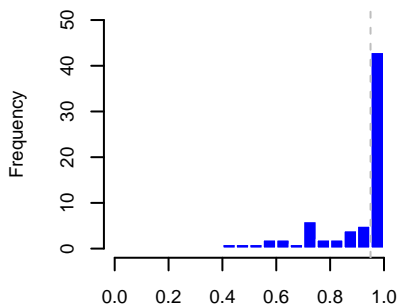
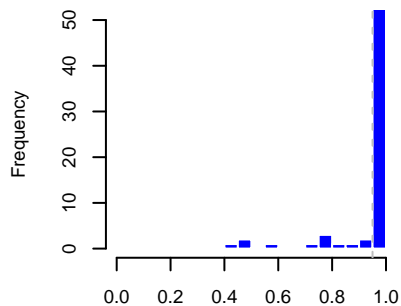
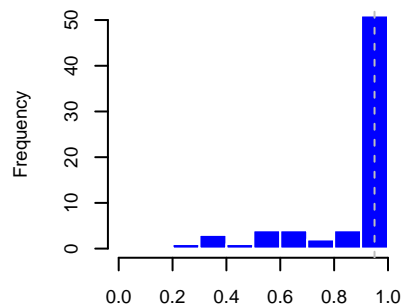


F-score

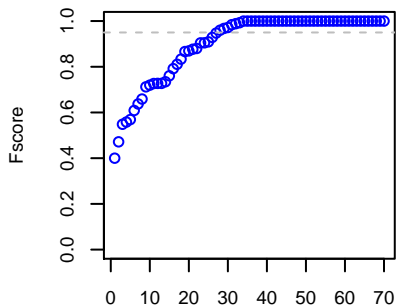
F-score
av = 0.921 (inc = 0.976)

Precision

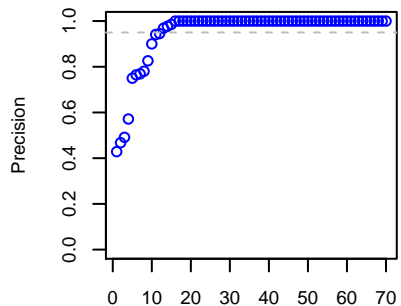
Precision
av = 0.959 (inc = 0.959)

Recall

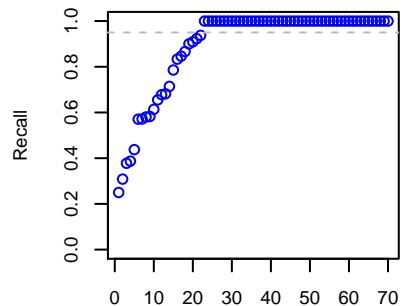
Recall
av = 0.908 (inc = 0.999)

Fscore

Index
n($\geq .95$) = 43

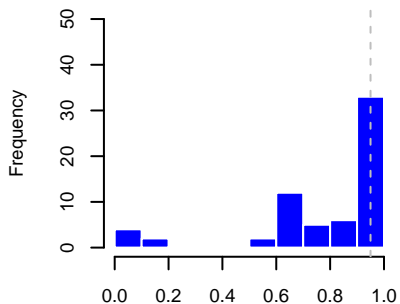
Precision

Index
n($\geq .95$) = 58

Recall

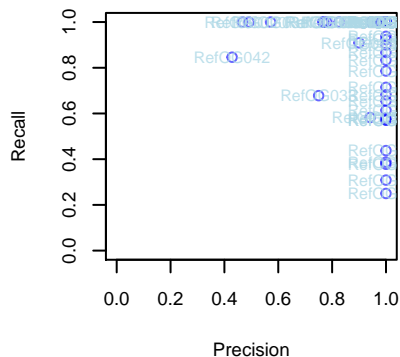
Index
n($\geq .95$) = 48

Rand index

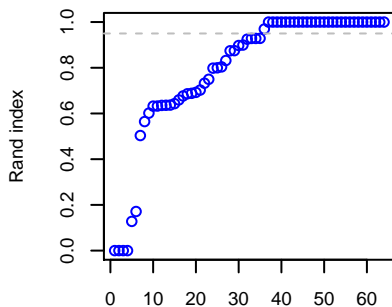


Rand index
av = 0.736 (inc = 0.736) | med = 0.927

Precision & recall

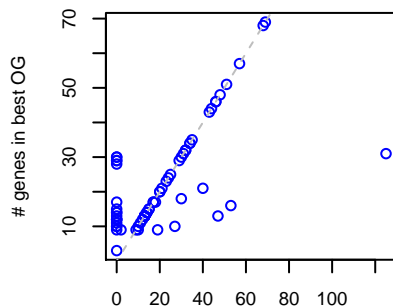


Rand index



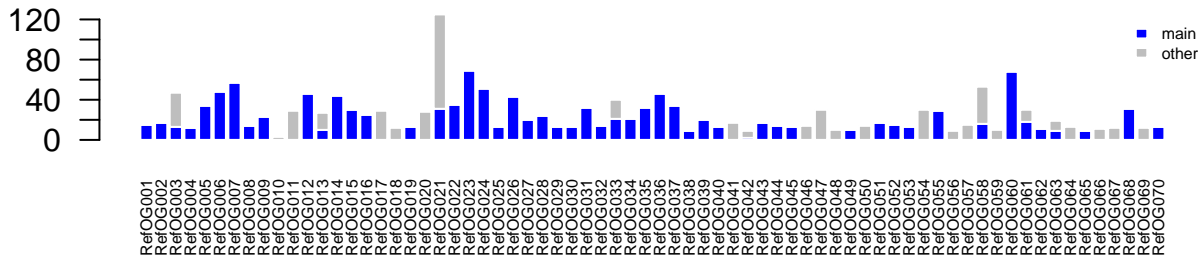
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 42 (p=0.966 | r=1.000)
n(best < all) = 9 (p=0.955 | r=0.689)

number of reference genes in main and other OGs



num OGs per refOG

