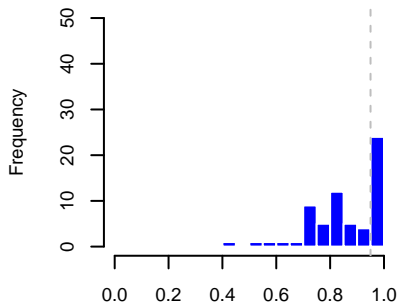
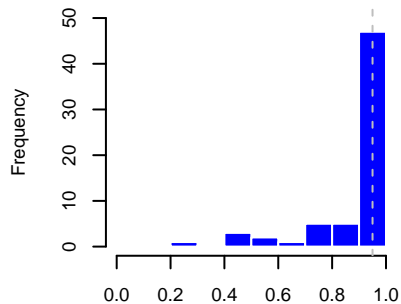
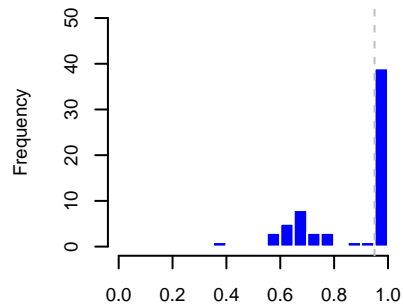


F-score

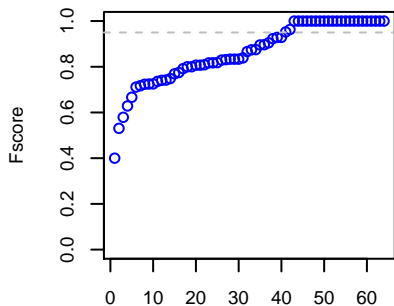
F-score
av = 0.829 (inc = 0.941)

Precision

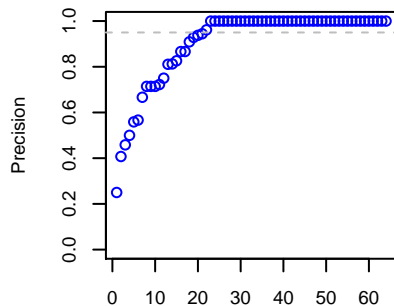
Precision
av = 0.911 (inc = 0.911)

Recall

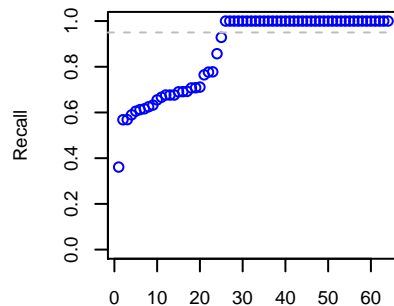
Recall
av = 0.799 (inc = 0.988)

Fscore

Index
n($\geq .95$) = 24

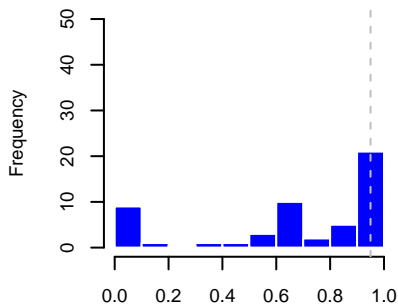
Precision

Index
n($\geq .95$) = 43

Recall

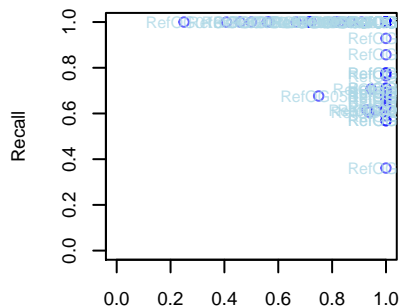
Index
n($\geq .95$) = 39

Rand index



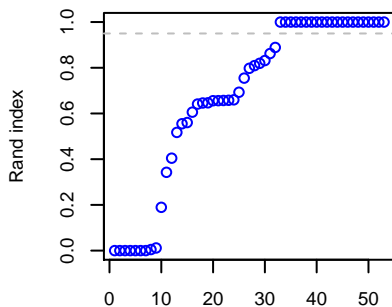
av = 0.695 (inc = 0.695) | med = 0.798

Precision & recall



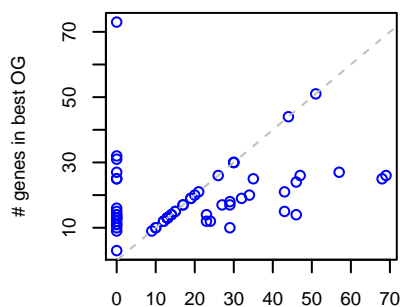
Precision

Rand index



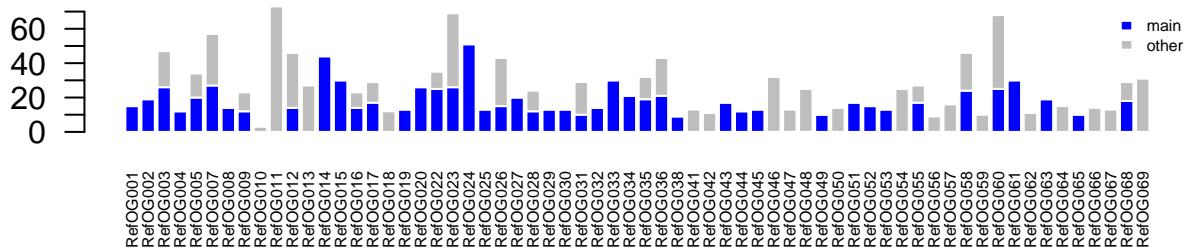
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 27 (p=0.861 | r=0.992)
n(best < all) = 18 (p=0.973 | r=0.664)

number of reference genes in main and other OGs



num OGs per refOG

