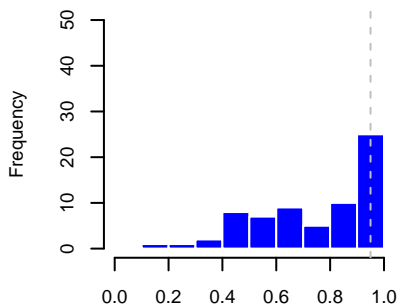
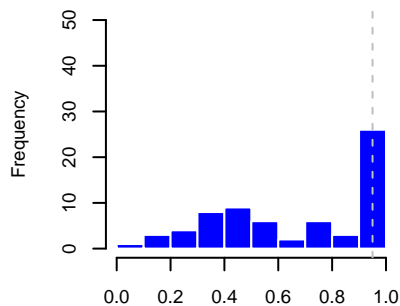
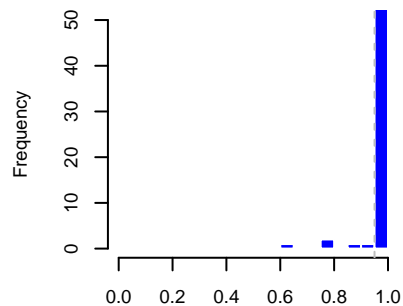


**F-score**

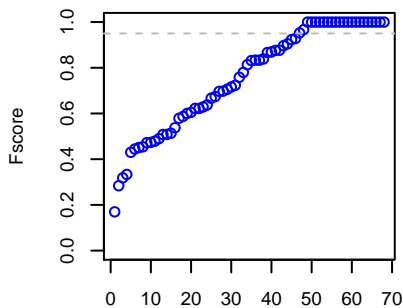
F-score  
av = 0.829 (inc = 0.831)

**Precision**

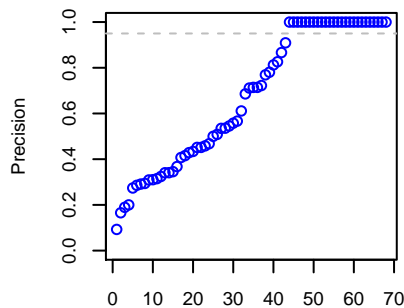
Precision  
av = 0.737 (inc = 0.729)

**Recall**

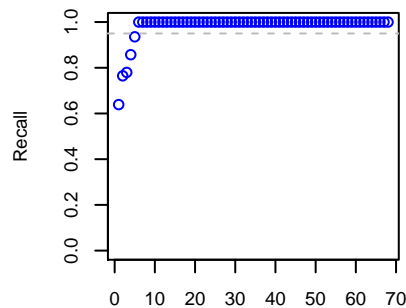
Recall  
av = 0.984 (inc = 0.996)

**Fscore**

Index  
n(>=.95) = 22

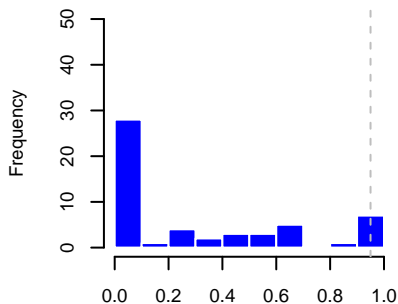
**Precision**

Index  
n(>=.95) = 25

**Recall**

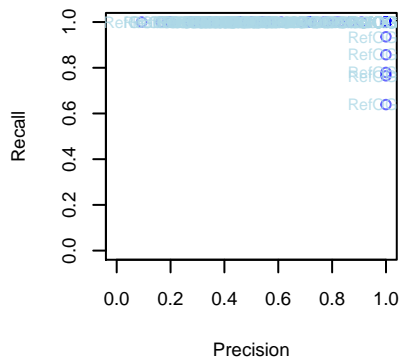
Index  
n(>=.95) = 63

### Rand index

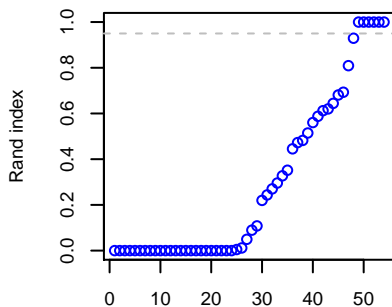


Rand index  
av = 0.373 (inc = 0.373) | med = 0.069

### Precision & recall

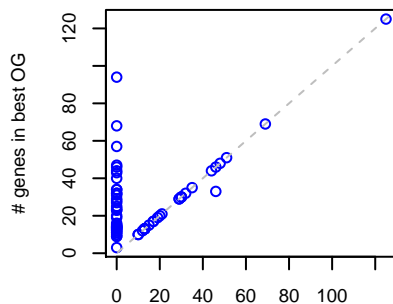


### Rand index



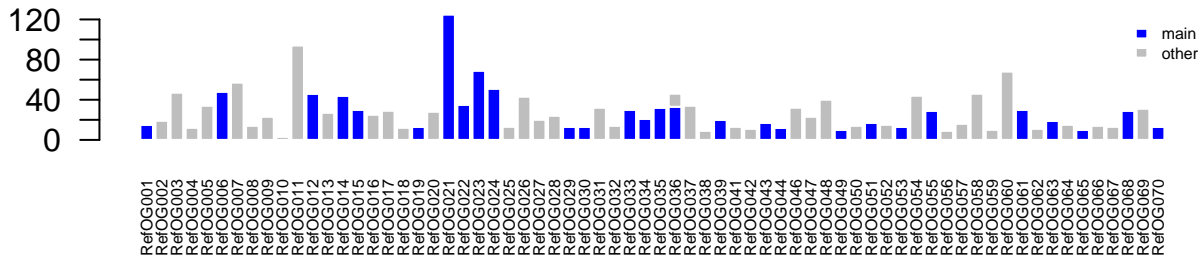
Index  
n(>=0.95) = NA

### size OGs



# genes in best OG  
n(best = all) = 27 (p=0.773 | r=0.992)  
n(best < all) = 1 (p=1.000 | r=0.780)

## number of reference genes in main and other OGs



## num OGs per refOG

