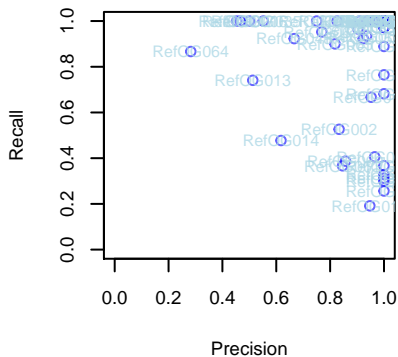
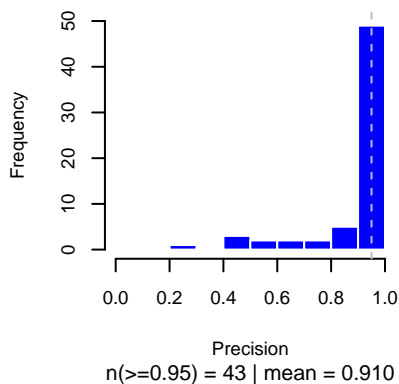


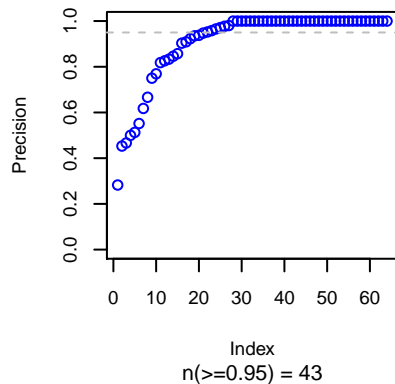
### Precision & recall



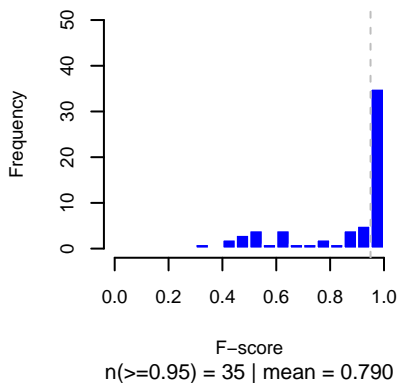
### Precision



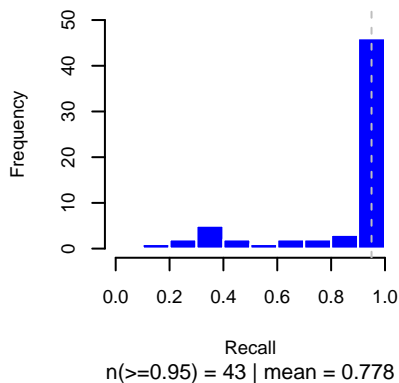
### Precision



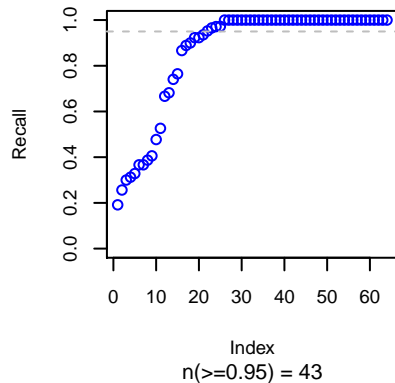
### F-score



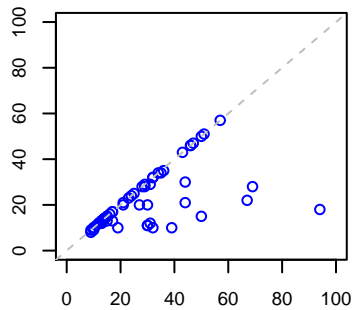
### Recall



### Recall

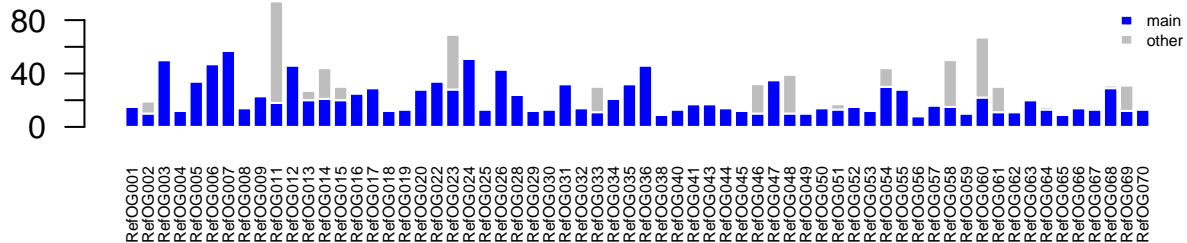


# genes in best OG



n(best = all) = 47 (p=0.920 | r=0.990)  
n(best = all) = 39 (p=0.930 | r=1.000)  
n(best < all) = 25 (p=0.873 | r=0.643)

## number of reference genes in main and other OGs



## num OGs per refOG

