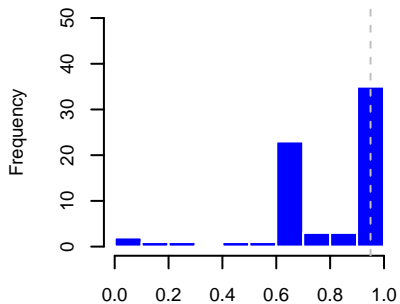
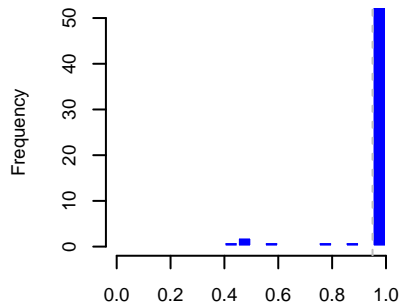
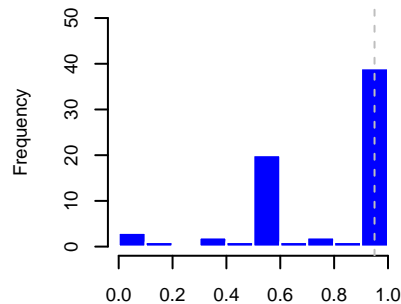


**F-score**

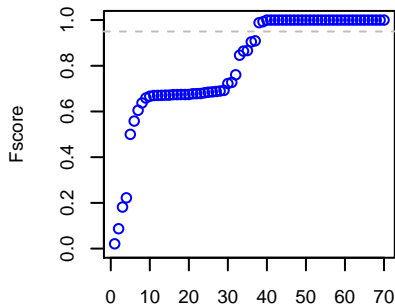
F-score  
av = 0.763 (inc = 0.879)

**Precision**

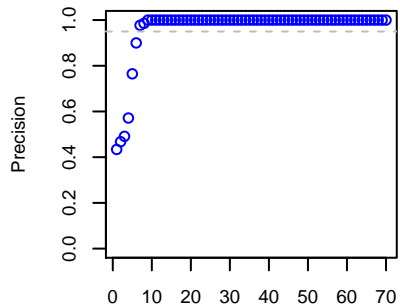
Precision  
av = 0.968 (inc = 0.967)

**Recall**

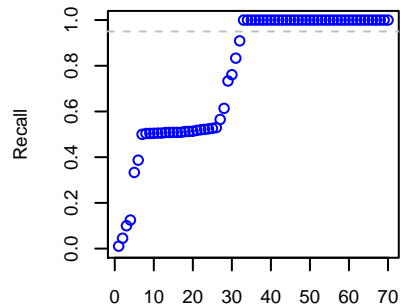
Recall  
av = 0.714 (inc = 0.891)

**Fscore**

Index  
n(>=.95) = 33

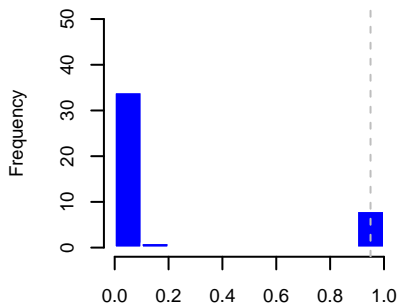
**Precision**

Index  
n(>=.95) = 64

**Recall**

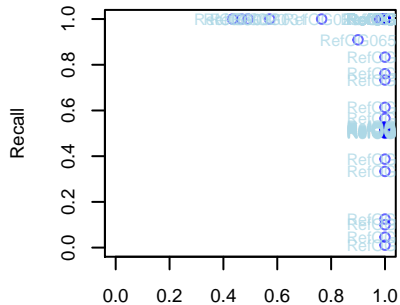
Index  
n(>=.95) = 38

### Rand index



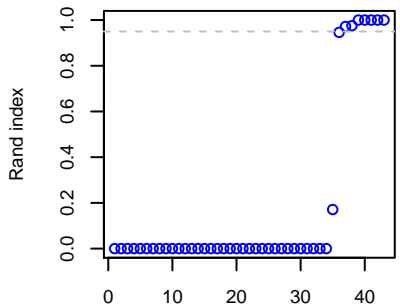
av = NaN (inc = NaN) | med = 0.000

### Precision & recall



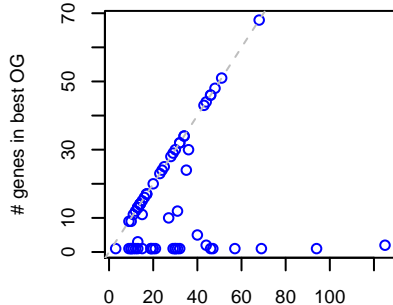
Precision

### Rand index



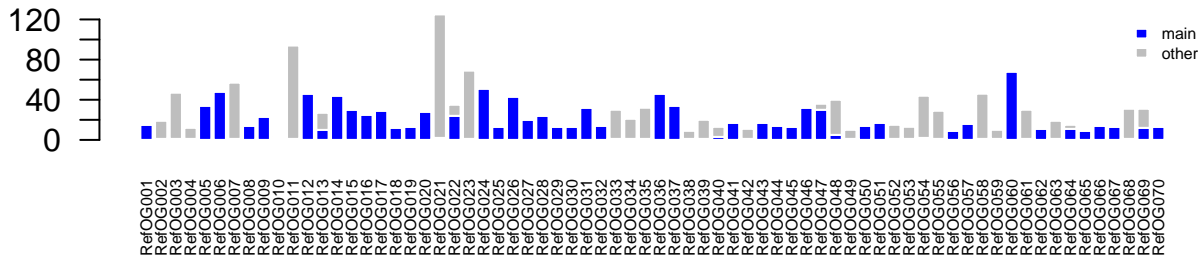
Index  
n(>=0.95) = NA

### size OGs



# genes in Ref OG  
n(best = all) = 38 (p=0.939 | r=0.998)  
n(best < all) = 32 (p=0.997 | r=0.489)

## number of reference genes in main and other OGs



## num OGs per refOG

