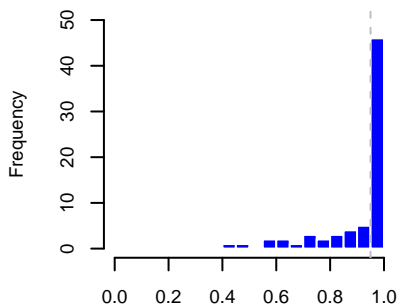
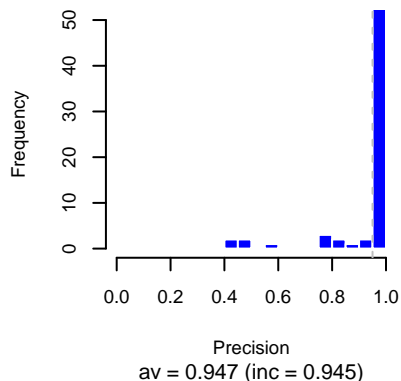
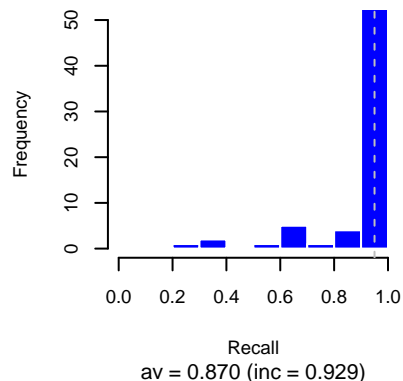
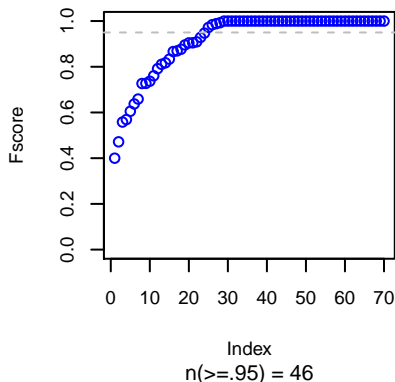
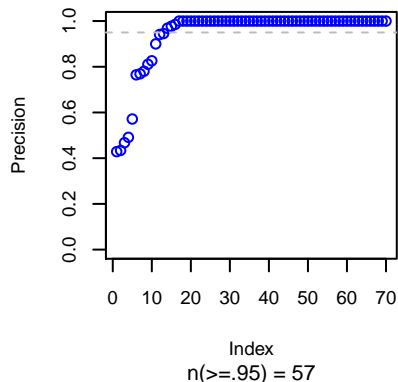
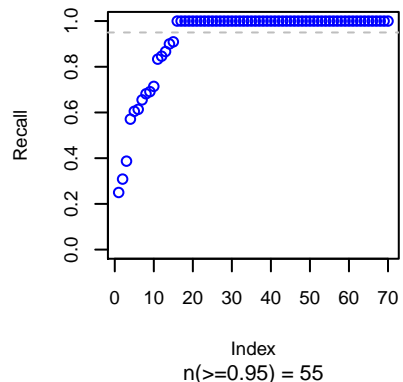
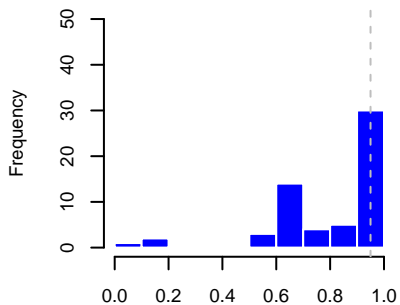


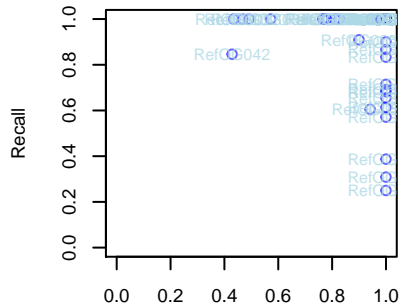
F-score**Precision****Recall****Fscore****Precision****Recall**

Rand index



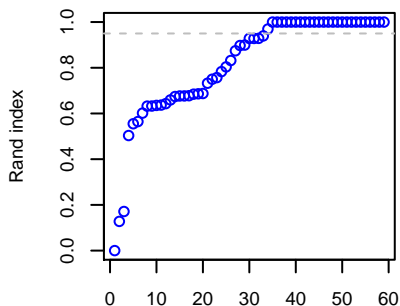
av = NaN (inc = NaN) | med = 0.928

Precision & recall



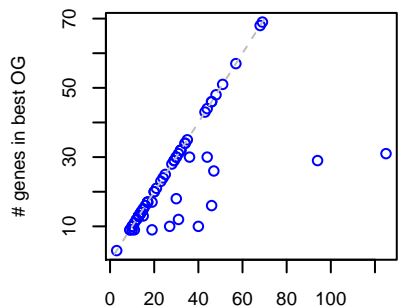
Precision

Rand index



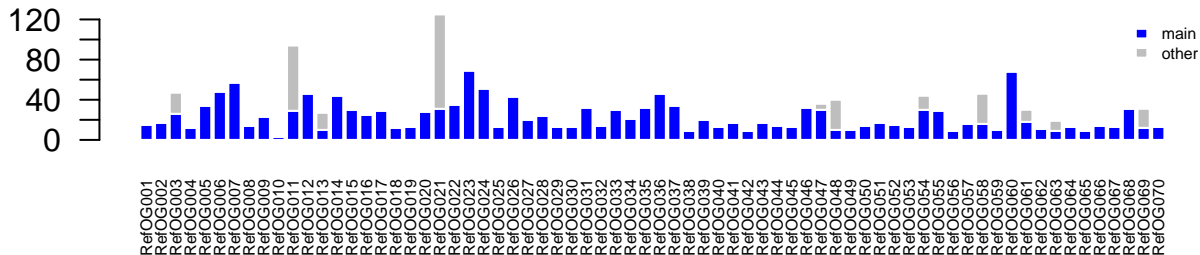
Index
n(>=0.95) = NA

size OGs



genes in best OG
n(best = all) = 55 (p=0.942 | r=1.000)
n(best < all) = 15 (p=0.951 | r=0.656)

number of reference genes in main and other OGs



num OGs per refOG

