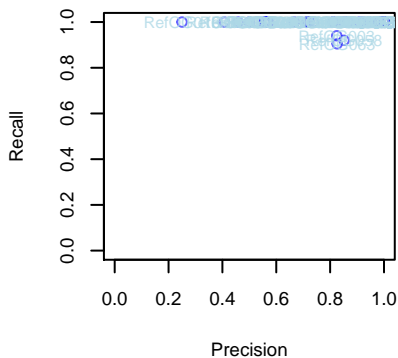
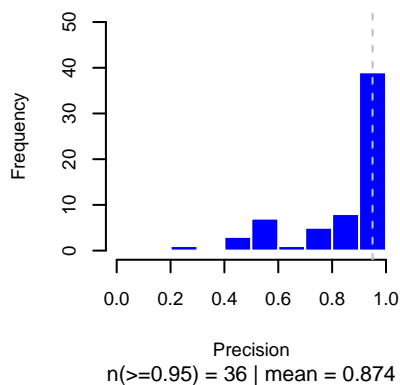


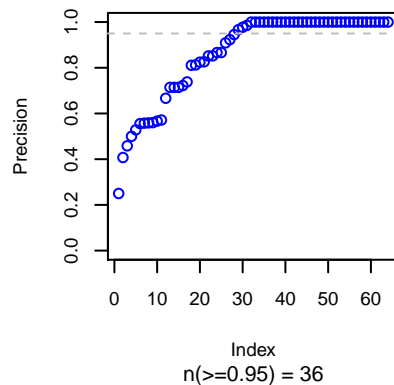
### Precision & recall



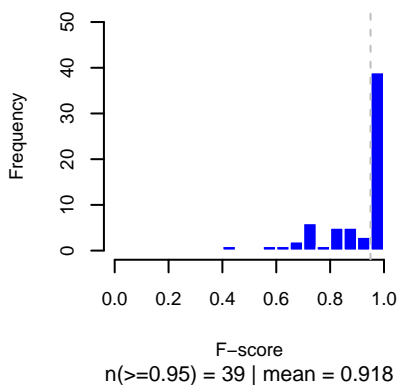
### Precision



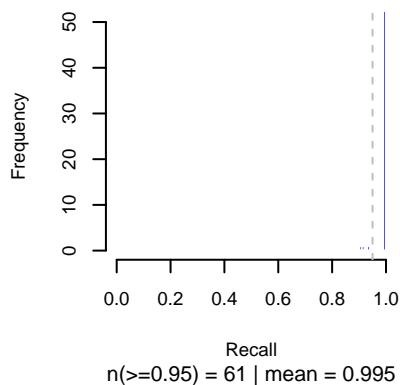
### Precision



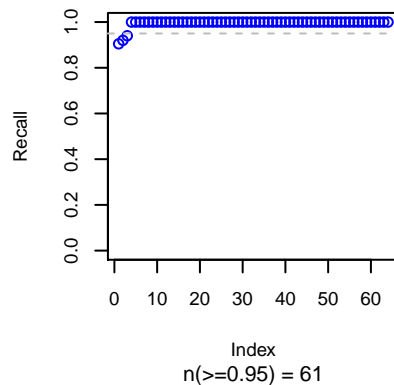
### F-score



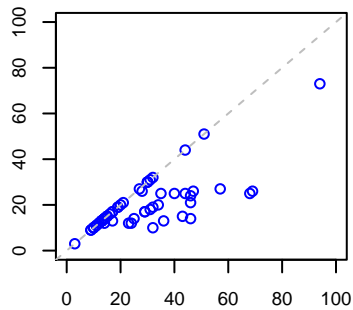
### Recall



### Recall

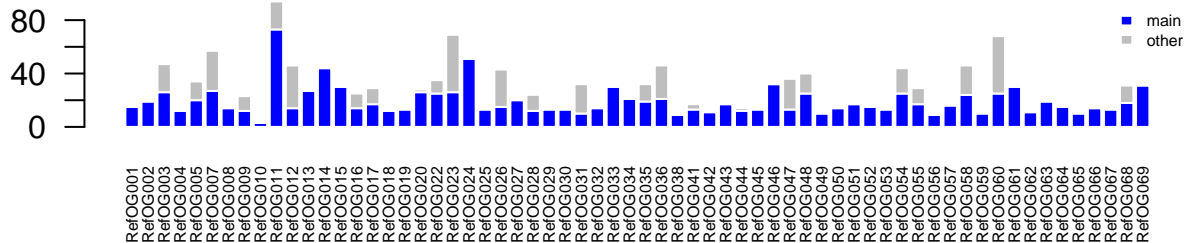


# genes in best OG



n(best < all) = 46 (p=0.998)  
n(best = all) = 39 (p=0.856 | r=0.998)  
n(best > all) = 25 (p=0.873 | r=0.994)

## number of reference genes in main and other OGs



## num OGs per refOG

