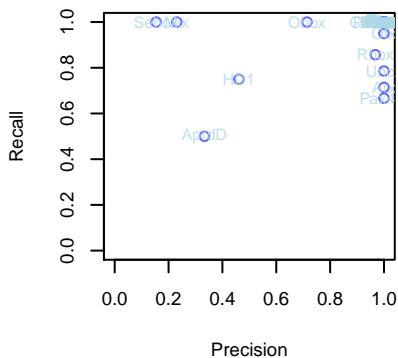
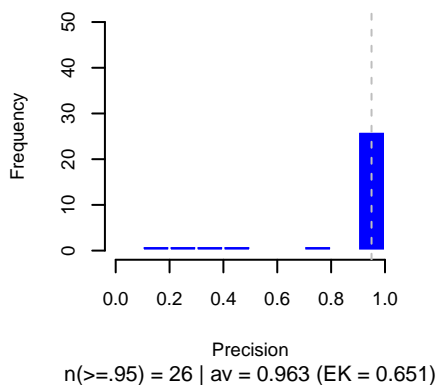
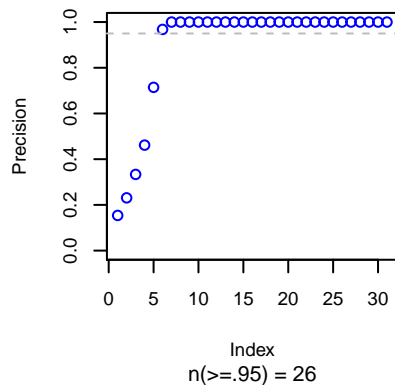
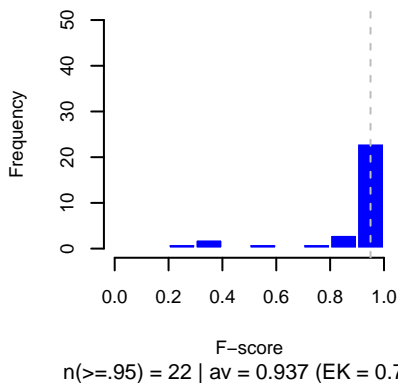
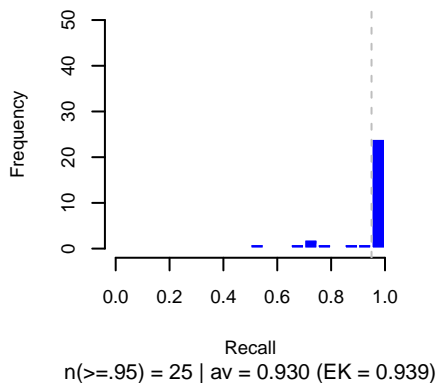
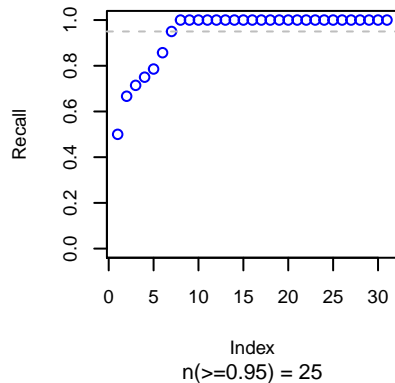
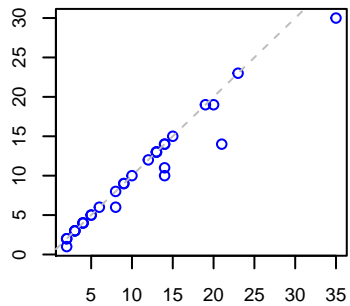


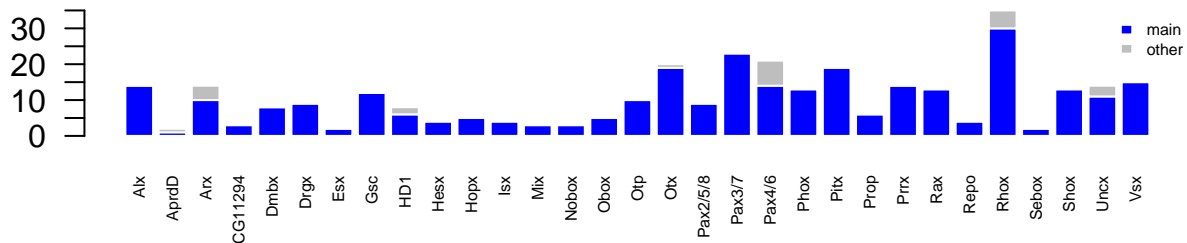
**Precision & recall****Precision****Precision****F-score****Recall****Recall**

# genes in best OG



n(best = 25) (p=0.921 | r=0.998)  
n(best = all) = 24 (p=0.921 | r=1.000)  
n(best < all) = 7 (p=0.823 | r=0.746)

## number of reference genes in main and other OGs



## num OGs per refOG

