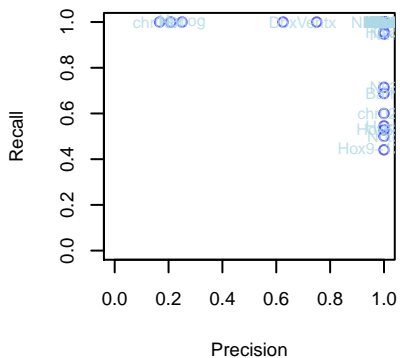
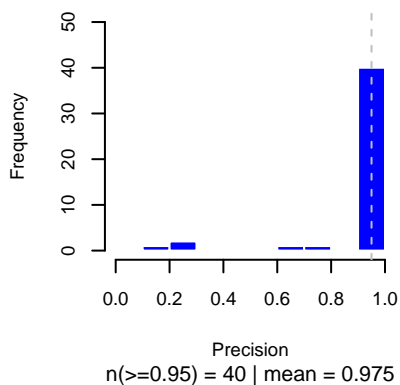


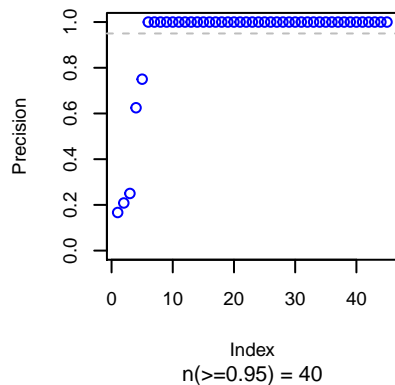
### Precision & recall



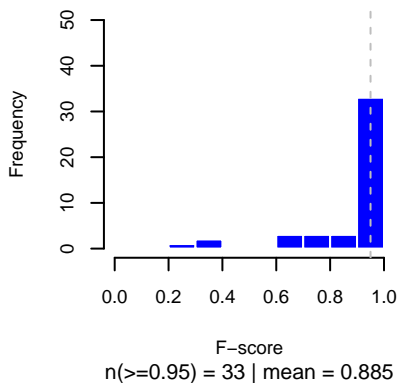
### Precision



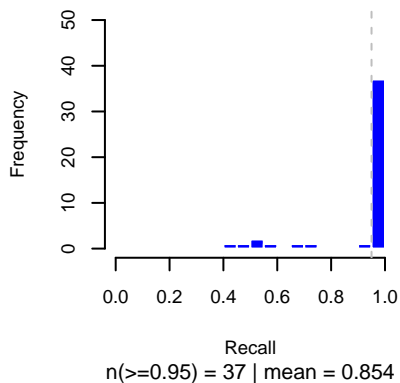
### Precision



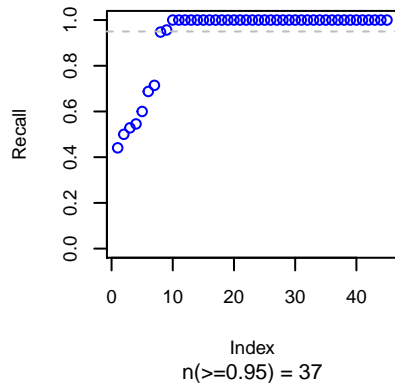
### F-score



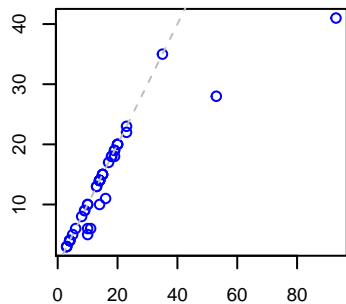
### Recall



### Recall

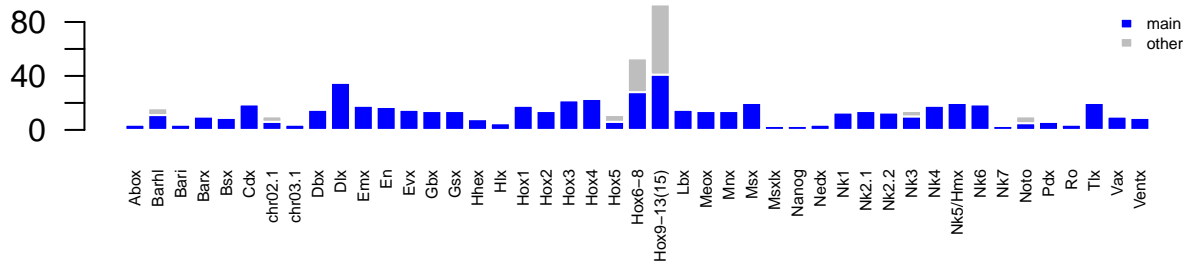


# genes in best OG



n(best = all) = 28 (p=0.920 | r=0.997)  
n(best = all) = 36 (p=0.917 | r=1.000)  
n(best < all) = 9 (p=1.000 | r=0.658)

## number of reference genes in main and other OGs



## num OGs per refOG

