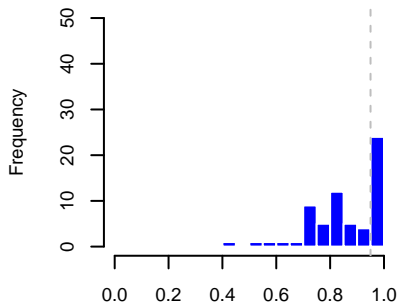
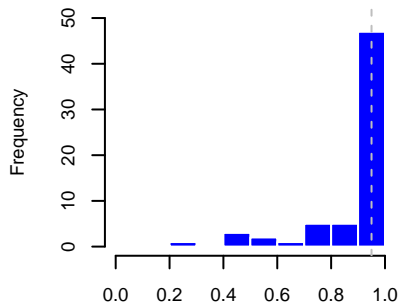
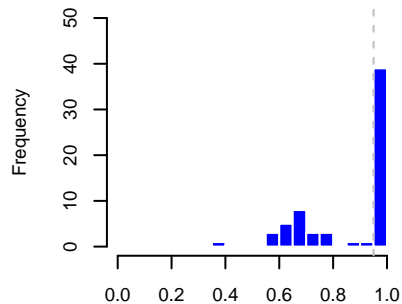


**F-score**

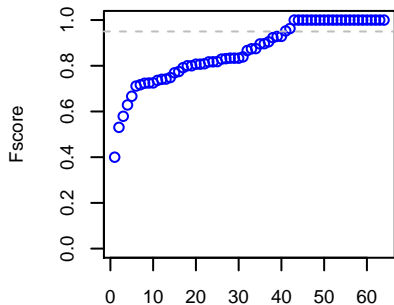
F-score  
av = 0.829 (inc = 0.941)

**Precision**

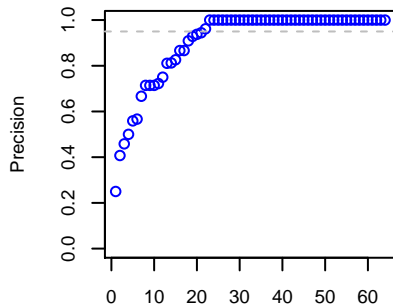
Precision  
av = 0.911 (inc = 0.911)

**Recall**

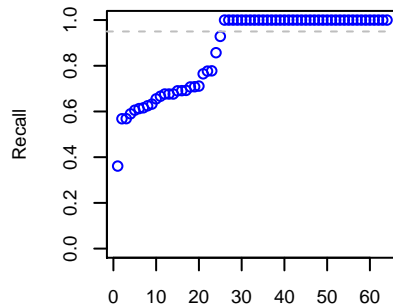
Recall  
av = 0.799 (inc = 0.988)

**Fscore**

Index  
n( $\geq .95$ ) = 24

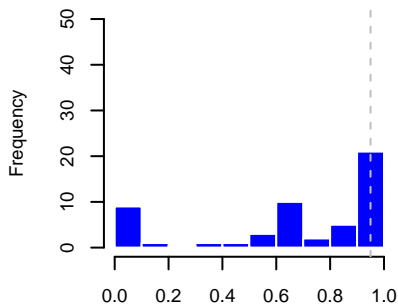
**Precision**

Index  
n( $\geq .95$ ) = 43

**Recall**

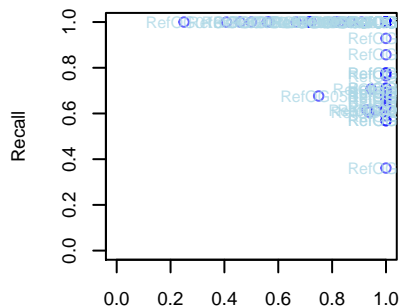
Index  
n( $\geq .95$ ) = 39

## Rand index



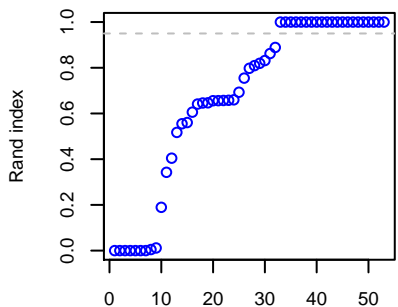
Rand index  
av = 0.695 (inc = 0.695) | med = 0.798

## Precision & recall



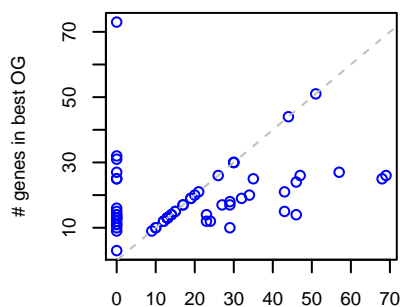
Precision

## Rand index



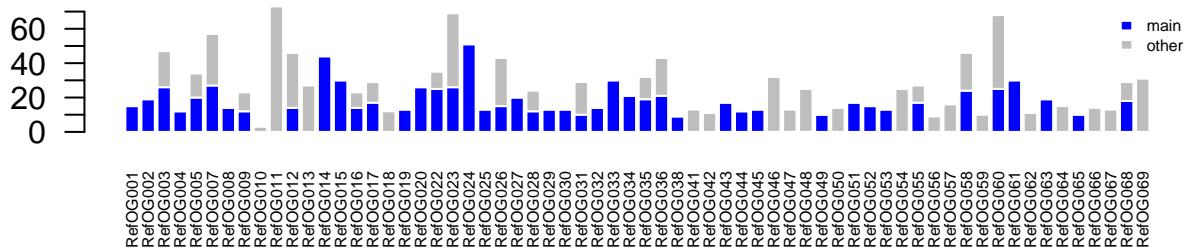
Index  
n(>=0.95) = NA

## size OGs



# genes in best OG  
n(best = all) = 27 (p=0.861 | r=0.992)  
n(best < all) = 18 (p=0.973 | r=0.664)

## number of reference genes in main and other OGs



## num OGs per refOG

