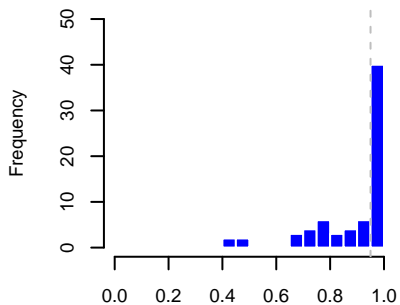
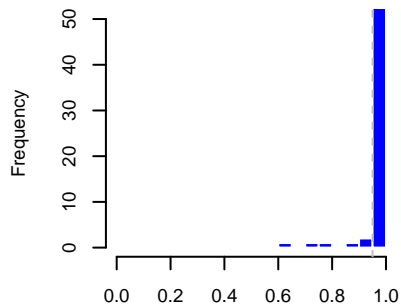
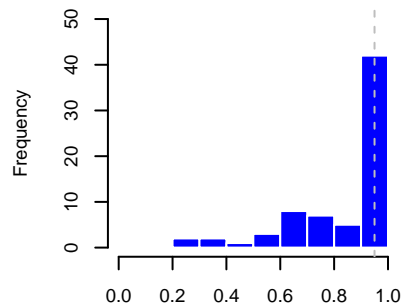


F-score

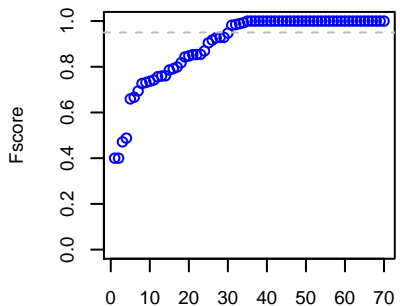
F-score
av = 0.897 (inc = 0.989)

Precision

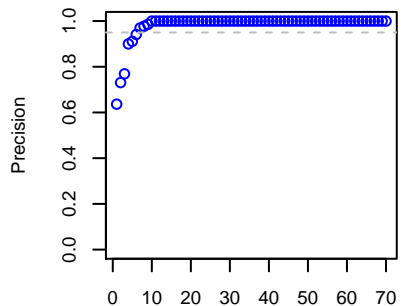
Precision
av = 0.980 (inc = 0.981)

Recall

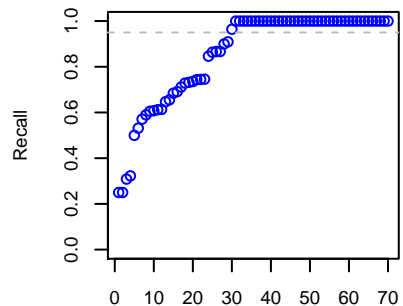
Recall
av = 0.843 (inc = 0.999)

Fscore

Index
n(>=.95) = 40

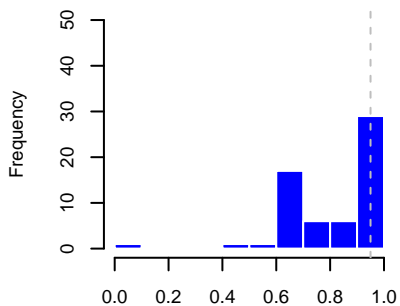
Precision

Index
n(>=.95) = 64

Recall

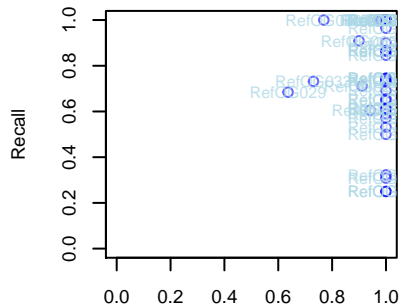
Index
n(>=.95) = 41

Rand index

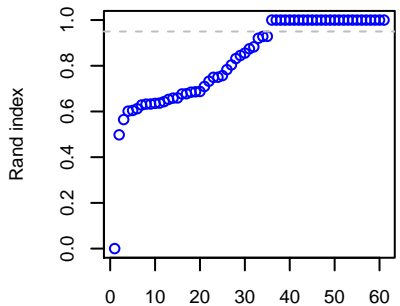


av = 0.785 (inc = 0.785) | med = 0.874

Precision & recall

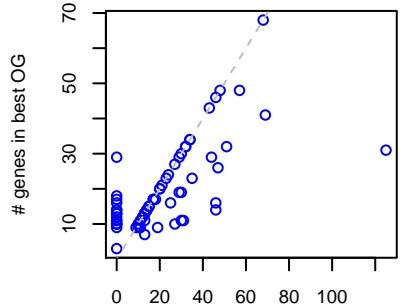


Rand index



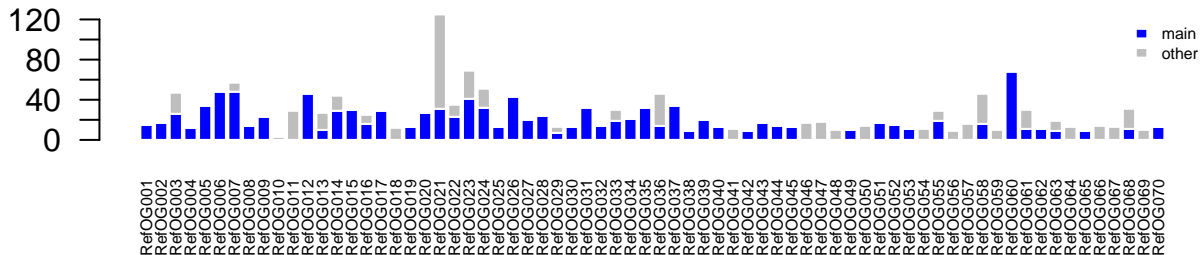
Index
n(>=0.95) = NA

size OGs



n(best = all) = 33 (p=0.991 | r=0.999)
n(best < all) = 21 (p=0.958 | r=0.722)

number of reference genes in main and other OGs



num OGs per refOG

