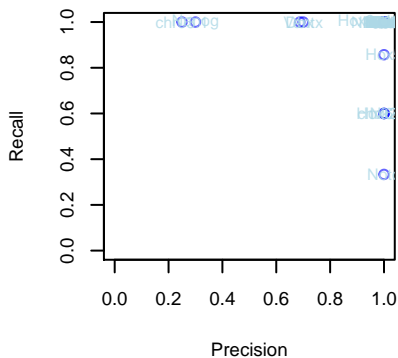
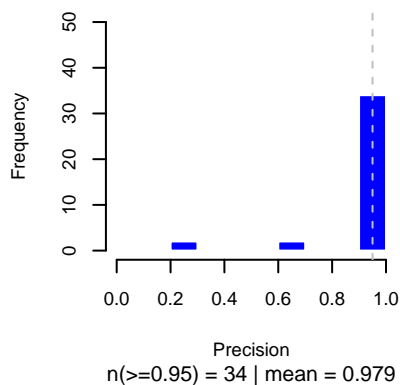


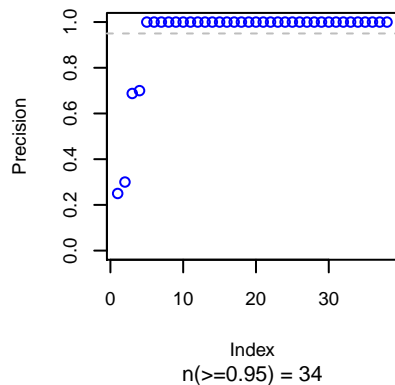
### Precision & recall



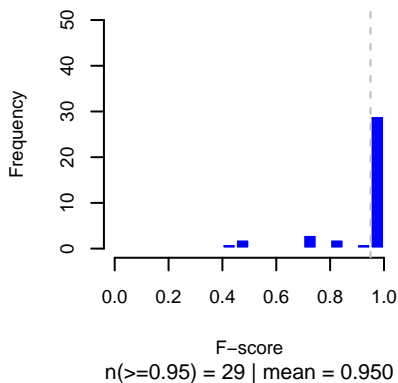
### Precision



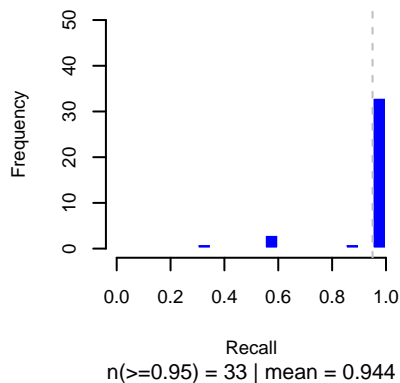
### Precision



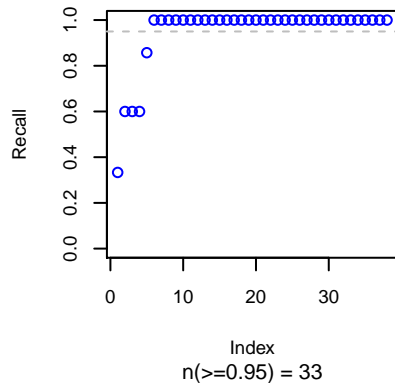
### F-score



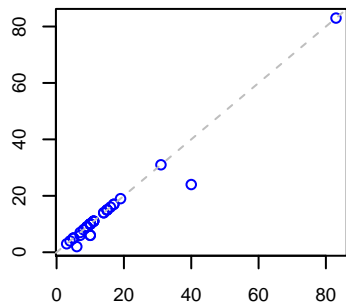
### Recall



### Recall

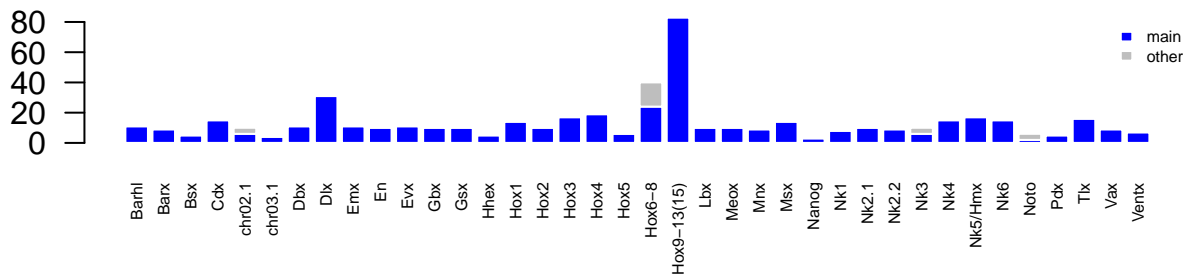


# genes in best OG



n(best = all) = 23 (p=1.000 | r=1.000)  
n(best = all) = 33 (p=0.938 | r=1.000)  
n(best < all) = 5 (p=1.000 | r=0.598)

## number of reference genes in main and other OGs



## num OGs per refOG

