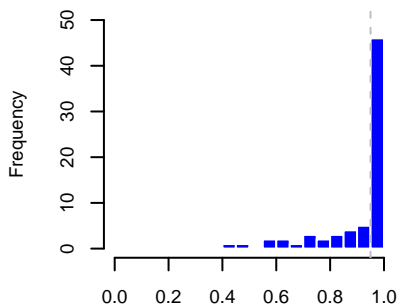
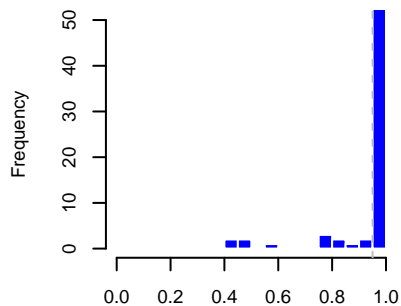
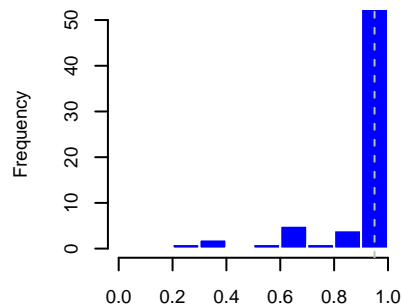


F-score

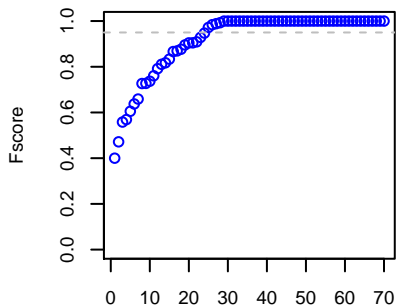
F-score
av = 0.929 (inc = 0.976)

Precision

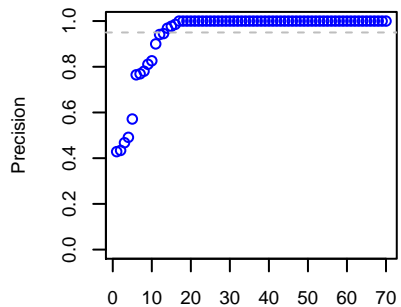
Precision
av = 0.962 (inc = 0.959)

Recall

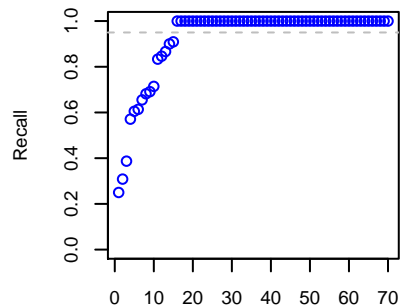
Recall
av = 0.920 (inc = 0.999)

Fscore

Index
n(>=.95) = 46

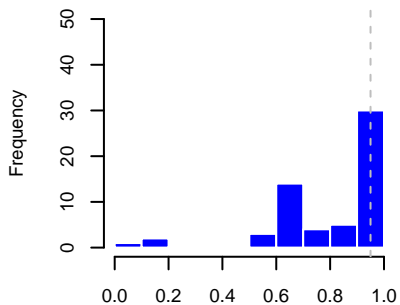
Precision

Index
n(>=.95) = 57

Recall

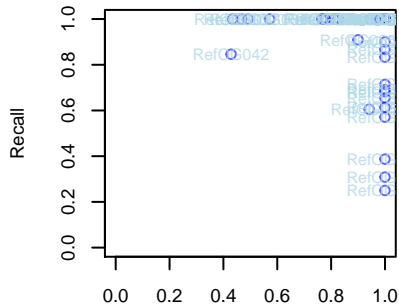
Index
n(>=.95) = 55

Rand index



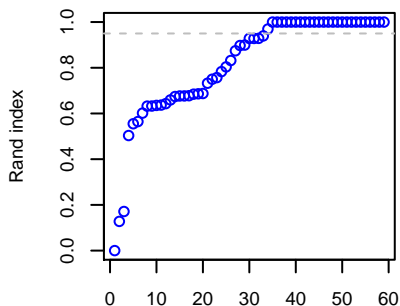
Rand index
av = 0.805 (inc = 0.805) | med = 0.928

Precision & recall



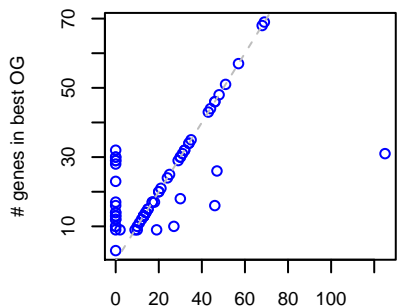
Precision

Rand index



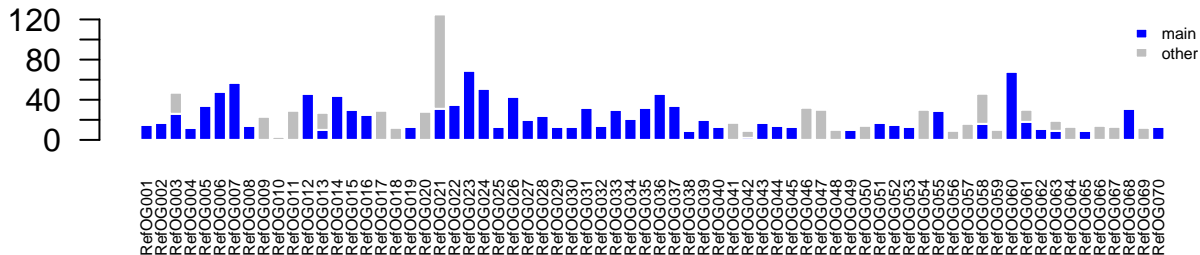
Index
n(≥ 0.95) = NA

size OGs



genes in best OG
n(best = all) = 42 (p=0.962 | r=1.000)
n(best < all) = 8 (p=0.980 | r=0.707)

number of reference genes in main and other OGs



num OGs per refOG

