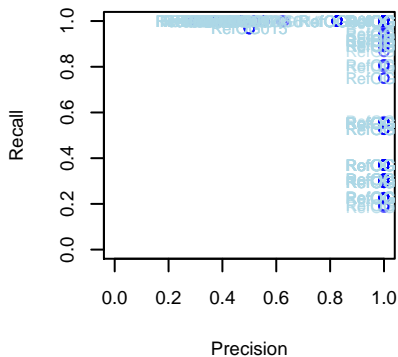
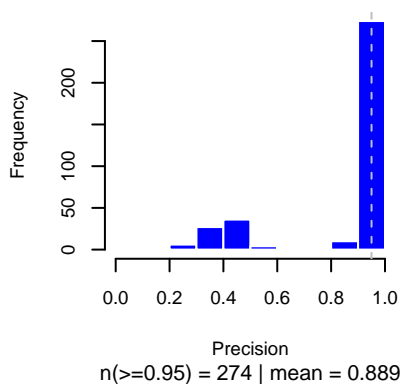


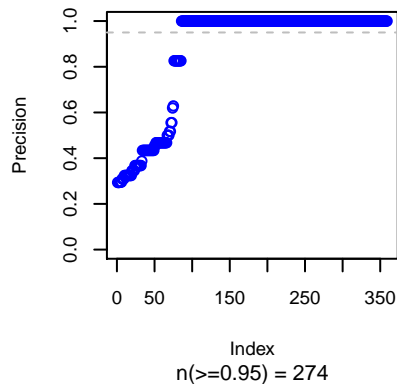
### Precision & recall



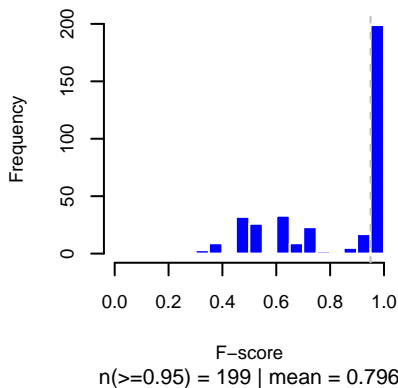
### Precision



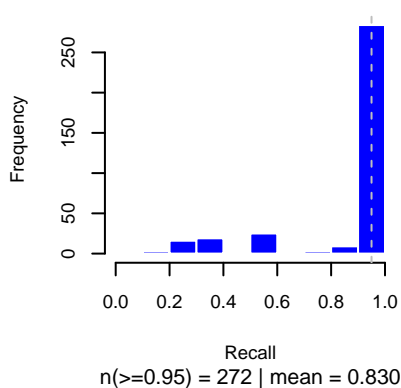
### Precision



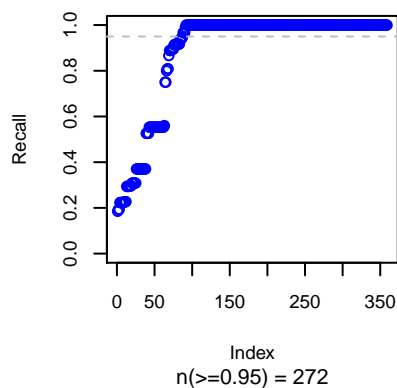
### F-score



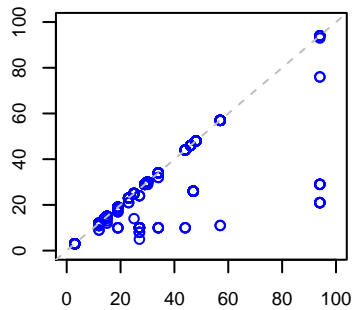
### Recall



### Recall

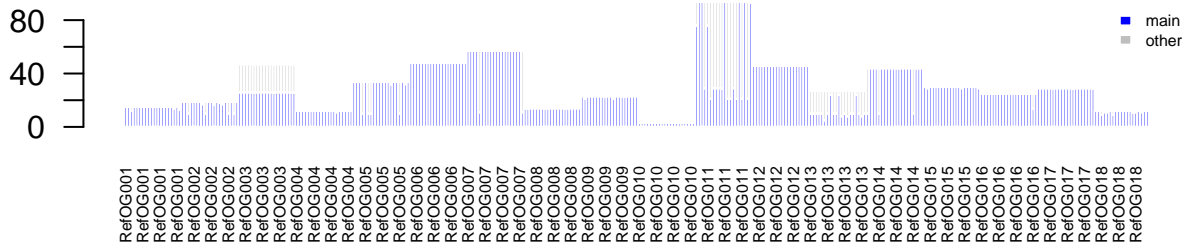


# genes in best OG



n(best > 0.9 all) = 294 (p=0.996)  
 n(best = all) = 268 (p=0.834 | r=1.000)  
 n(best < all) = 91 (p=0.984 | r=0.551)

## number of reference genes in main and other OGs



## num OGs per refOG

