other OGs



num OGs per refOG

