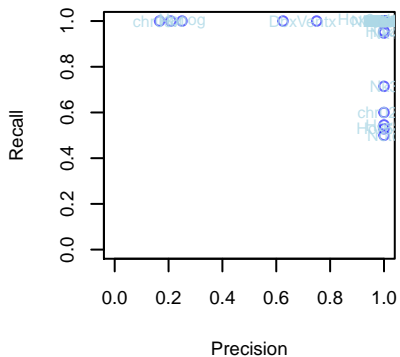
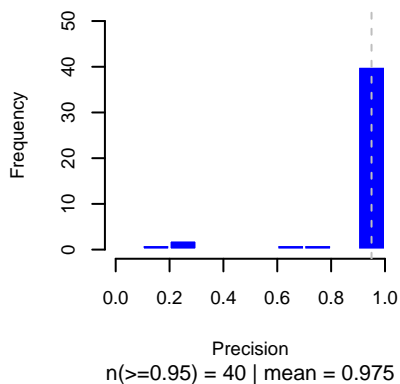


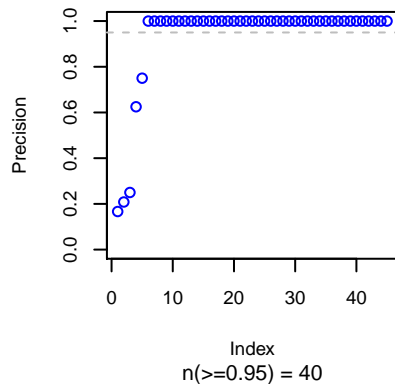
Precision & recall



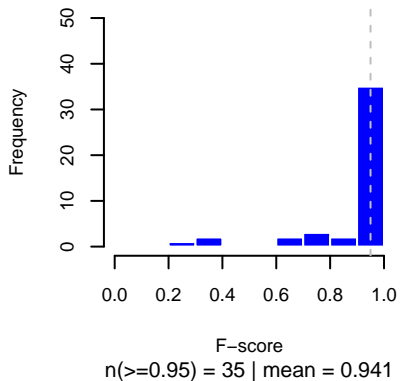
Precision



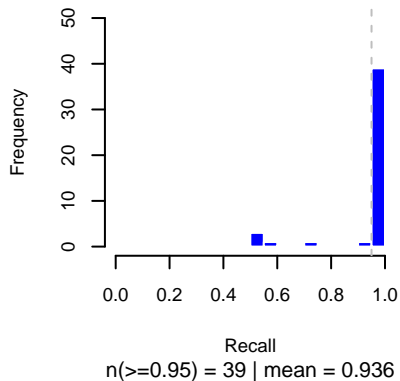
Precision



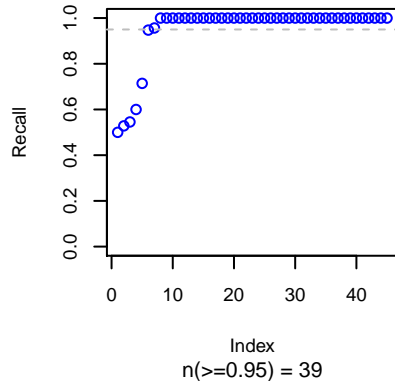
F-score

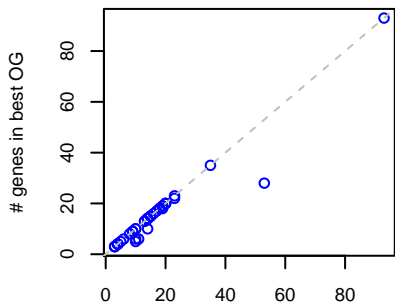


Recall



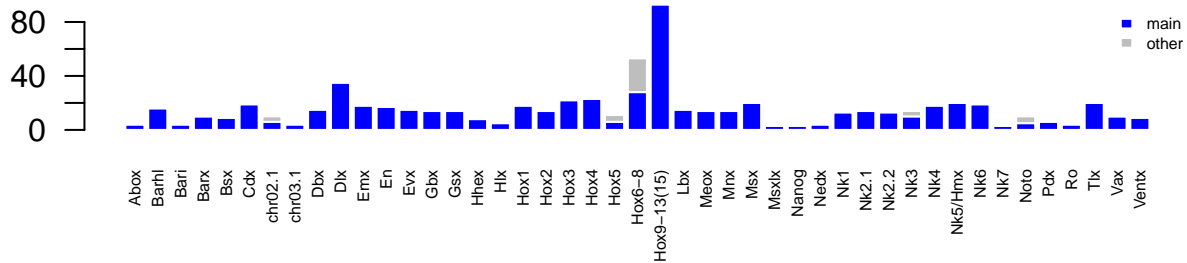
Recall





n(best = all) = 46 (p=0.925 | r=0.998)
 n(best = all) = 38 (p=0.921 | r=1.000)
 n(best < all) = 7 (p=1.000 | r=0.685)

number of reference genes in main and other OGs



num OGs per refOG

