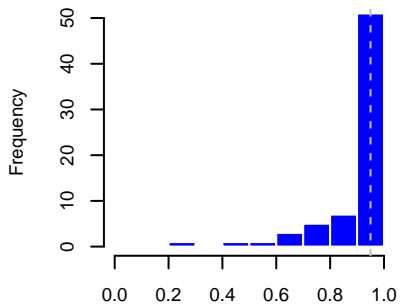
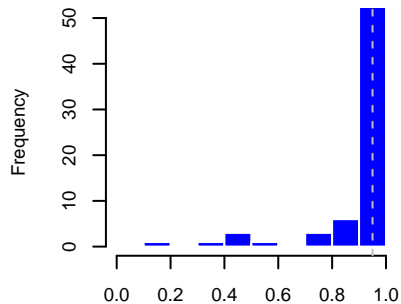
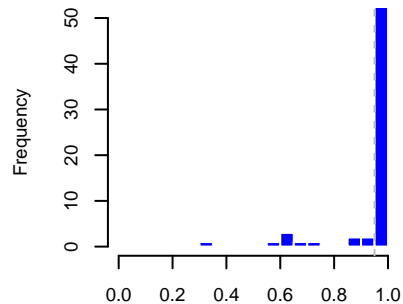


F-score

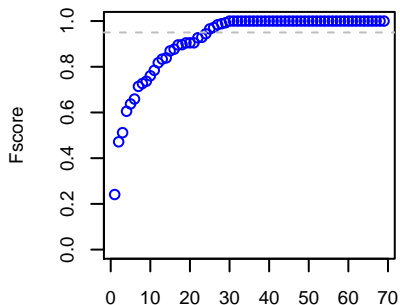
F-score
av = 0.930 (inc = 0.974)

Precision

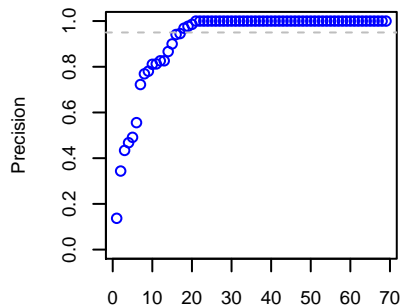
Precision
av = 0.958 (inc = 0.955)

Recall

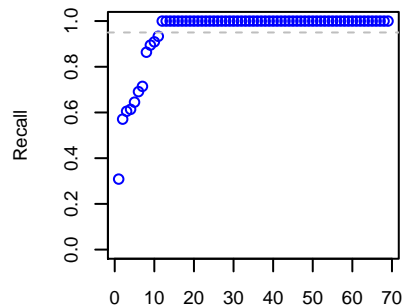
Recall
av = 0.924 (inc = 0.999)

Fscore

Index
n(>=.95) = 45

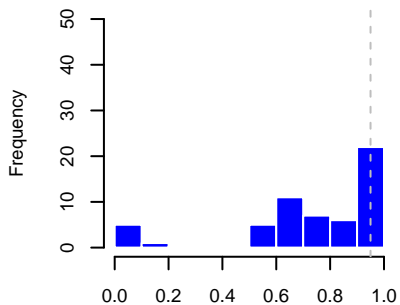
Precision

Index
n(>=.95) = 52

Recall

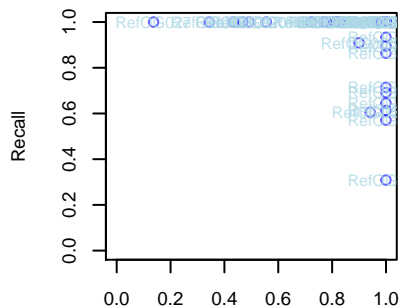
Index
n(>=.95) = 58

Rand index



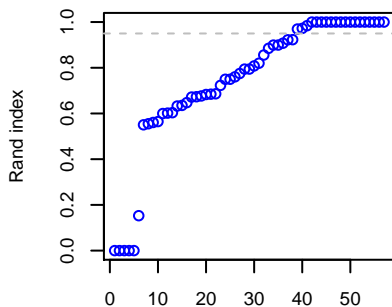
Rand index
av = 0.750 (inc = 0.750) | med = 0.794

Precision & recall



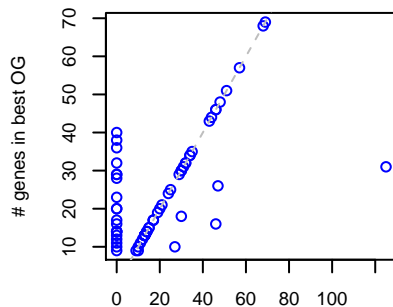
Precision

Rand index



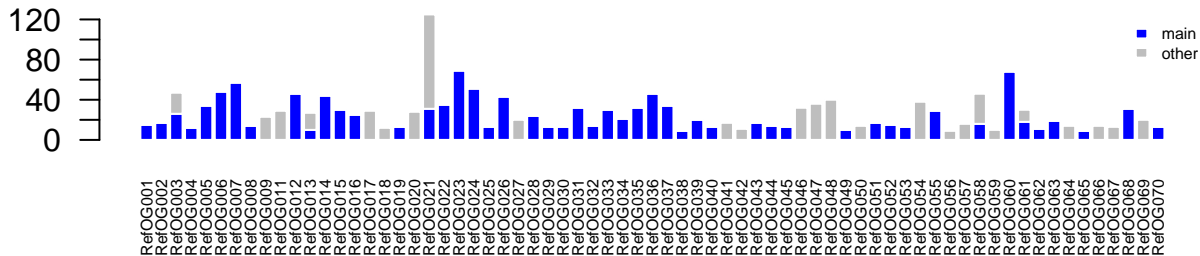
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 43 (p=0.950 | r=0.998)
n(best < all) = 6 (p=0.974 | r=0.684)

number of reference genes in main and other OGs



num OGs per refOG

