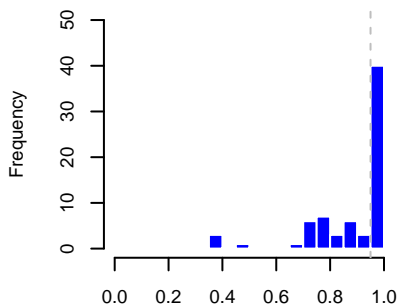
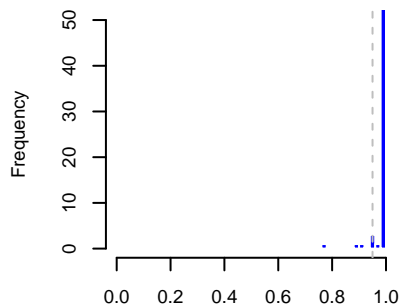
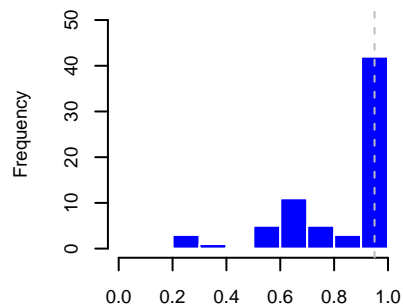


F-score

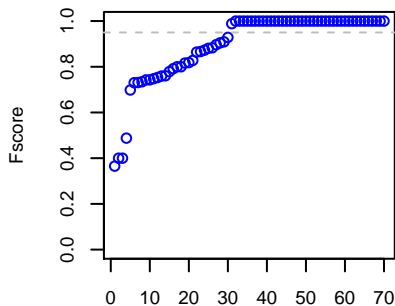
F-score
av = 0.876 (inc = 0.994)

Precision

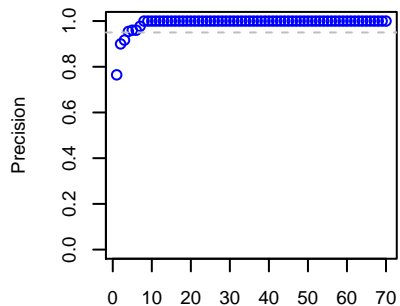
Precision
av = 0.990 (inc = 0.991)

Recall

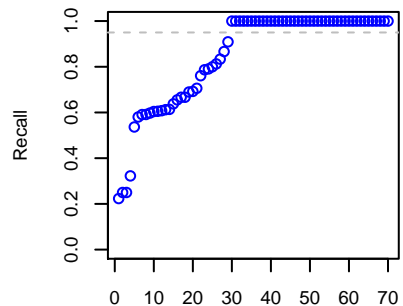
Recall
av = 0.806 (inc = 0.997)

Fscore

Index
n(>=.95) = 40

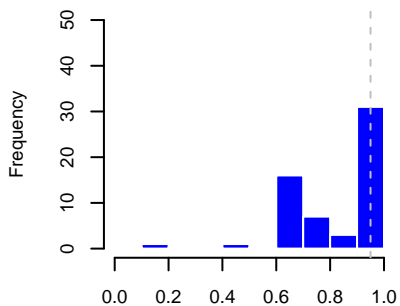
Precision

Index
n(>=.95) = 67

Recall

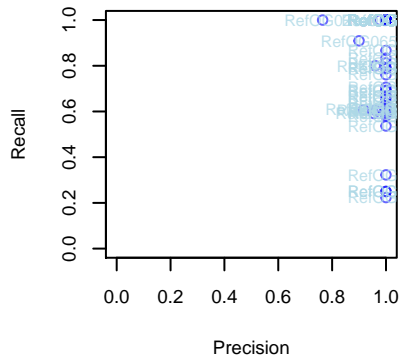
Index
n(>=.95) = 41

Rand index

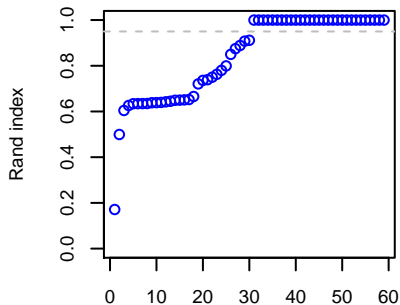


av = 0.833 (inc = 0.833) | med = 0.912

Precision & recall

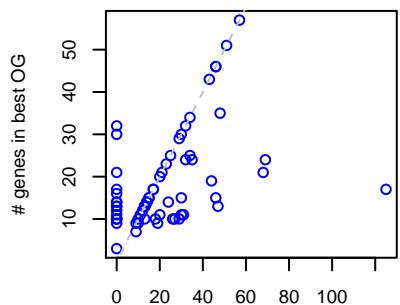


Rand index



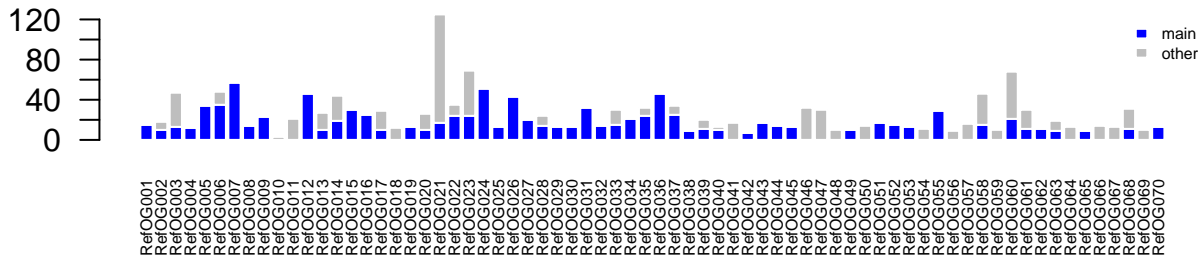
n(>=0.95) = NA

size OGs



n(best = all) = 48 (p=0.990 | r=0.930)
n(best = all) = 31 (p=0.992 | r=1.000)
n(best < all) = 23 (p=0.987 | r=0.675)

number of reference genes in main and other OGs



num OGs per refOG

