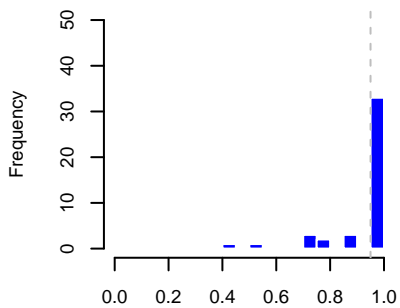
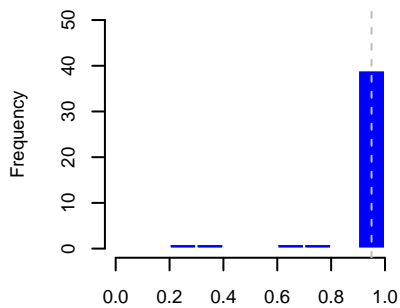
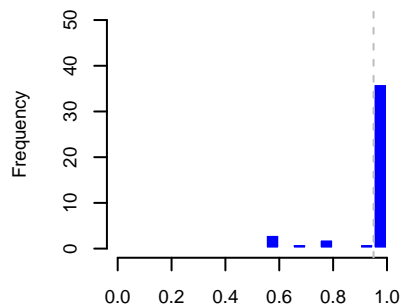


**F-score**

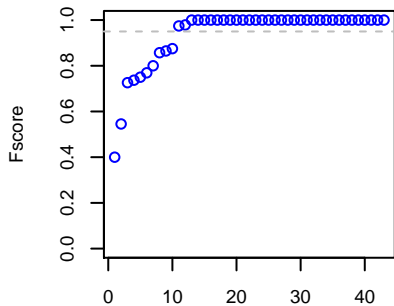
F-score  
av = 0.937 (inc = 0.967)

**Precision**

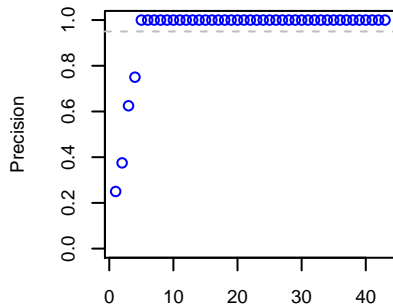
Precision  
av = 0.953 (inc = 0.953)

**Recall**

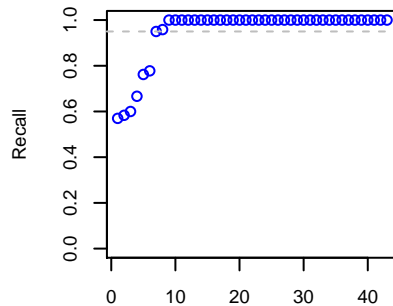
Recall  
av = 0.950 (inc = 1.000)

**Fscore**

Index  
n(>=.95) = 33

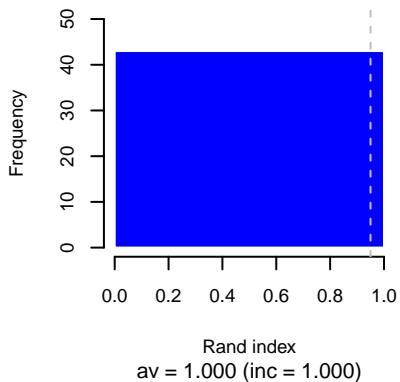
**Precision**

Index  
n(>=.95) = 39

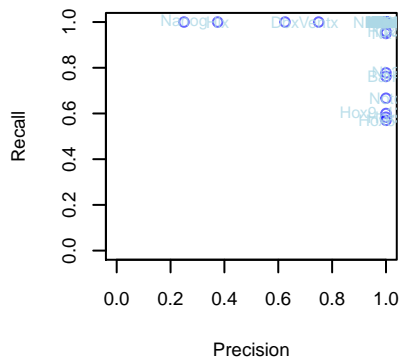
**Recall**

Index  
n(>=.95) = 37

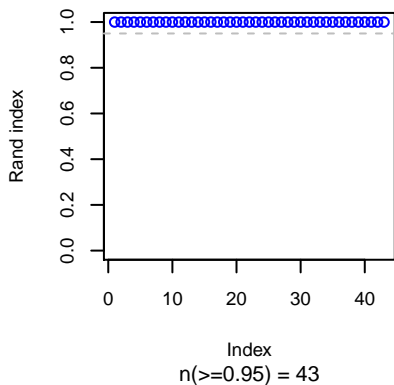
### Rand index



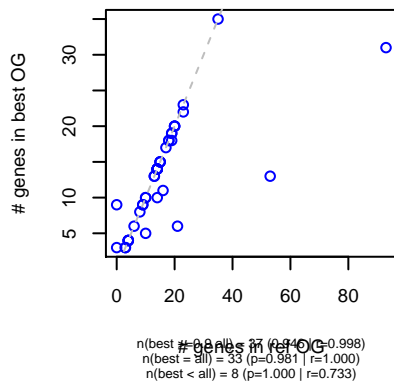
### Precision & recall



### Rand index

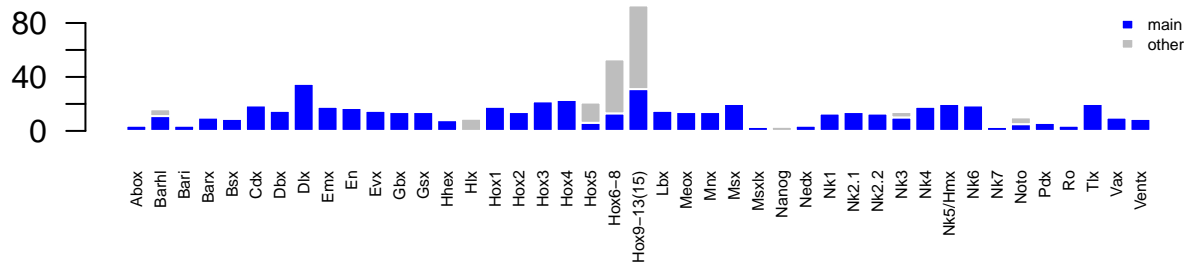


### size OGs



n(best = all) = 27 (p=0.998 | r=0.998)  
 n(best = all) = 33 (p=0.981 | r=1.000)  
 n(best < all) = 8 (p=1.000 | r=0.733)

## number of reference genes in main and other OGs



## num OGs per refOG

