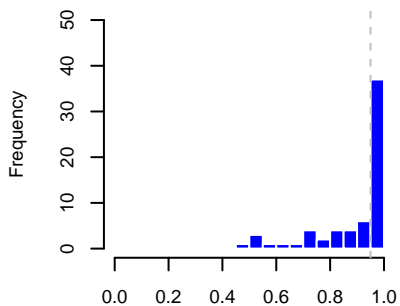
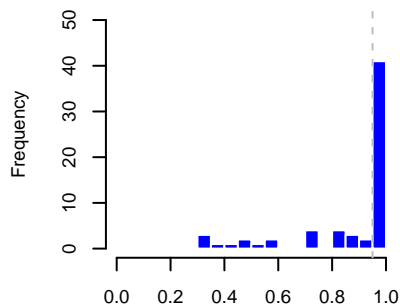
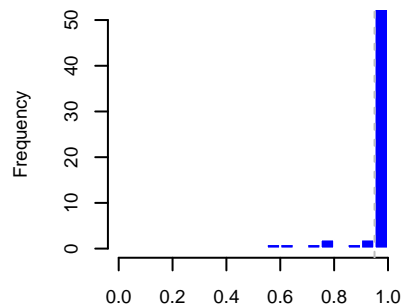


F-score

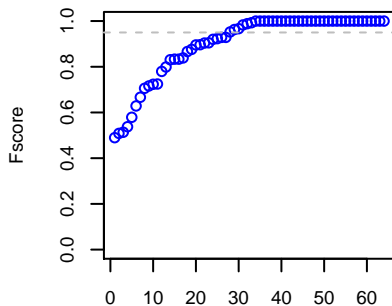
F-score
av = 0.928 (inc = 0.939)

Precision

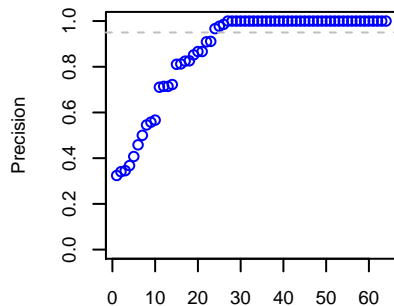
Precision
av = 0.905 (inc = 0.902)

Recall

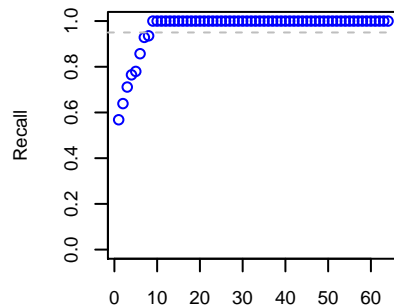
Recall
av = 0.972 (inc = 0.996)

Fscore

Index
n(>=.95) = 37

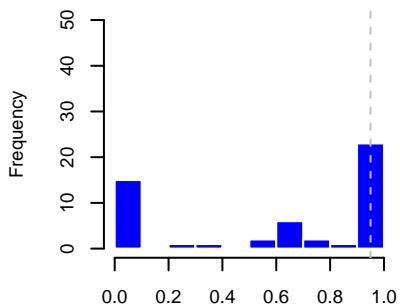
Precision

Index
n(>=.95) = 41

Recall

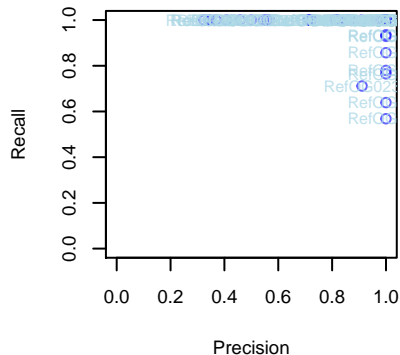
Index
n(>=.95) = 56

Rand index

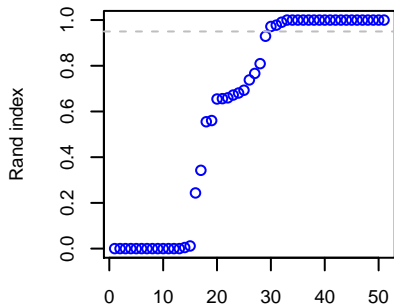


av = 0.718 (inc = 0.718) | med = 0.738

Precision & recall

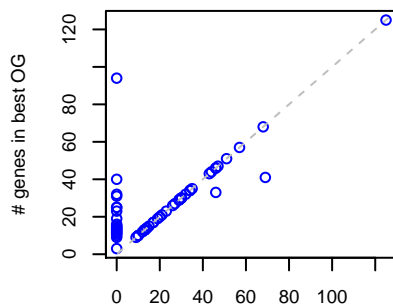


Rand index



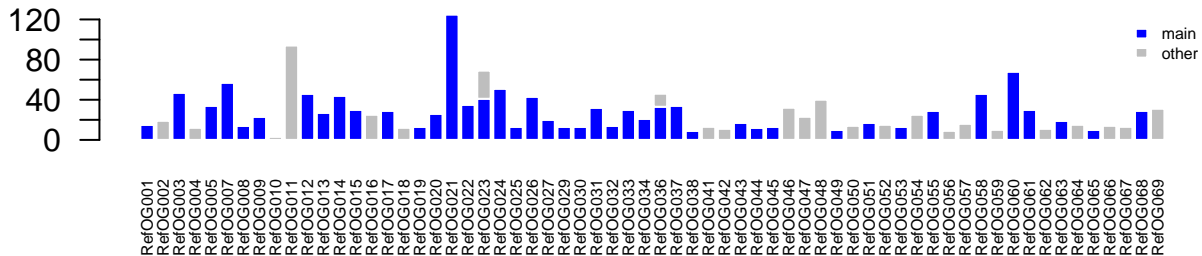
n(>=0.95) = NA

size OGs



n(best = all) = 40 (p=0.893 | r=0.993)
n(best < all) = 2 (p=0.956 | r=0.746)

number of reference genes in main and other OGs



num OGs per refOG

