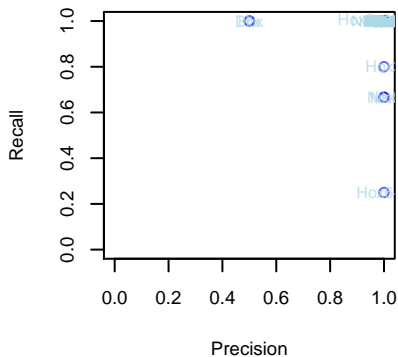
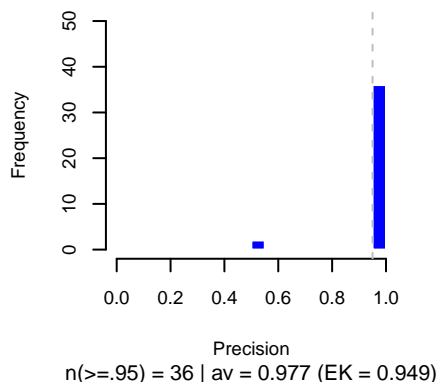


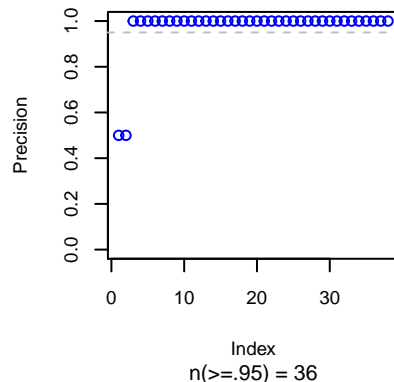
### Precision & recall



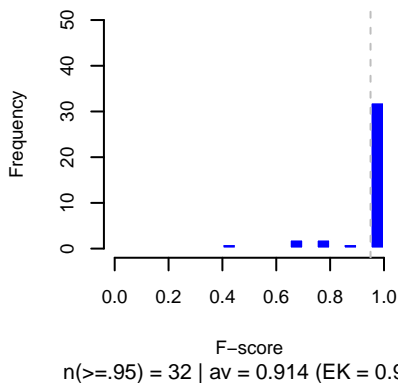
### Precision



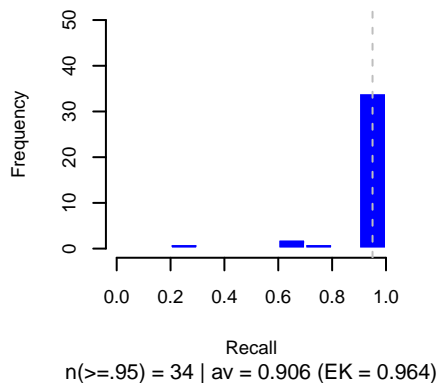
### Precision



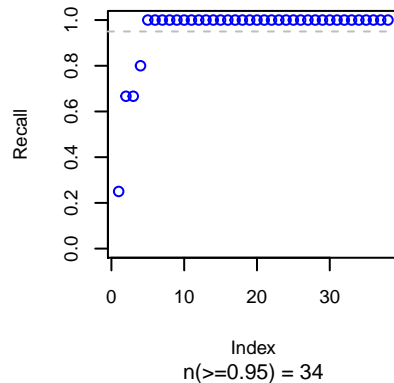
### F-score



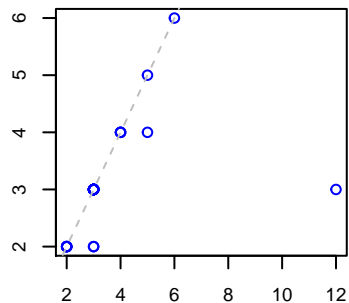
### Recall



### Recall

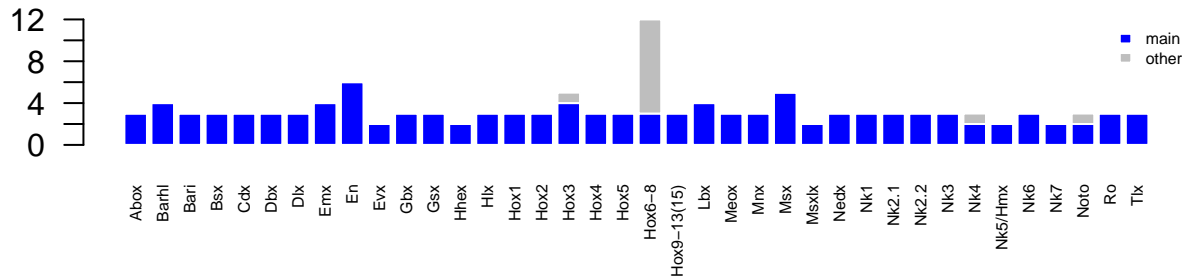


# genes in best OG



n(best = 2) = 2 (p=1.000 | r=1.000)  
n(best = all) = 34 (p=0.971 | r=1.000)  
n(best < all) = 4 (p=1.000 | r=0.596)

## number of reference genes in main and other OGs



## num OGs per refOG

