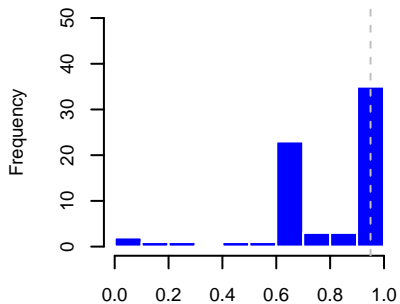
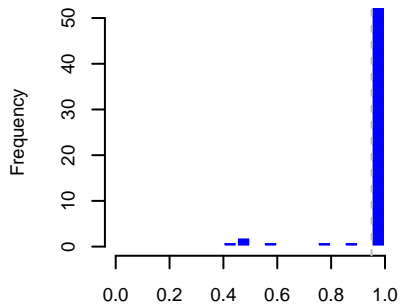
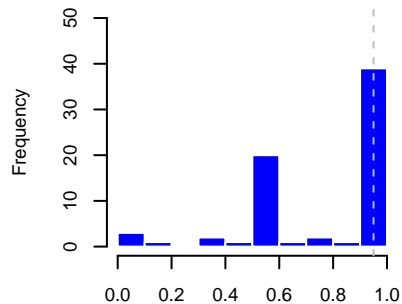


**F-score**

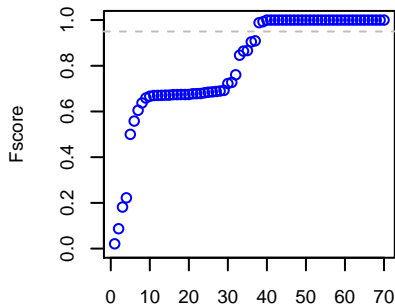
F-score  
av = 0.837 (inc = 0.991)

**Precision**

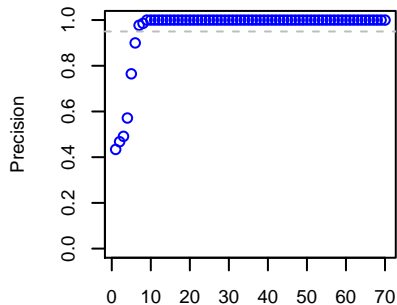
Precision  
av = 0.986 (inc = 0.986)

**Recall**

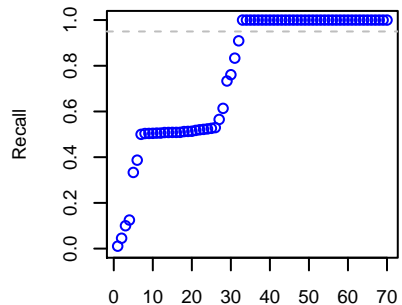
Recall  
av = 0.765 (inc = 0.999)

**Fscore**

Index  
n(>=.95) = 33

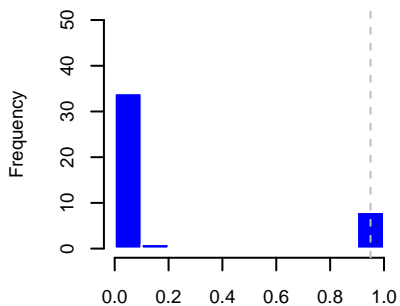
**Precision**

Index  
n(>=.95) = 64

**Recall**

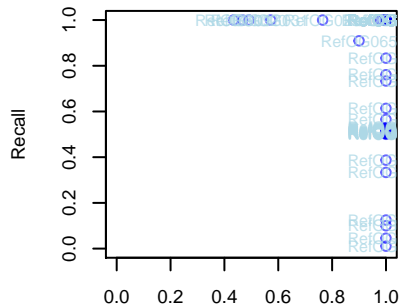
Index  
n(>=.95) = 38

### Rand index



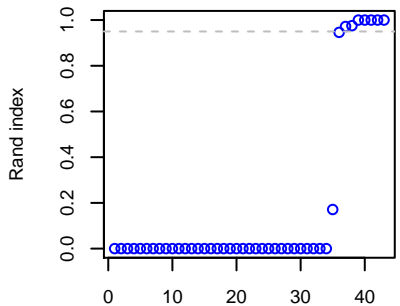
Rand index  
av = NaN (inc = NaN) | med = 0.000

### Precision & recall



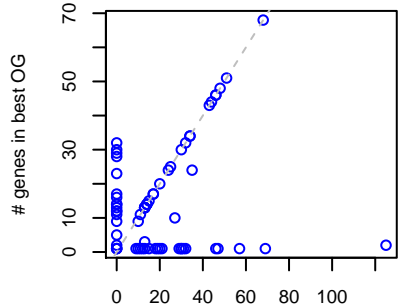
Precision

### Rand index



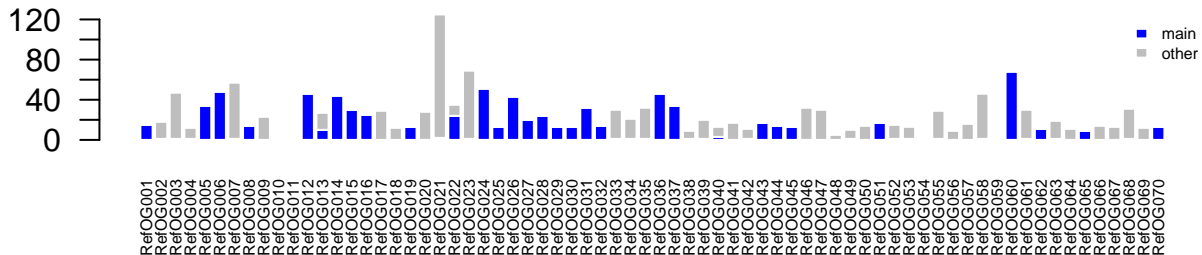
Index  
n(>=0.95) = NA

### size OGs



# genes in best OG  
n(best < all) = 17 (p=0.882)  
n(best = all) = 27 (p=0.974 | r=1.000)  
n(best > all) = 24 (p=0.996 | r=0.545)

## number of reference genes in main and other OGs



## num OGs per refOG

