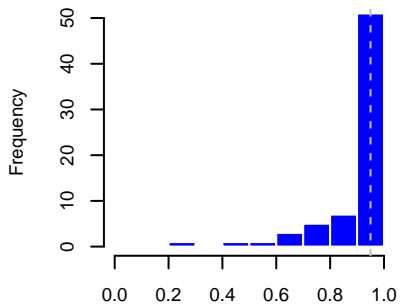
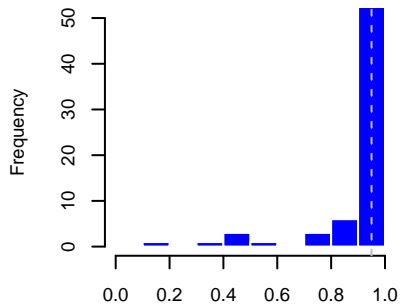
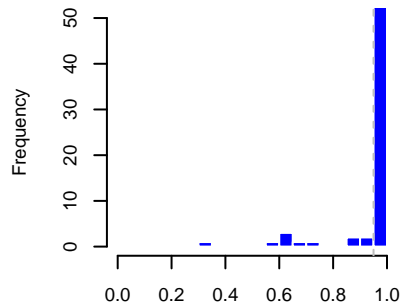


**F-score**

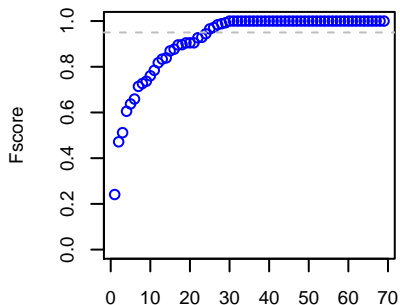
F-score  
av = 0.930 (inc = 0.974)

**Precision**

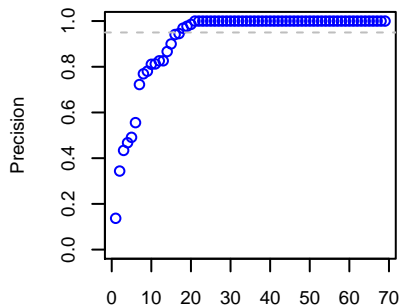
Precision  
av = 0.958 (inc = 0.955)

**Recall**

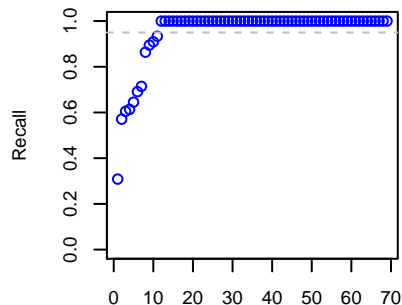
Recall  
av = 0.924 (inc = 0.999)

**Fscore**

Index  
n(>=.95) = 45

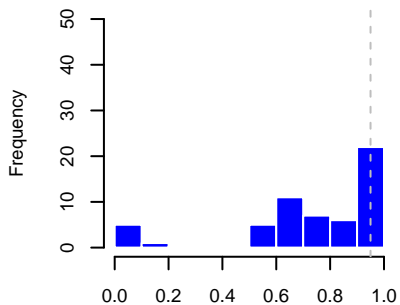
**Precision**

Index  
n(>=.95) = 52

**Recall**

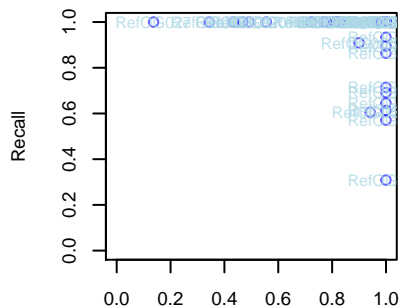
Index  
n(>=.95) = 58

### Rand index

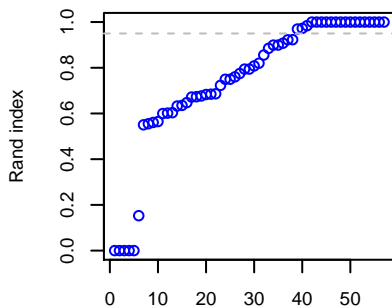


av = 0.750 (inc = 0.750) | med = 0.794

### Precision & recall

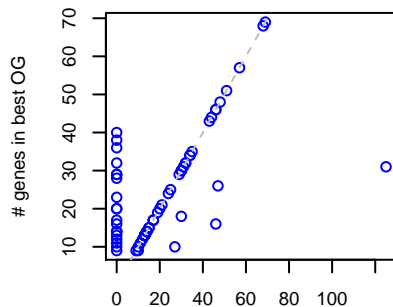


### Rand index



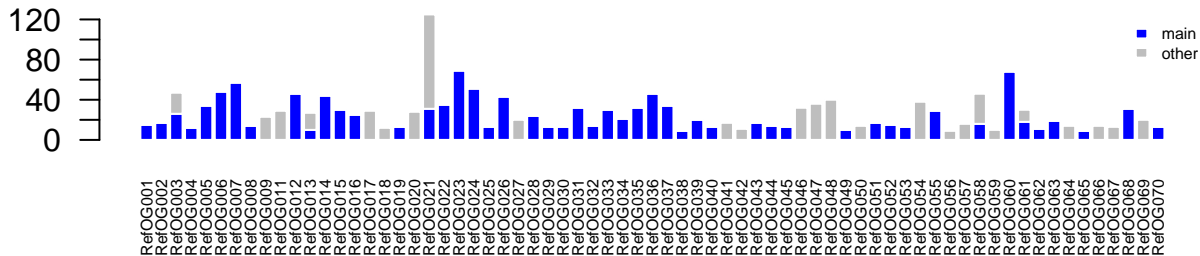
Index  
n(>=0.95) = NA

### size OGs



# genes in ref OG  
n(best = all) = 43 (p=0.950 | r=0.998)  
n(best < all) = 6 (p=0.974 | r=0.684)

## number of reference genes in main and other OGs



## num OGs per refOG

