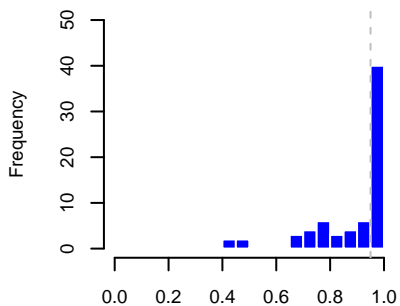
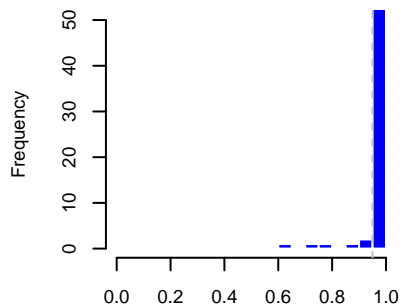
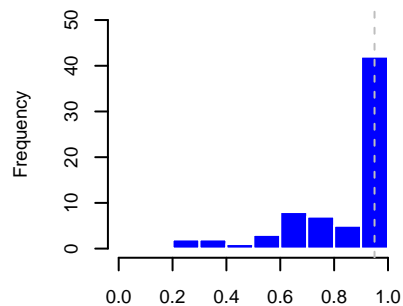


**F-score**

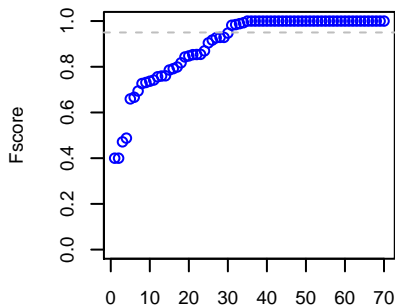
F-score  
av = 0.897 (inc = 0.946)

**Precision**

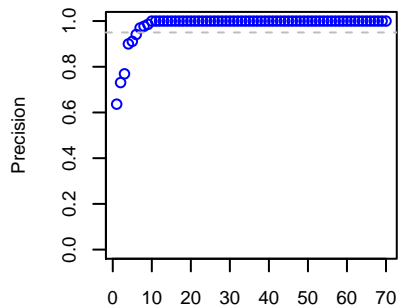
Precision  
av = 0.983 (inc = 0.985)

**Recall**

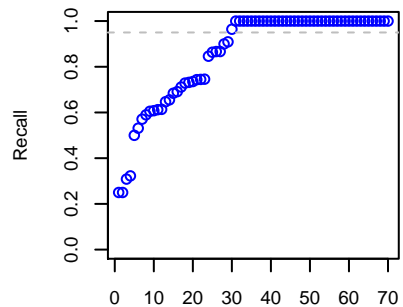
Recall  
av = 0.854 (inc = 0.937)

**Fscore**

Index  
n(>=.95) = 40

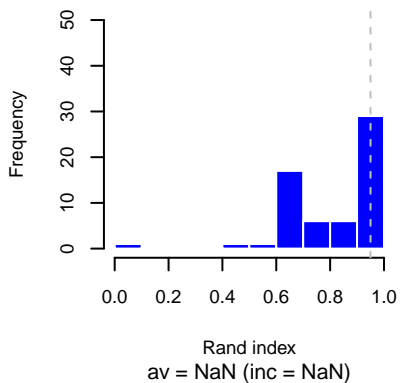
**Precision**

Index  
n(>=.95) = 64

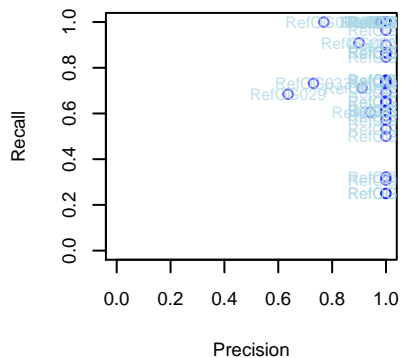
**Recall**

Index  
n(>=.95) = 41

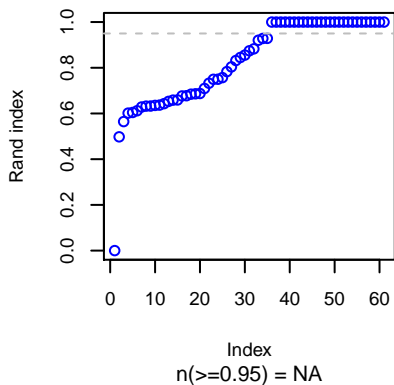
### Rand index



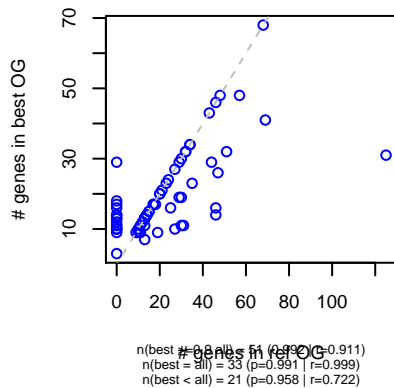
### Precision & recall



### Rand index

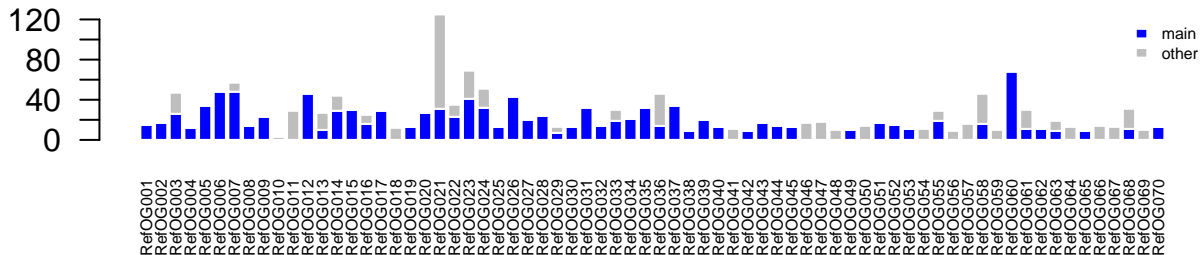


### size OGs



n(best = all) = 33 (p=0.991 | r=0.999)  
 n(best < all) = 21 (p=0.958 | r=0.722)

## number of reference genes in main and other OGs



## num OGs per refOG

