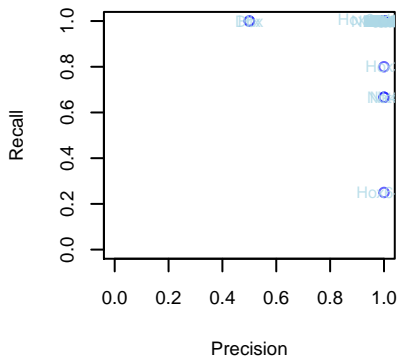
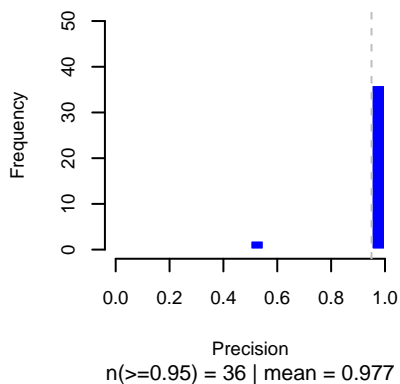


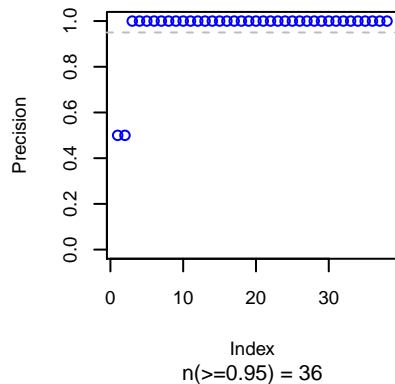
Precision & recall



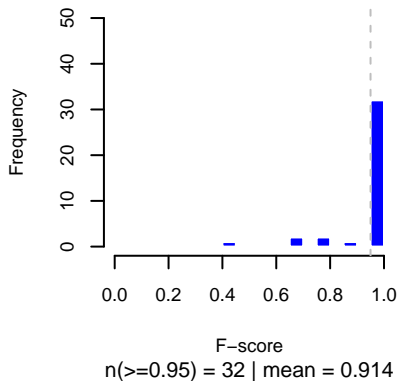
Precision



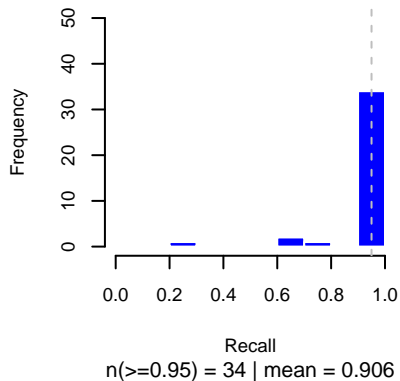
Precision



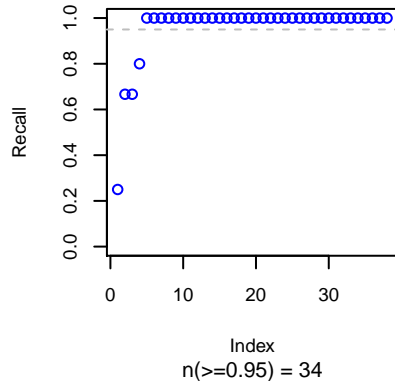
F-score



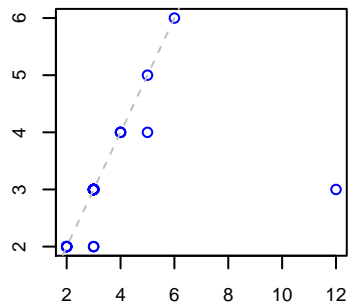
Recall



Recall

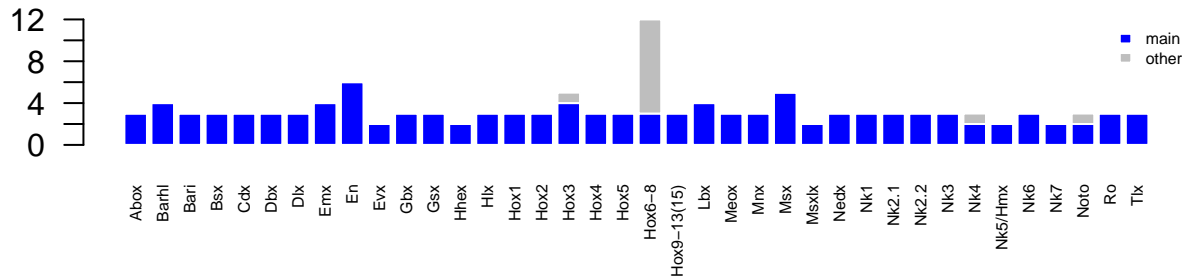


genes in best OG



n(best = 2) = 2 (p=1.000 | r=1.000)
n(best = all) = 34 (p=0.971 | r=1.000)
n(best < all) = 4 (p=1.000 | r=0.596)

number of reference genes in main and other OGs



num OGs per refOG

