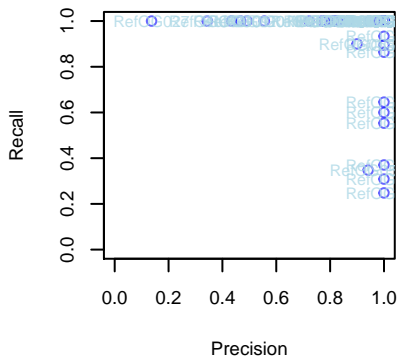
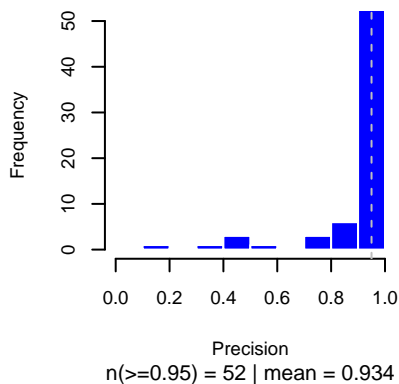


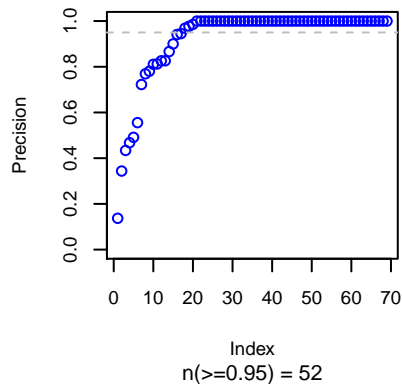
### Precision & recall



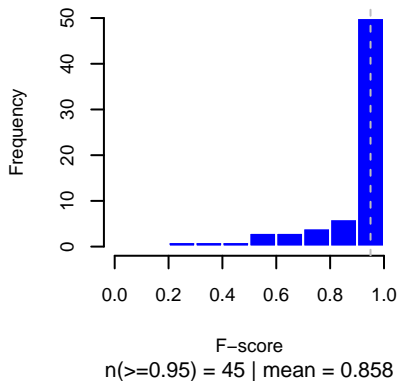
### Precision



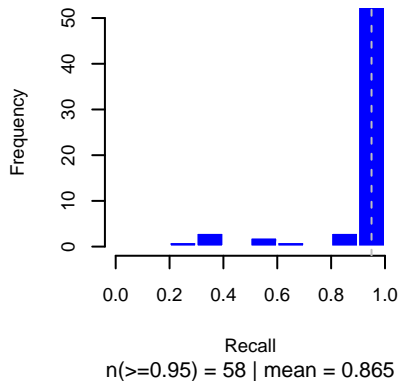
### Precision



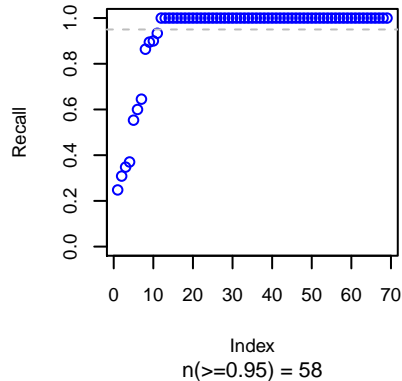
### F-score



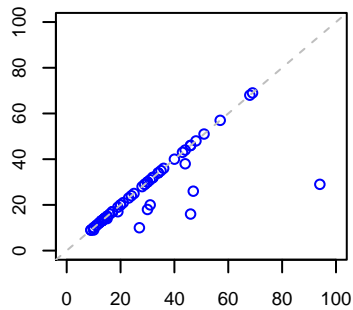
### Recall



### Recall

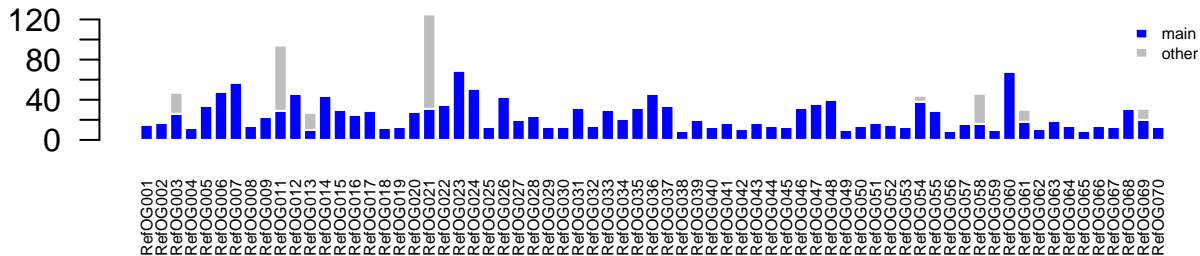


# genes in best OG



n(best = all) = 66 (p=0.997 | r=0.997)  
n(best = all) = 58 (p=0.909 | r=1.000)  
n(best < all) = 11 (p=0.986 | r=0.606)

## number of reference genes in main and other OGs



## num OGs per refOG

