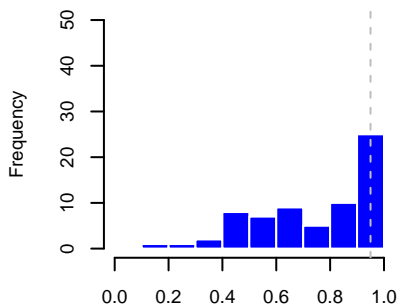
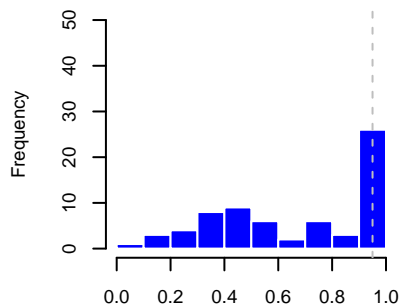
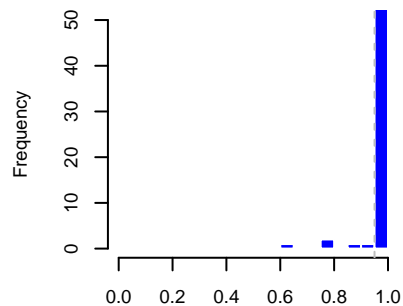


F-score

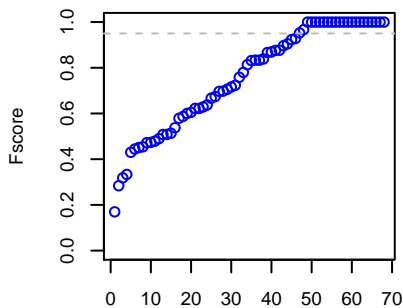
F-score
av = 0.736 (inc = 0.737)

Precision

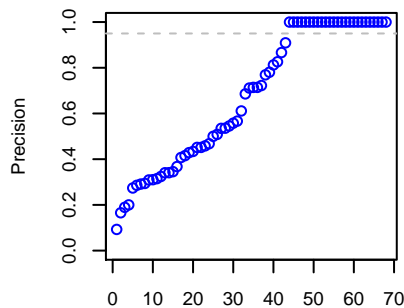
Precision
av = 0.647 (inc = 0.644)

Recall

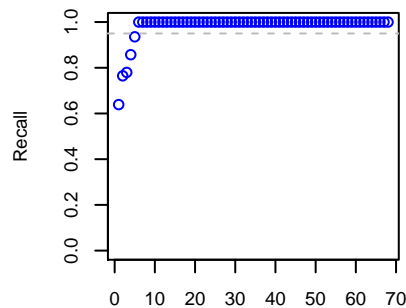
Recall
av = 0.984 (inc = 0.989)

Fscore

Index
n(>=.95) = 22

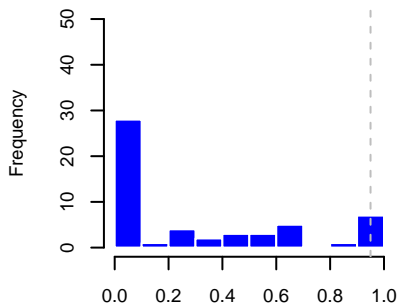
Precision

Index
n(>=.95) = 25

Recall

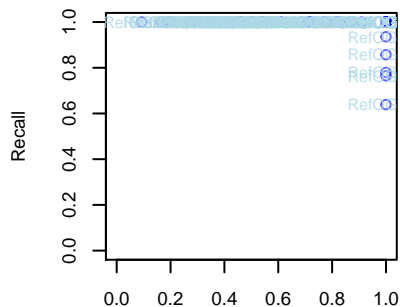
Index
n(>=.95) = 63

Rand index



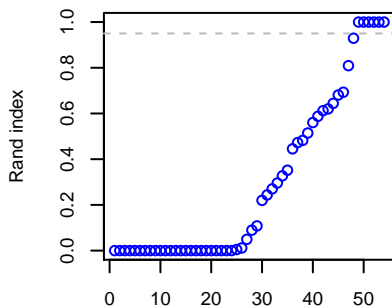
Rand index
av = NaN (inc = NaN) | med = 0.069

Precision & recall



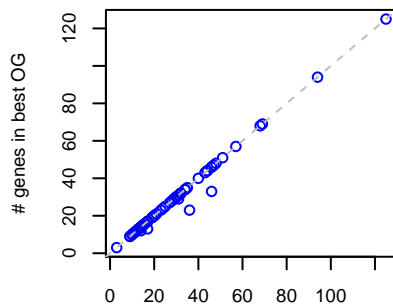
Precision

Rand index



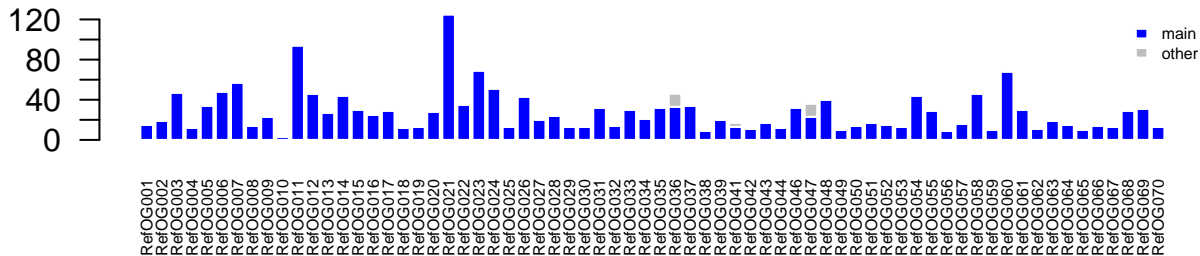
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 63 (p=0.950 | r=0.999)
n(best < all) = 5 (p=1.000 | r=0.795)

number of reference genes in main and other OGs



num OGs per refOG

