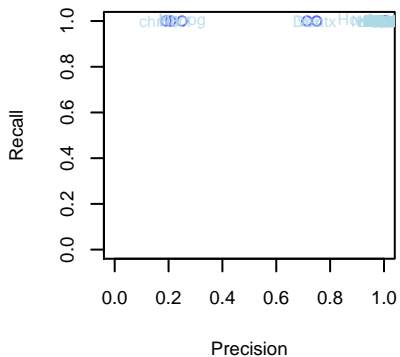
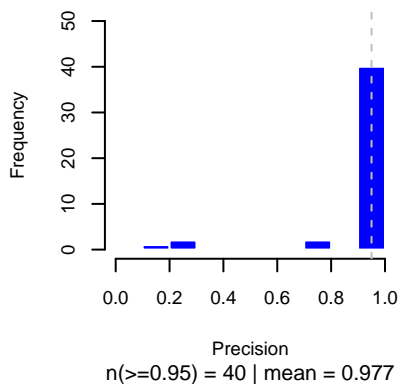


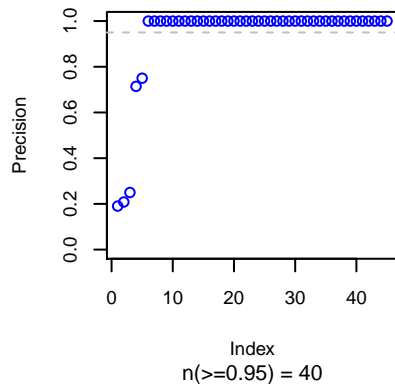
### Precision & recall



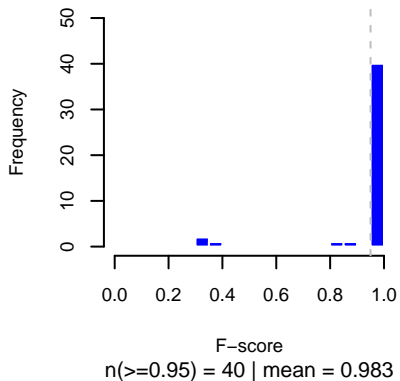
### Precision



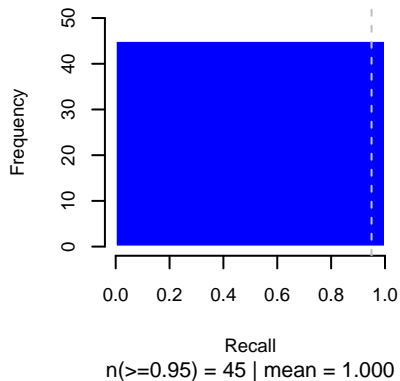
### Precision



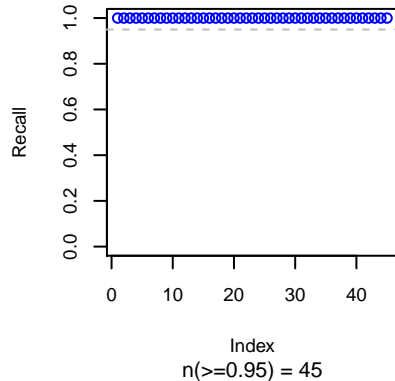
### F-score



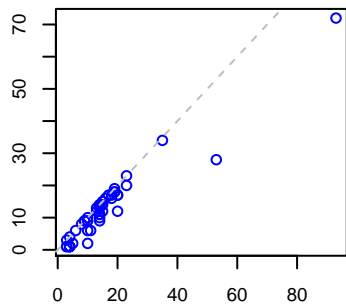
### Recall



### Recall

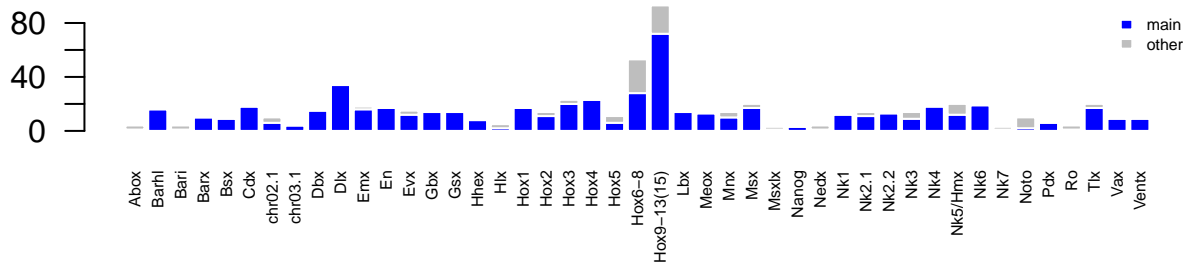


# genes in best OG



n(best = all) = 23 (p=0.000 | r=1.000)  
n(best = all) = 15 (p=0.860 | r=1.000)  
n(best < all) = 30 (p=0.974 | r=1.000)

## number of reference genes in main and other OGs



## num OGs per refOG

