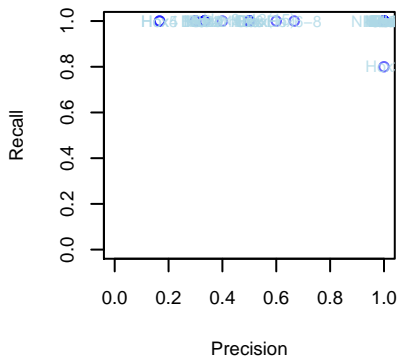
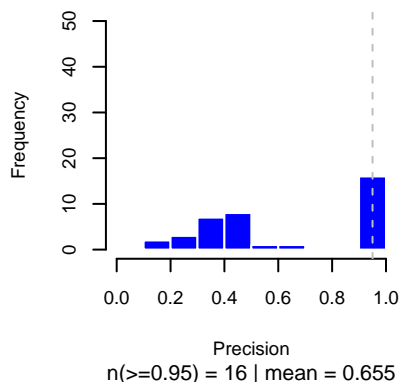


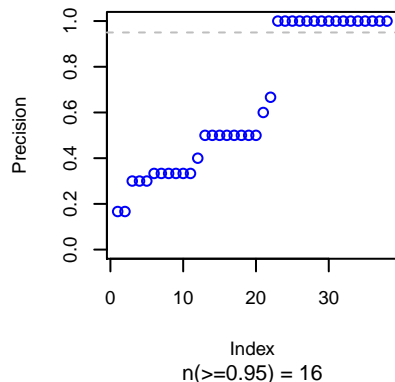
Precision & recall



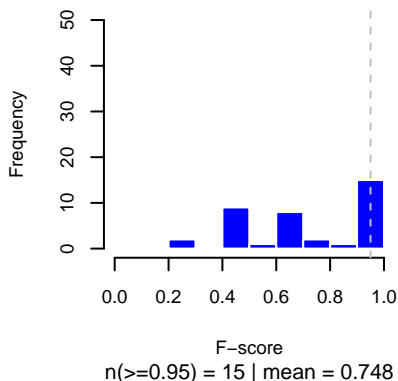
Precision



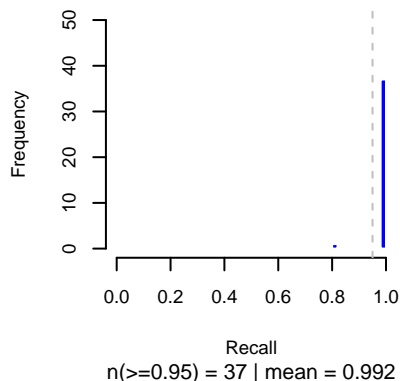
Precision



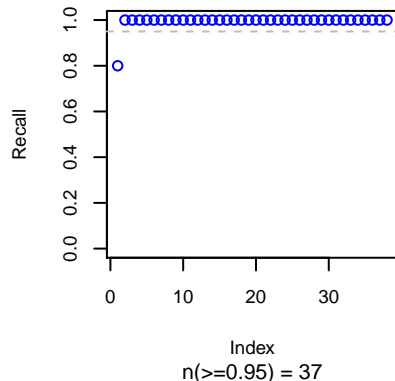
F-score



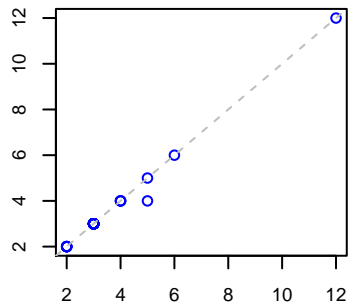
Recall



Recall

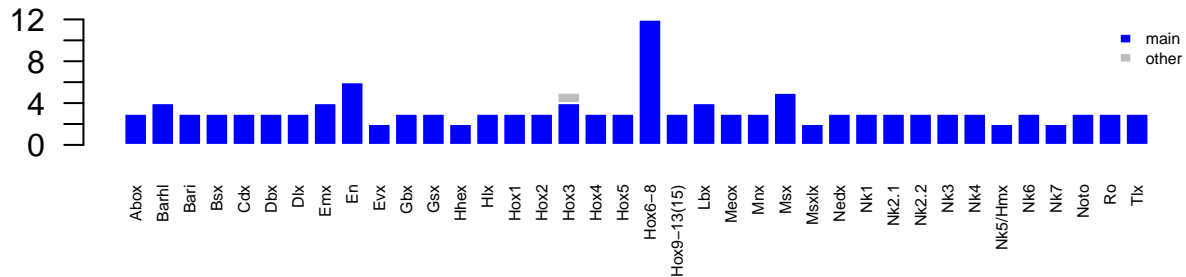


genes in best OG



n(best = all) = 27 (p=1.000 | r=1.000)
n(best = all) = 37 (p=0.646 | r=1.000)
n(best < all) = 1 (p=1.000 | r=0.800)

number of reference genes in main and other OGs



num OGs per refOG

