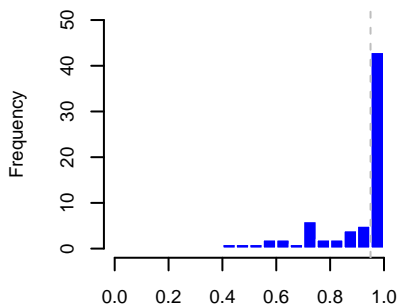
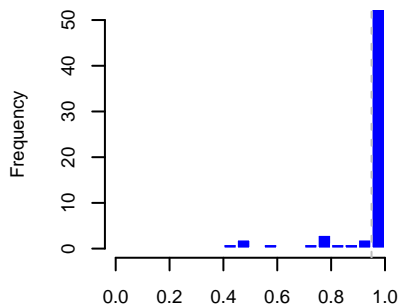
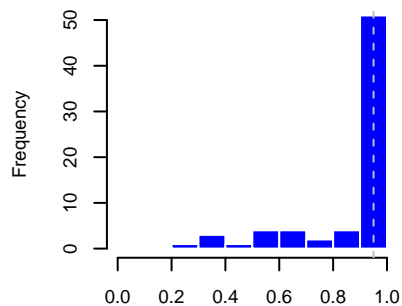


F-score

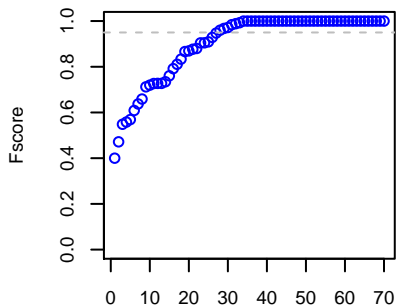
F-score
av = 0.864 (inc = 0.905)

Precision

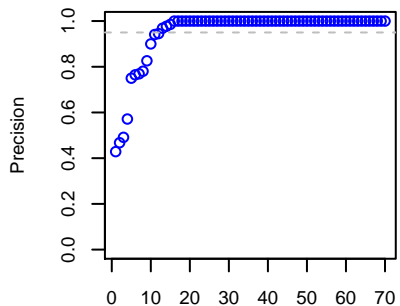
Precision
av = 0.952 (inc = 0.953)

Recall

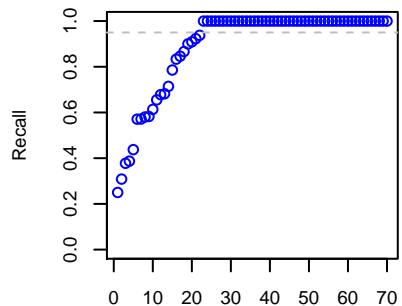
Recall
av = 0.841 (inc = 0.909)

Fscore

Index
n(>=.95) = 43

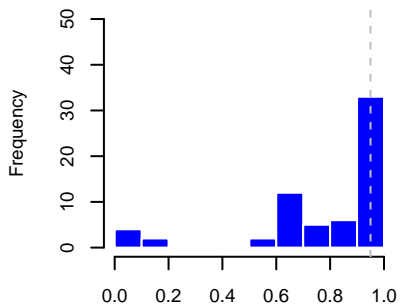
Precision

Index
n(>=.95) = 58

Recall

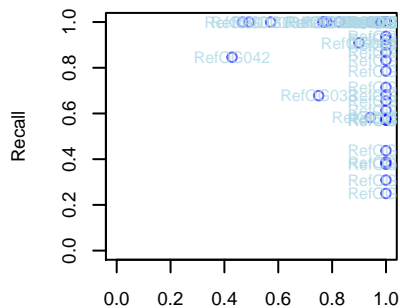
Index
n(>=.95) = 48

Rand index



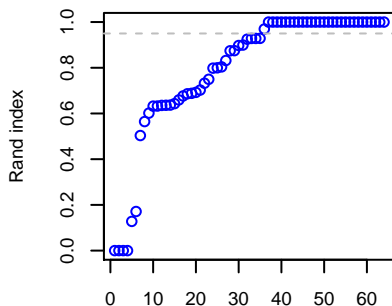
av = NaN (inc = NaN) | med = 0.927

Precision & recall



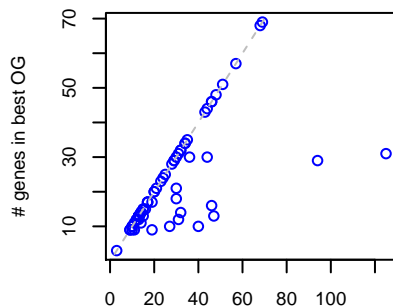
Precision

Rand index



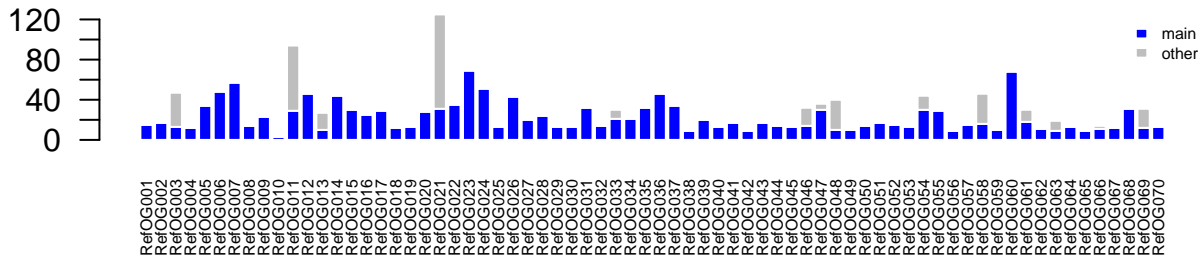
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 50 (p=0.951 | r=0.976)
n(best = all) = 50 (p=0.951 | r=0.979)
n(best < all) = 20 (p=0.951 | r=0.673)

number of reference genes in main and other OGs



num OGs per refOG

