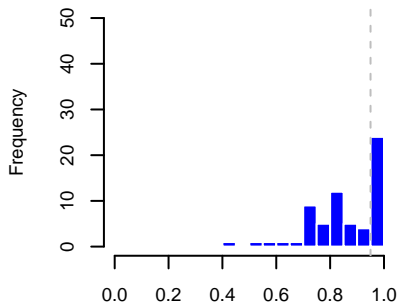
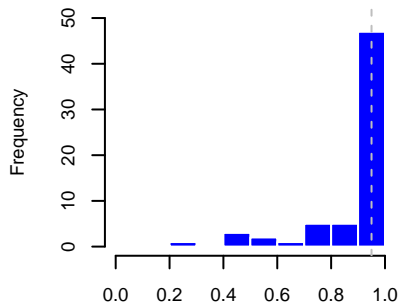
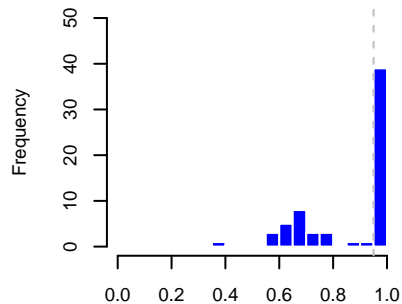


**F-score**

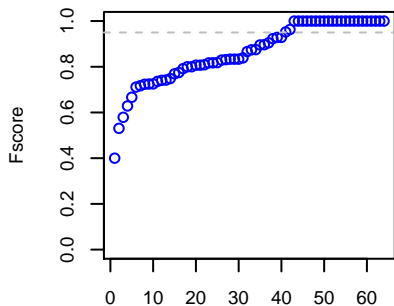
F-score  
av = 0.829 (inc = 0.911)

**Precision**

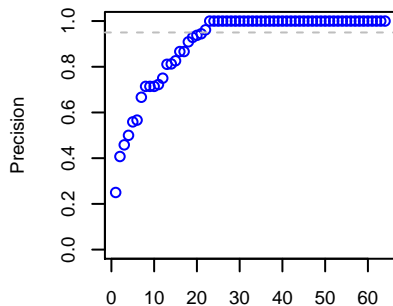
Precision  
av = 0.916 (inc = 0.916)

**Recall**

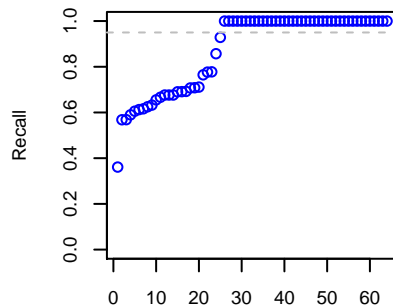
Recall  
av = 0.803 (inc = 0.942)

**Fscore**

Index  
n(>=.95) = 24

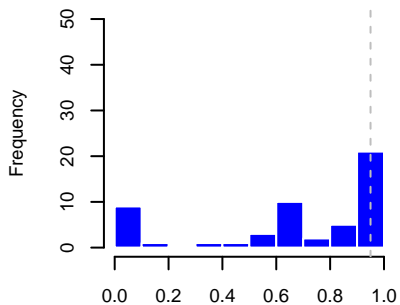
**Precision**

Index  
n(>=.95) = 43

**Recall**

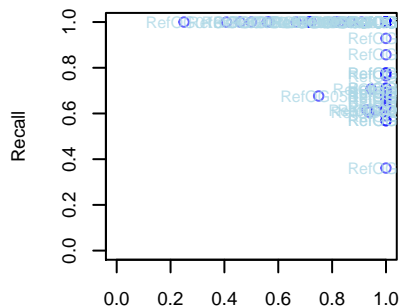
Index  
n(>=.95) = 39

## Rand index



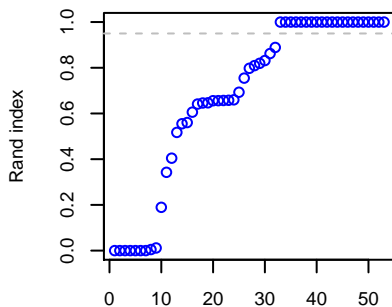
av = NaN (inc = NaN) | med = 0.798

## Precision & recall



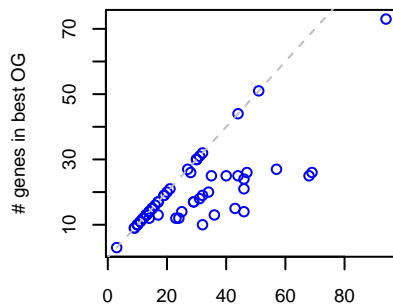
Precision

## Rand index



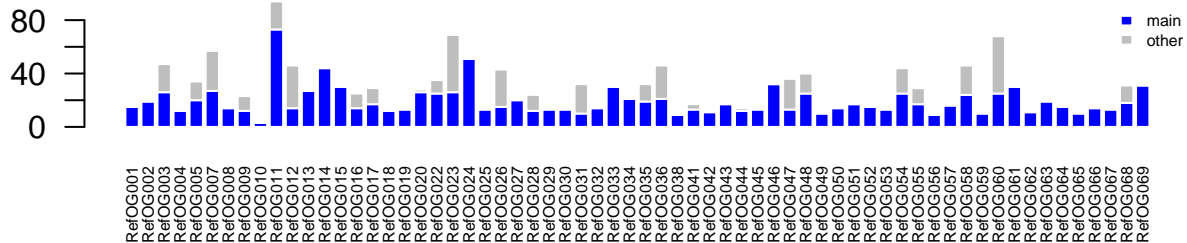
Index  
n(>=0.95) = NA

## size OGs



# genes in Ref OG  
n(best = all) = 39 (p=0.856 | r=1.000)  
n(best < all) = 25 (p=0.981 | r=0.673)

## number of reference genes in main and other OGs



## num OGs per refOG

