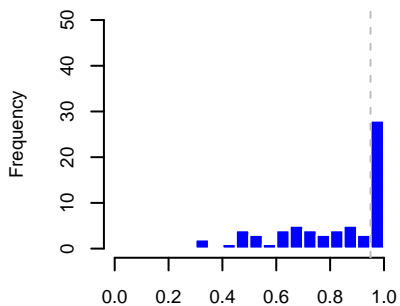
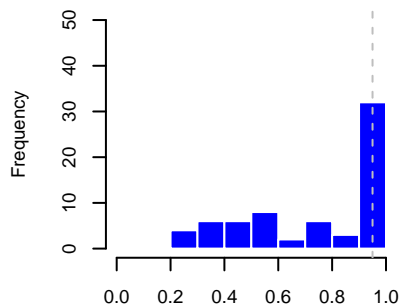
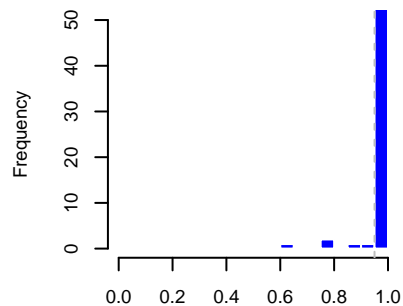


F-score

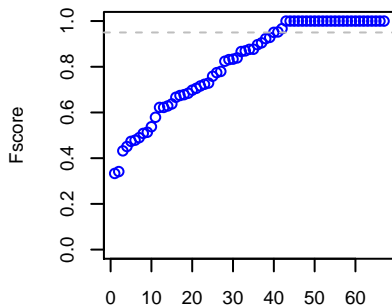
F-score
av = 0.820 (inc = 0.821)

Precision

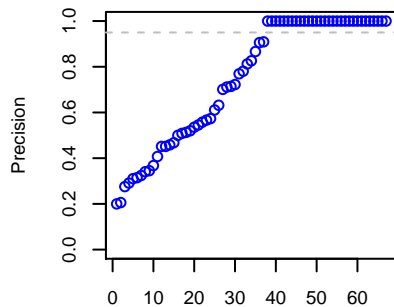
Precision
av = 0.751 (inc = 0.747)

Recall

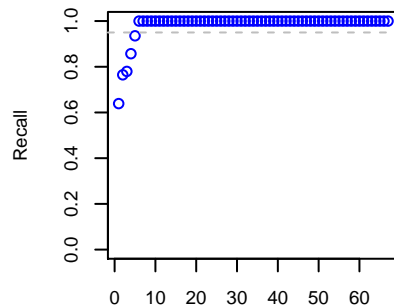
Recall
av = 0.983 (inc = 0.988)

Fscore

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

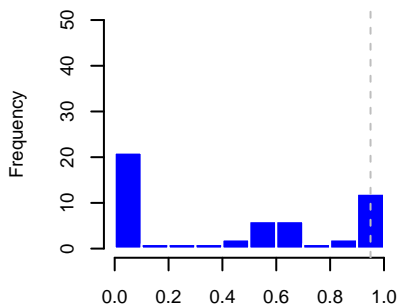
Precision

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

Recall

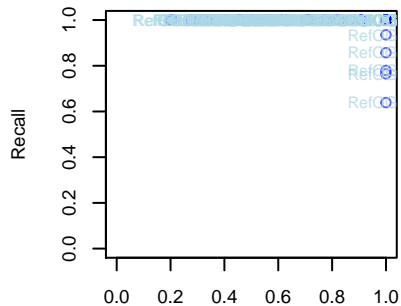
Index
 $n(\geq .95) = 62$

Rand index

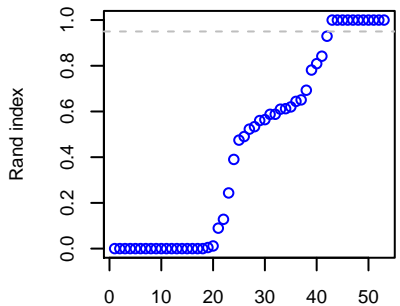


av = NaN (inc = NaN) | med = 0.523

Precision & recall

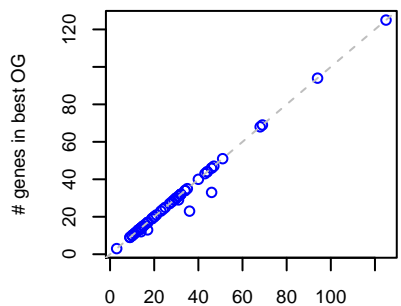


Rand index



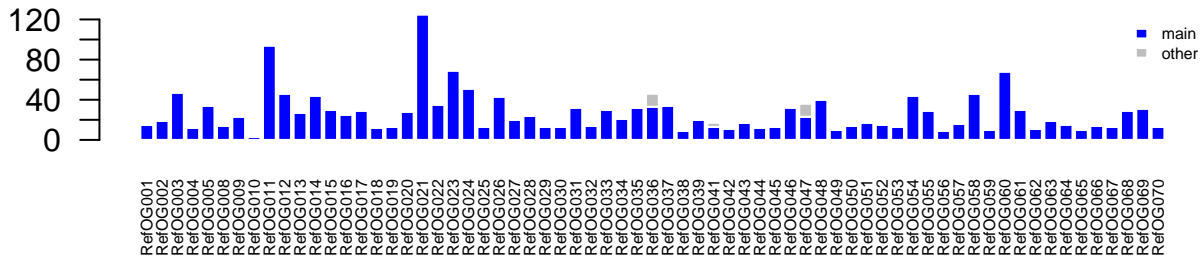
Index
n(>=0.95) = NA

size OGs



n(best = all) = 62 (p=0.726 | r=1.000)
n(best < all) = 5 (p=1.000 | r=0.795)

number of reference genes in main and other OGs



num OGs per refOG

