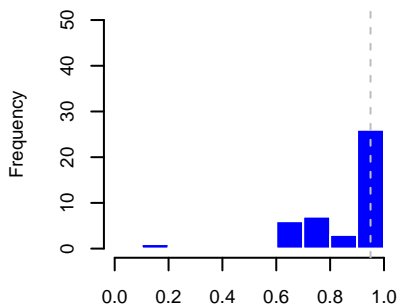
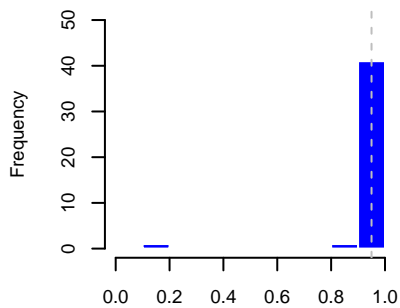
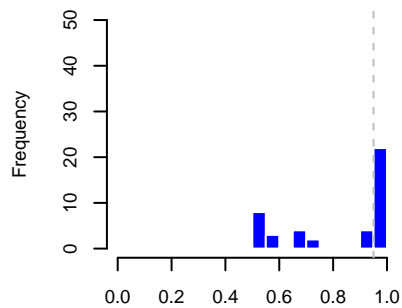


F-score

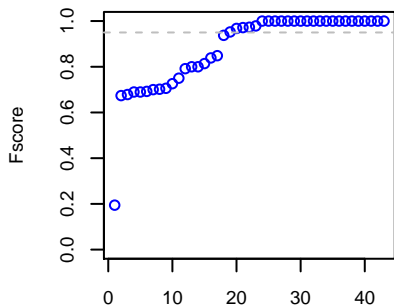
F-score
av = 0.849 (inc = 0.993)

Precision

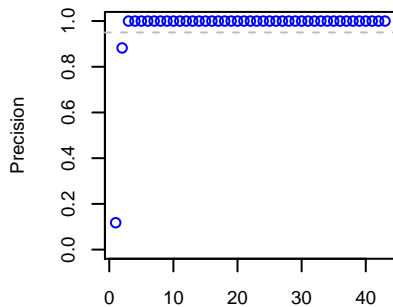
Precision
av = 0.986 (inc = 0.989)

Recall

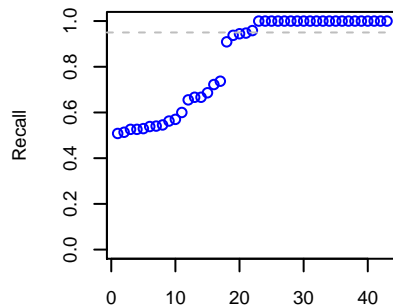
Recall
av = 0.776 (inc = 1.000)

Fscore

Index
n(>=.95) = 25

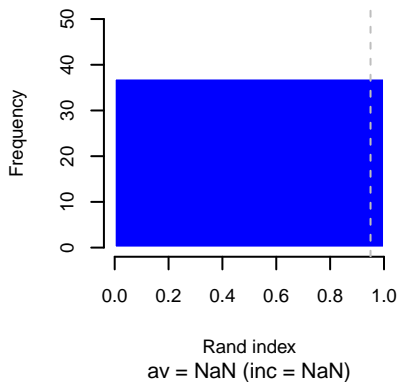
Precision

Index
n(>=.95) = 41

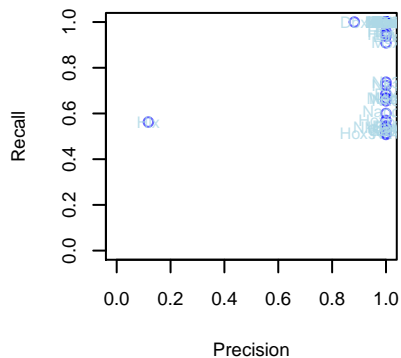
Recall

Index
n(>=.95) = 22

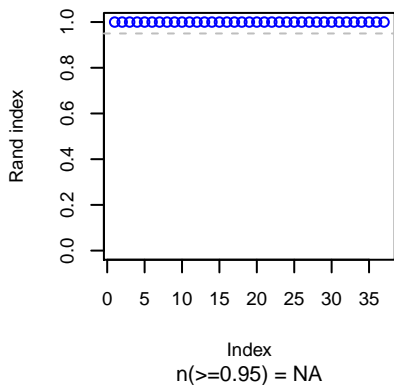
Rand index



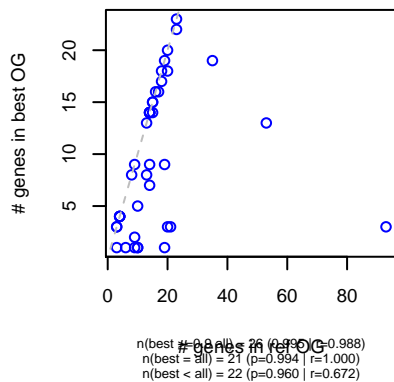
Precision & recall



Rand index

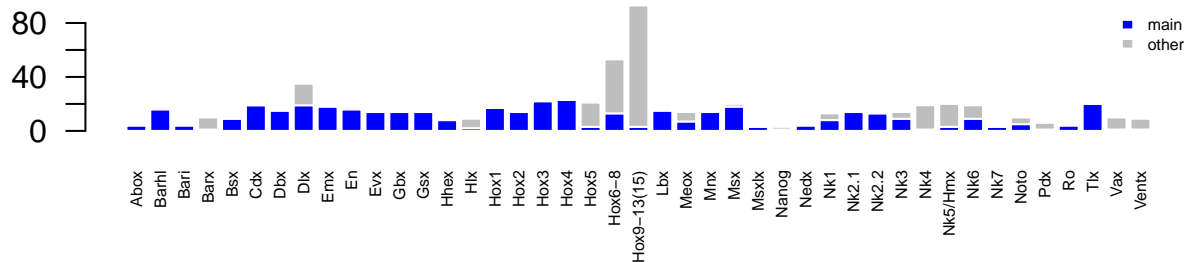


size OGs



$n(\text{best} = \text{all}) = 26$ ($p=0.990$ | $r=0.988$)
 $n(\text{best} < \text{all}) = 21$ ($p=0.994$ | $r=1.000$)
 $n(\text{best} > \text{all}) = 22$ ($p=0.960$ | $r=0.672$)

number of reference genes in main and other OGs



num OGs per refOG

