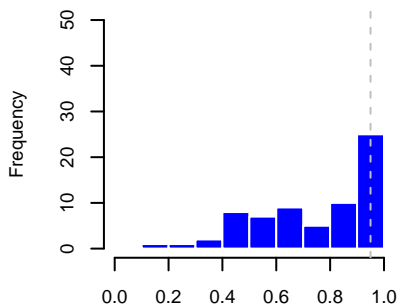
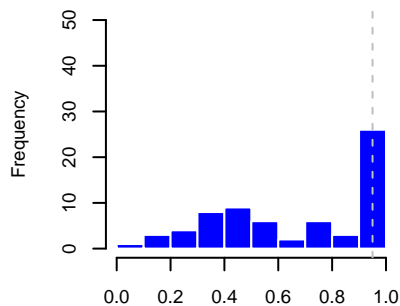
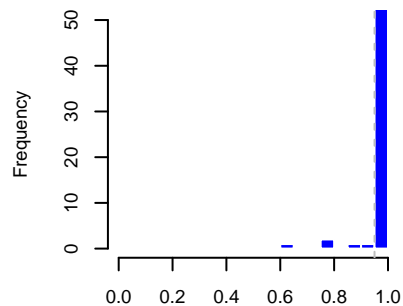


F-score

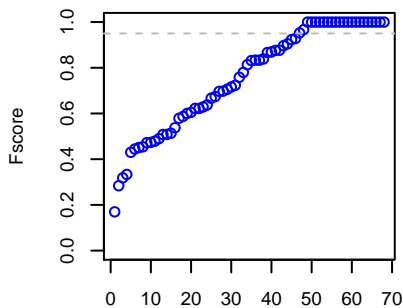
F-score
av = 0.829 (inc = 0.831)

Precision

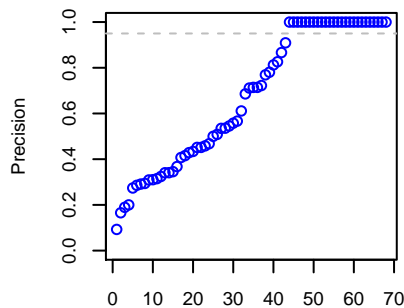
Precision
av = 0.737 (inc = 0.729)

Recall

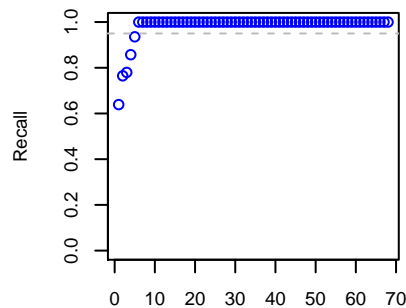
Recall
av = 0.984 (inc = 0.996)

Fscore

Index
n(>=.95) = 22

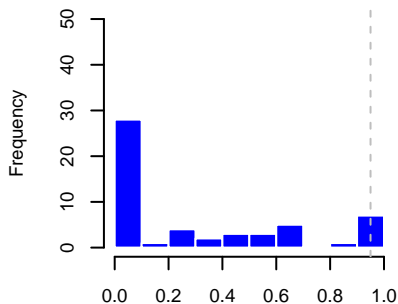
Precision

Index
n(>=.95) = 25

Recall

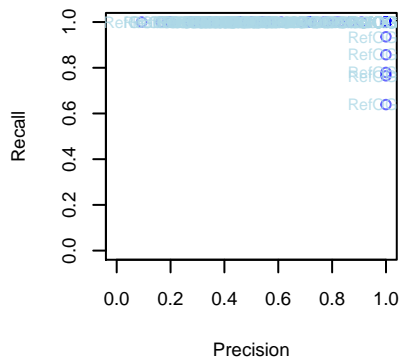
Index
n(>=.95) = 63

Rand index

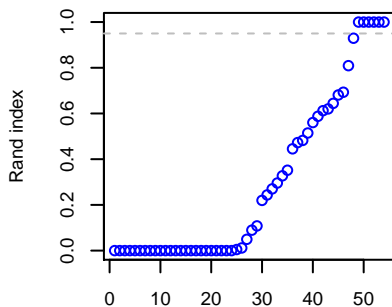


Rand index
av = 0.373 (inc = 0.373) | med = 0.069

Precision & recall

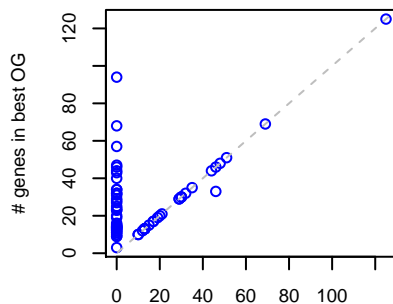


Rand index



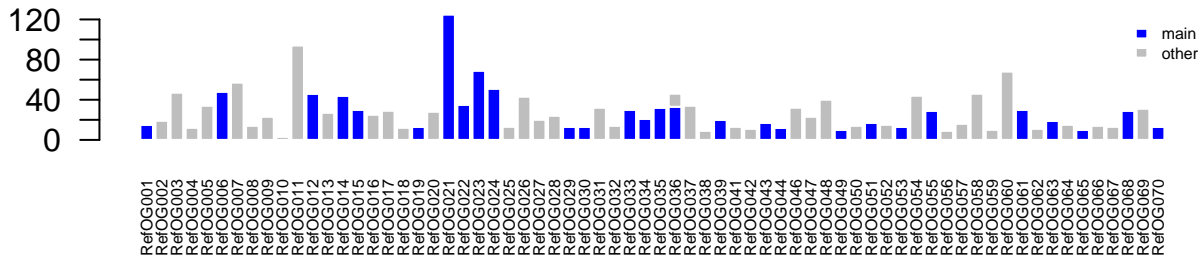
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 27 (p=0.773 | r=0.992)
n(best < all) = 1 (p=1.000 | r=0.780)

number of reference genes in main and other OGs



num OGs per refOG

