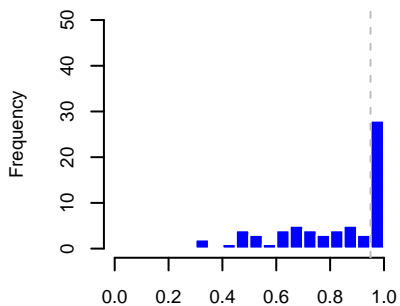
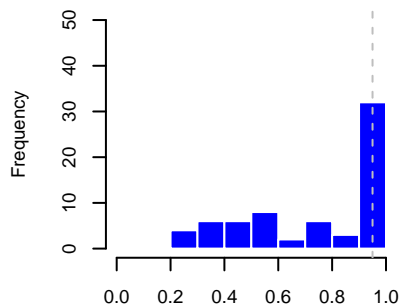
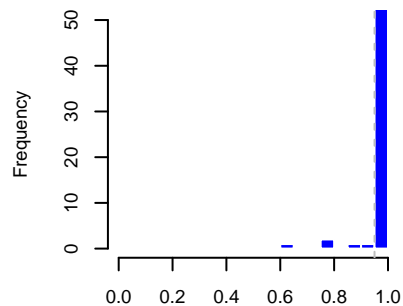


**F-score**

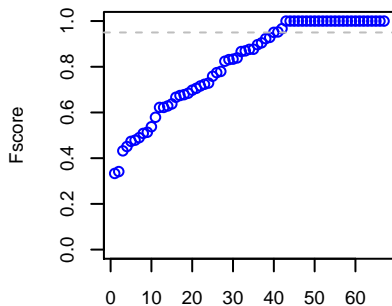
F-score  
av = 0.861 (inc = 0.863)

**Precision**

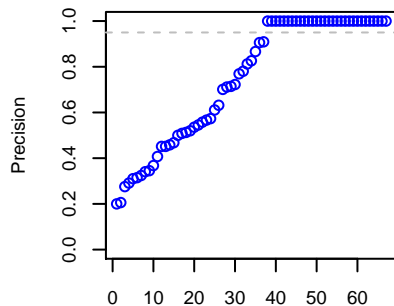
Precision  
av = 0.787 (inc = 0.781)

**Recall**

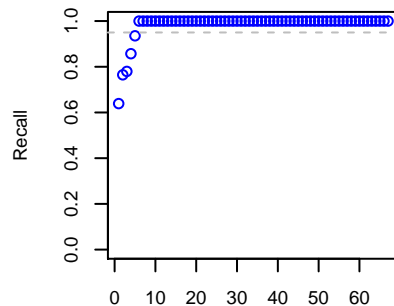
Recall  
av = 0.987 (inc = 0.997)

**Fscore**

Index  
n(>=.95) = 28

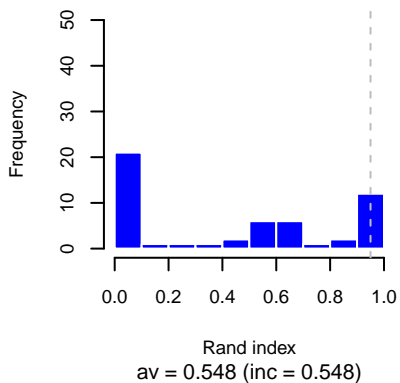
**Precision**

Index  
n(>=.95) = 30

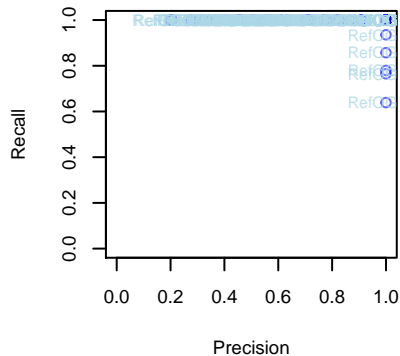
**Recall**

Index  
n(>=.95) = 62

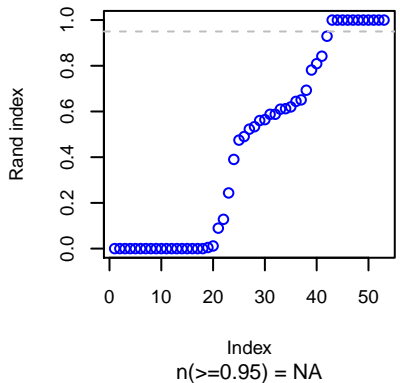
# Rand index



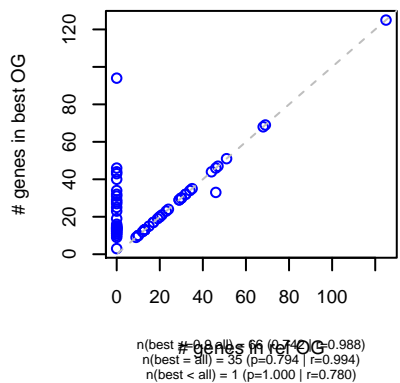
# Precision & recall



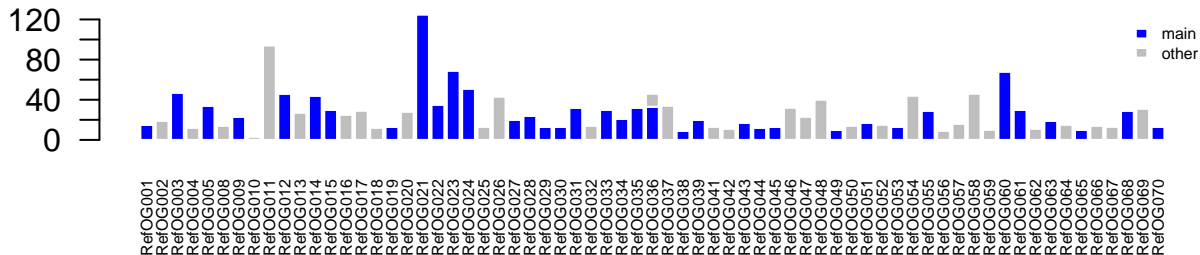
# Rand index



# size OGs



## number of reference genes in main and other OGs



## num OGs per refOG

