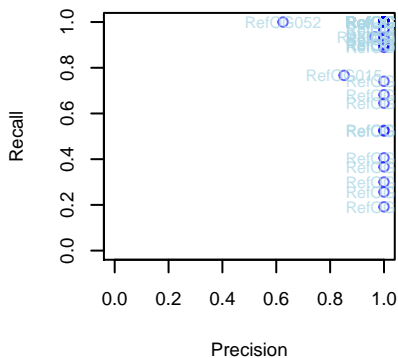
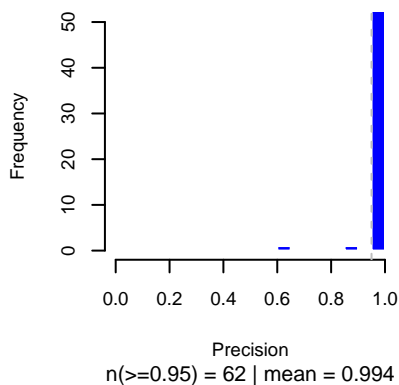


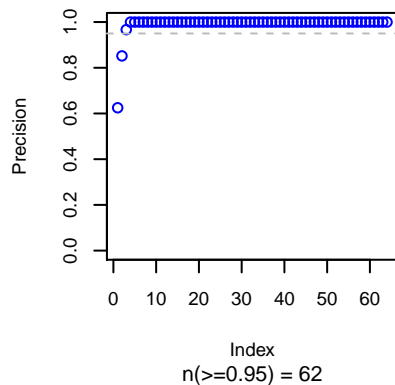
### Precision & recall



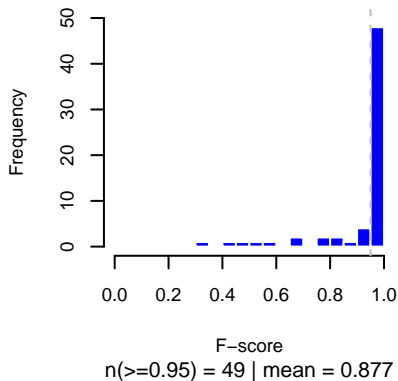
### Precision



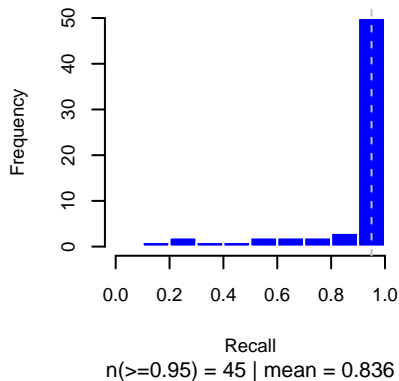
### Precision



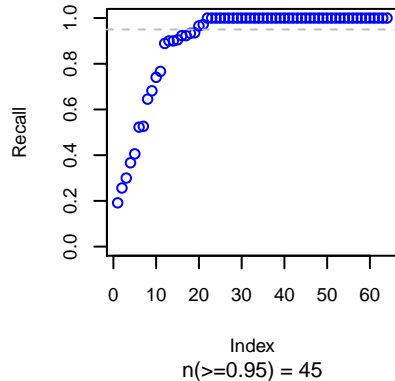
### F-score



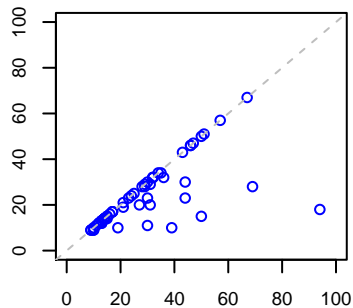
### Recall



### Recall

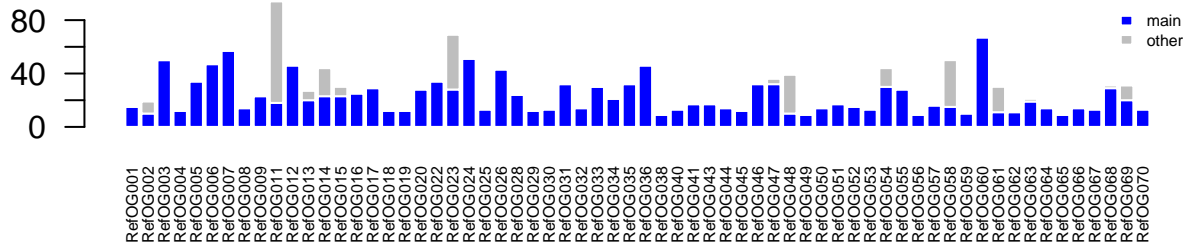


# genes in best OG



n(best = all) = 52 (p=0.992 | r=0.988)  
n(best = all) = 43 (p=0.991 | r=1.000)  
n(best < all) = 21 (p=0.991 | r=0.698)

## number of reference genes in main and other OGs



## num OGs per refOG

