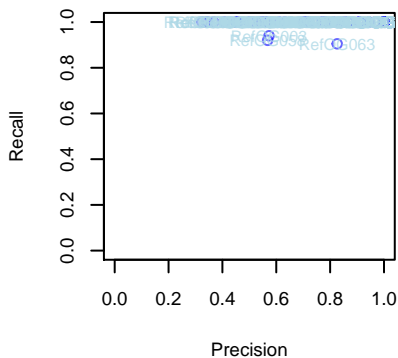
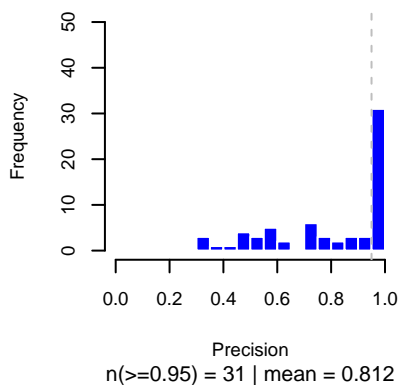


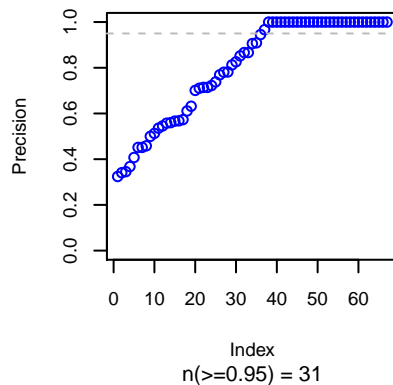
Precision & recall



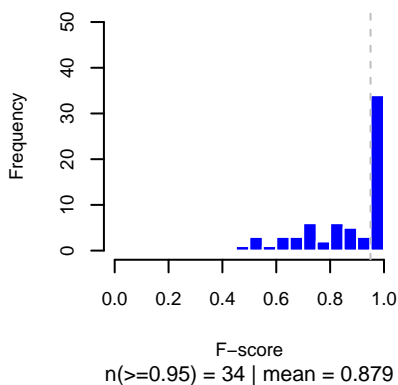
Precision



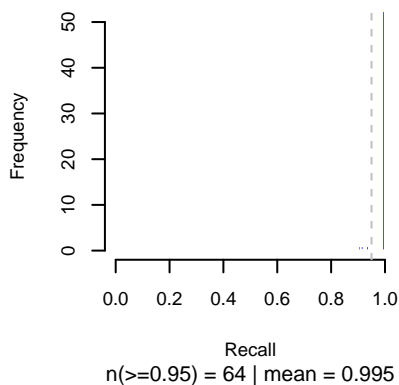
Precision



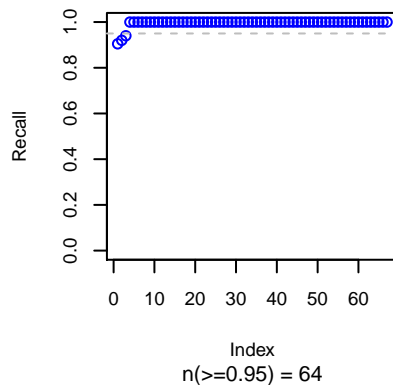
F-score



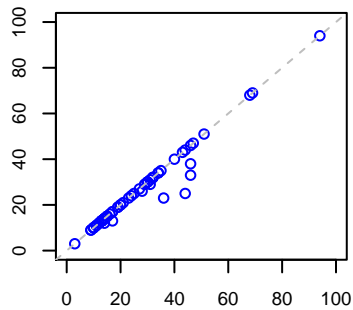
Recall



Recall

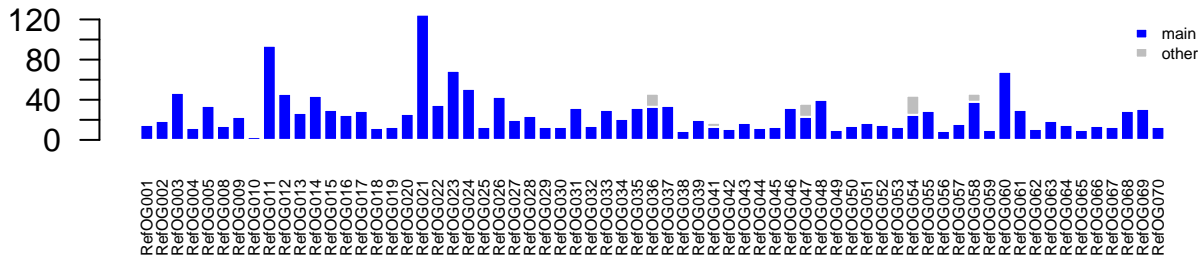


genes in best OG



n(best = all) = 61 (p=0.002 | r=0.997)
 n(best = all) = 59 (p=0.809 | r=0.997)
 n(best < all) = 8 (p=0.771 | r=0.990)

number of reference genes in main and other OGs



num OGs per refOG

