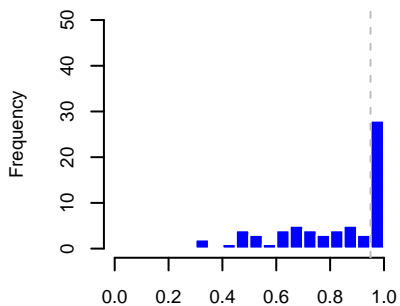
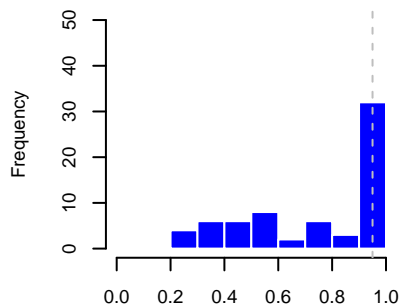
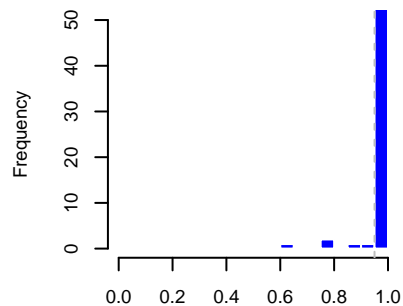


F-score

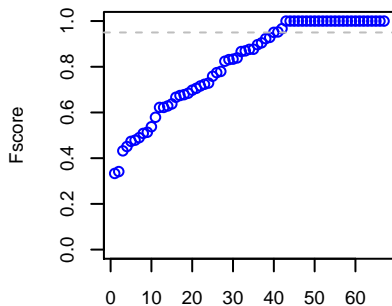
F-score
av = 0.861 (inc = 0.863)

Precision

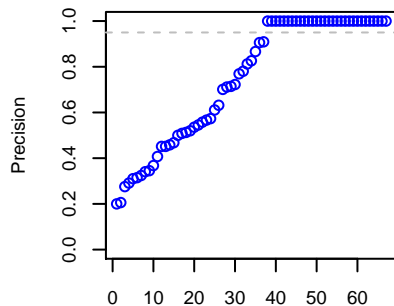
Precision
av = 0.787 (inc = 0.781)

Recall

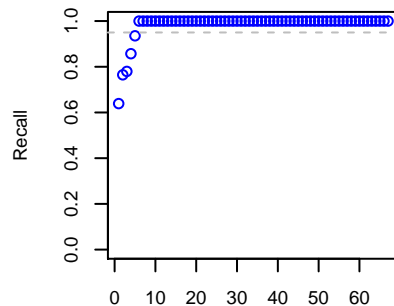
Recall
av = 0.987 (inc = 0.997)

Fscore

Index
 $n(\geq .95) = 28$

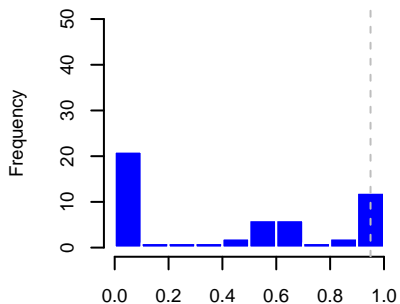
Precision

Index
 $n(\geq .95) = 30$

Recall

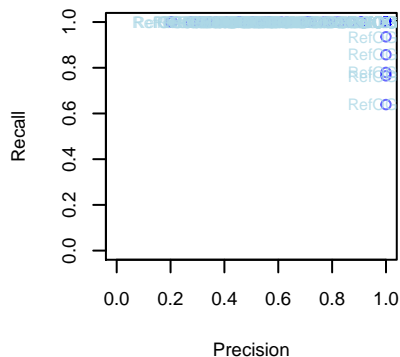
Index
 $n(\geq .95) = 62$

Rand index

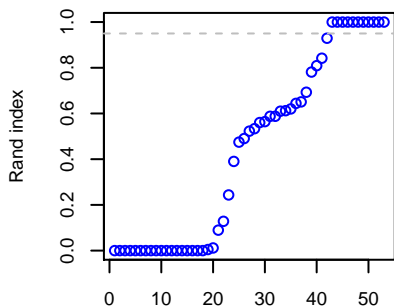


av = 0.548 (inc = 0.548) | med = 0.523

Precision & recall

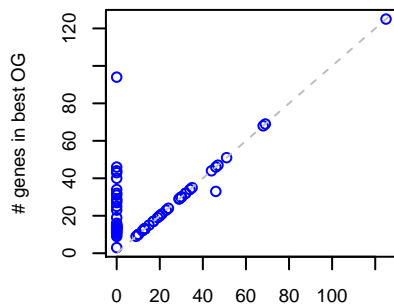


Rand index



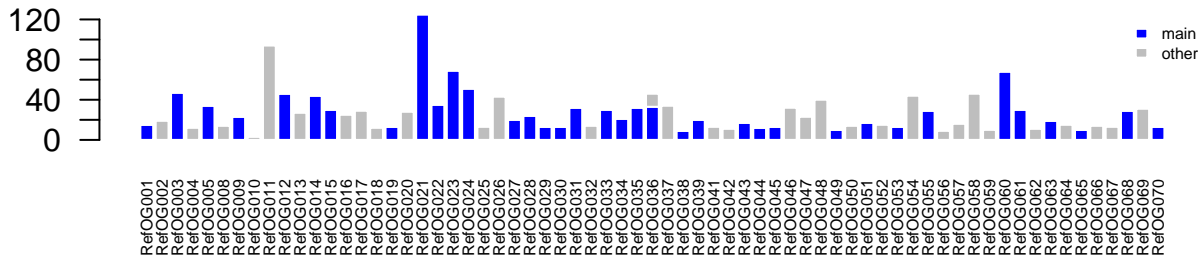
Index
n(>=0.95) = NA

size OGs



n(best = all) = 35 (p=0.794 | r=0.994)
n(best < all) = 1 (p=1.000 | r=0.780)

number of reference genes in main and other OGs



num OGs per refOG

