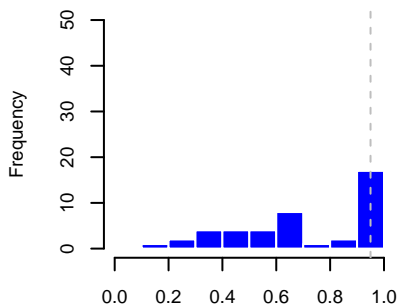
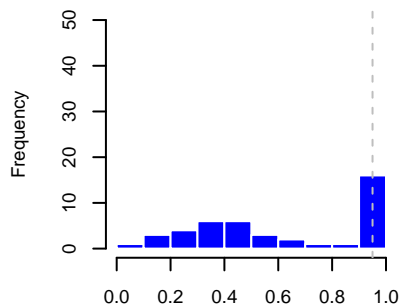
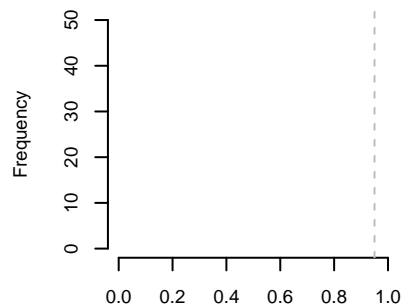


**F-score**

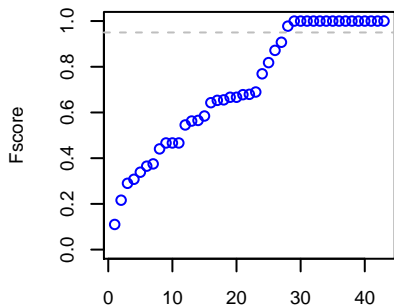
F-score  
av = 0.716 (inc = 0.716)

**Precision**

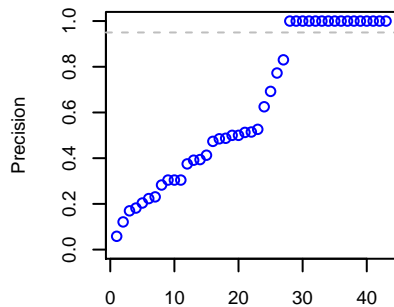
Precision  
av = 0.625 (inc = 0.625)

**Recall**

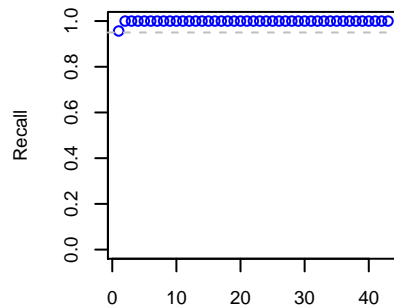
Recall  
av = 0.999 (inc = 0.999)

**Fscore**

Index  
n(>=.95) = 16

**Precision**

Index  
n(>=.95) = 16

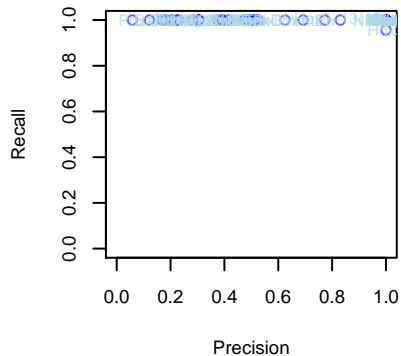
**Recall**

Index  
n(>=0.95) = 43

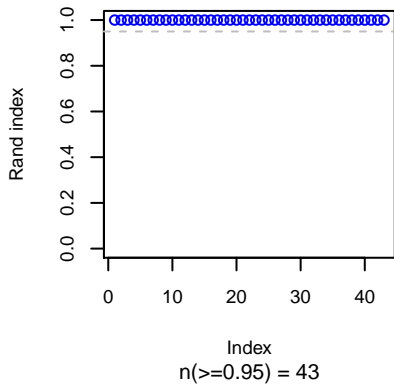
### Rand index



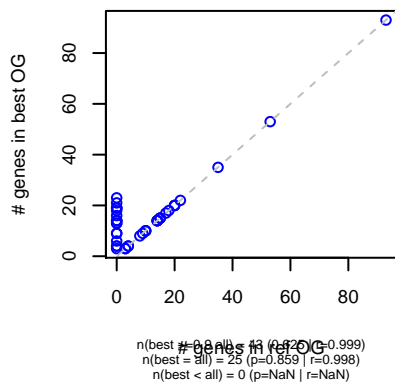
### Precision & recall



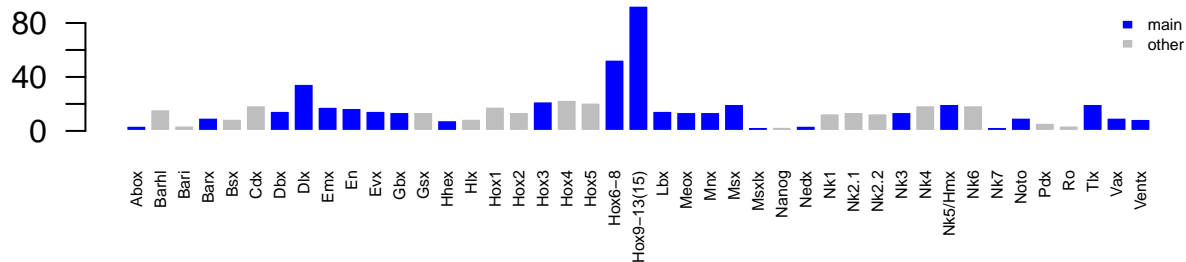
### Rand index



### size OGs



## number of reference genes in main and other OGs



## num OGs per refOG

