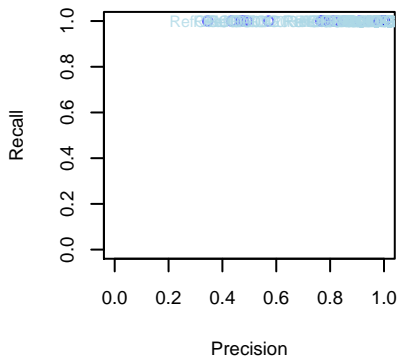
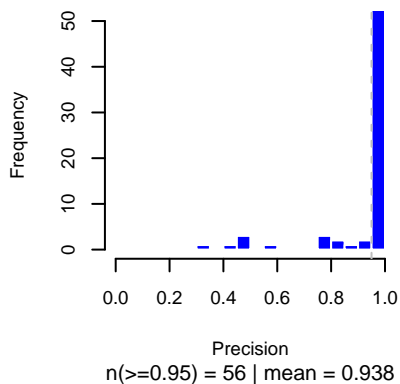


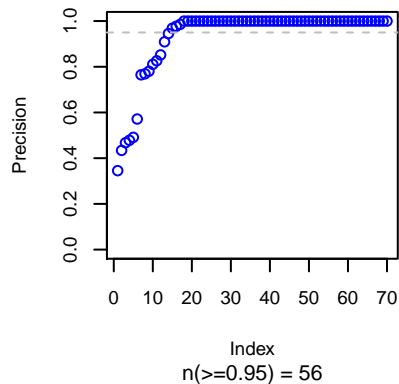
### Precision & recall



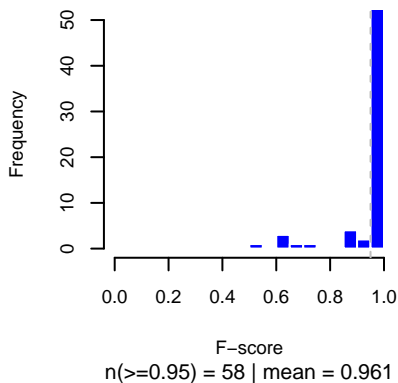
### Precision



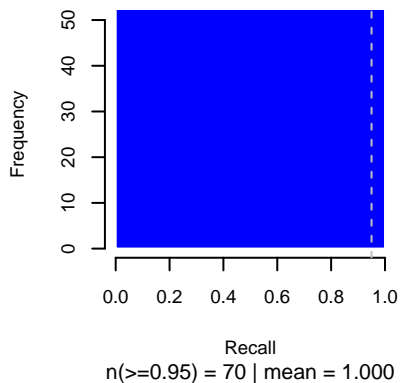
### Precision



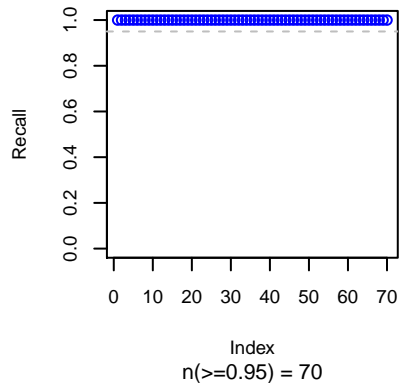
### F-score



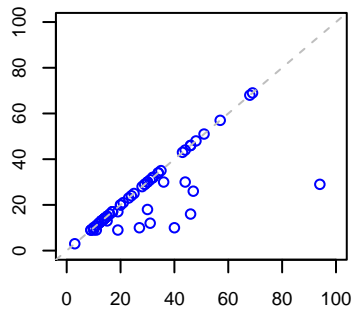
### Recall



### Recall

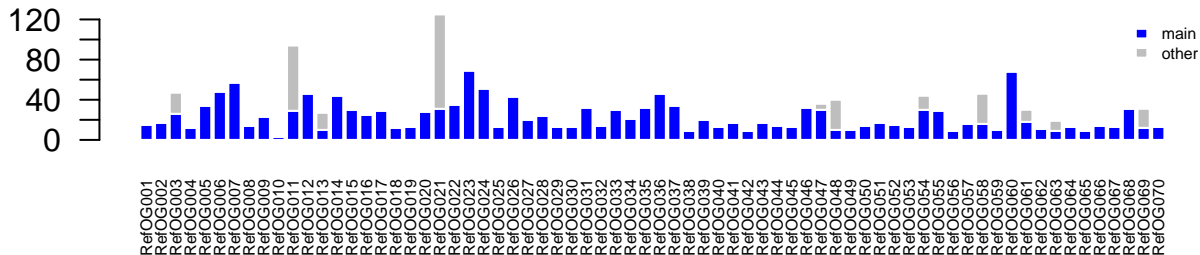


# genes in best OG



n(best = all) = 56 (p=1.000 | r=1.000)  
n(best = all) = 55 (p=0.942 | r=1.000)  
n(best < all) = 15 (p=0.906 | r=1.000)

## number of reference genes in main and other OGs



## num OGs per refOG

