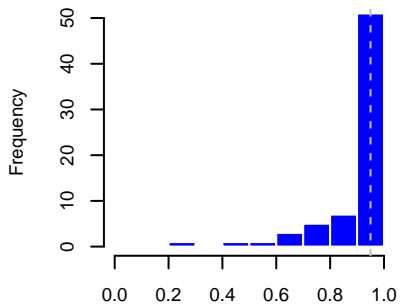
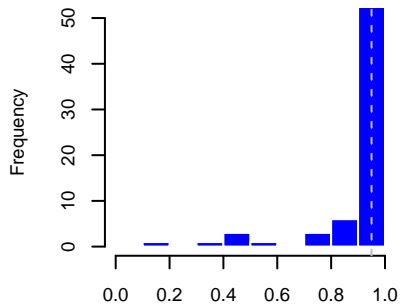
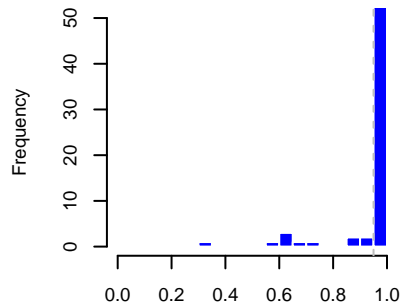


F-score

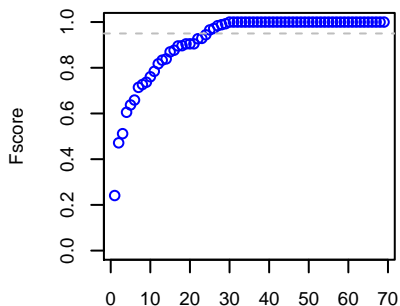
F-score
av = 0.891 (inc = 0.924)

Precision

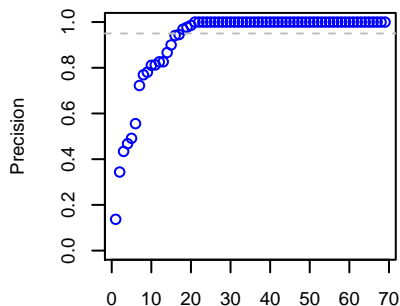
Precision
av = 0.934 (inc = 0.932)

Recall

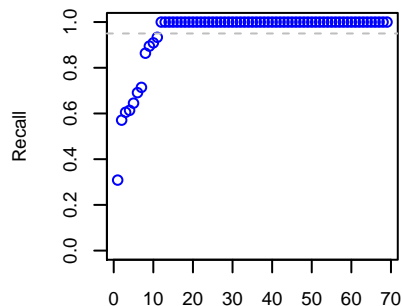
Recall
av = 0.901 (inc = 0.956)

Fscore

Index
n(>=.95) = 45

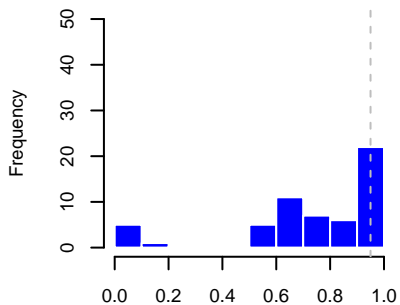
Precision

Index
n(>=.95) = 52

Recall

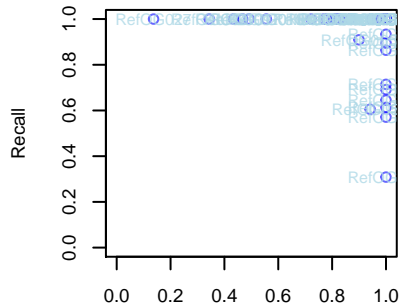
Index
n(>=.95) = 58

Rand index



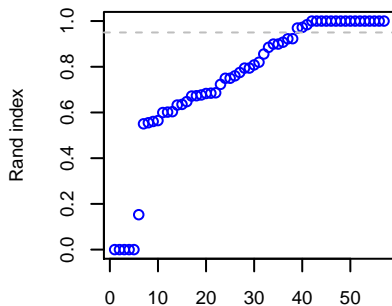
Rand index
av = NaN (inc = NaN) | med = 0.794

Precision & recall



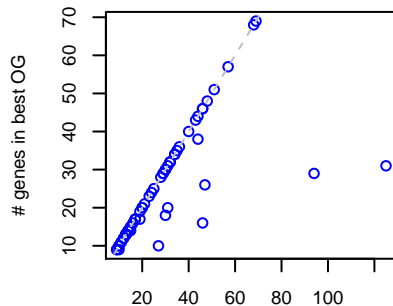
Precision

Rand index



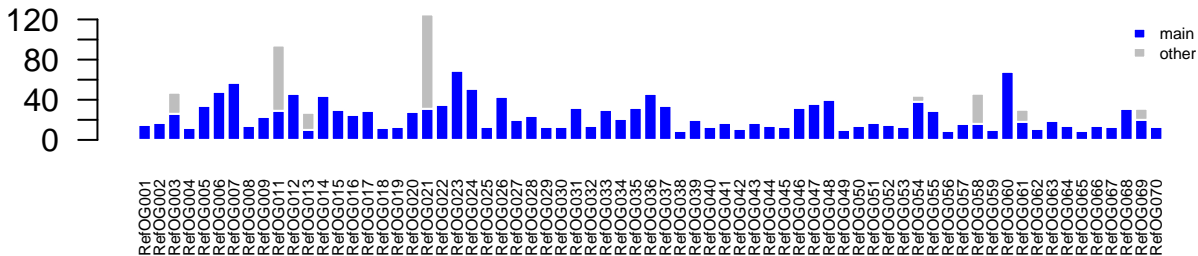
Index
n(>=0.95) = NA

size OGs



genes in ref OG
n(best < all) = 58 (p=0.909 | r=1.000)
n(best > all) = 11 (p=0.986 | r=0.705)

number of reference genes in main and other OGs



num OGs per refOG

