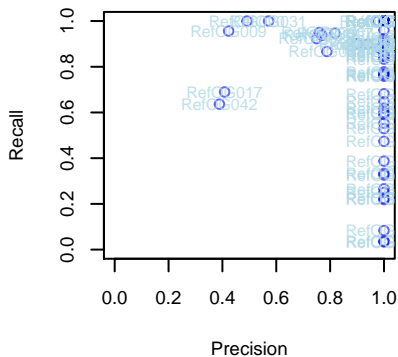
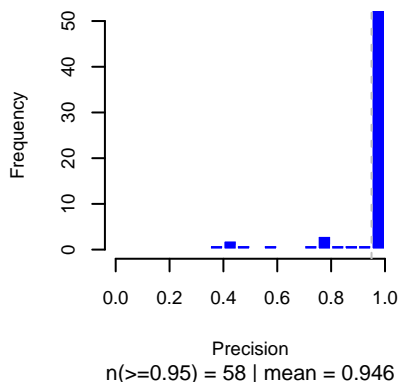


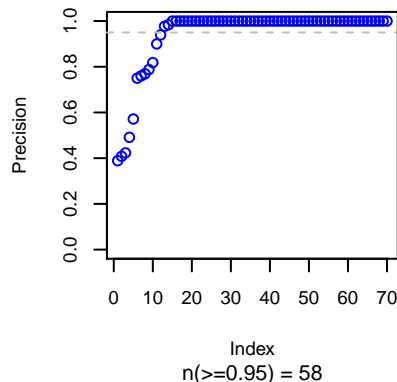
Precision & recall



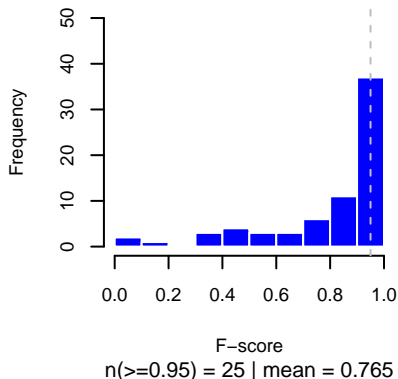
Precision



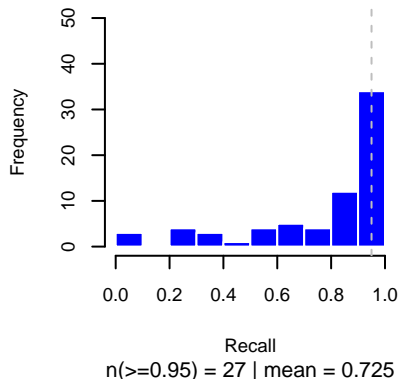
Precision



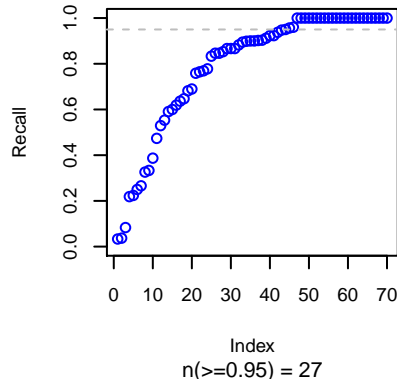
F-score



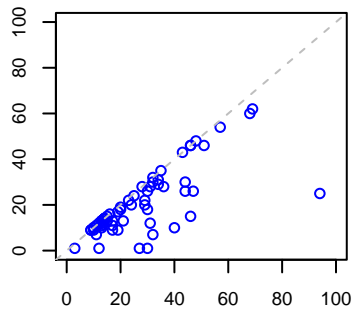
Recall



Recall

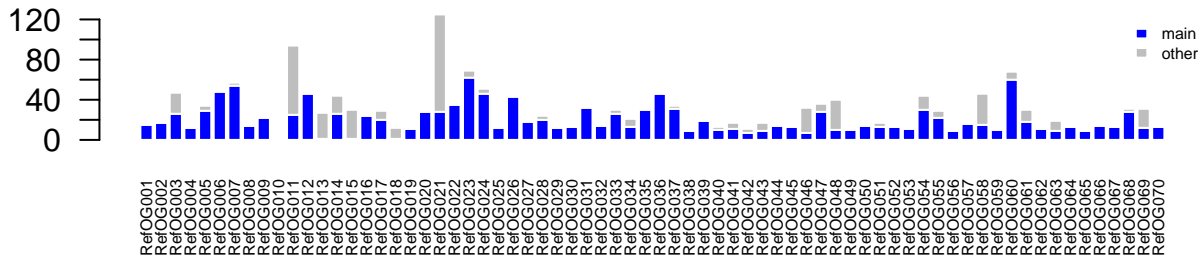


genes in best OG



n(best = all) = 26 (p=0.975)
n(best = all) = 24 (p=0.960 | r=1.000)
n(best < all) = 46 (p=0.933 | r=0.674)

number of reference genes in main and other OGs



num OGs per refOG

