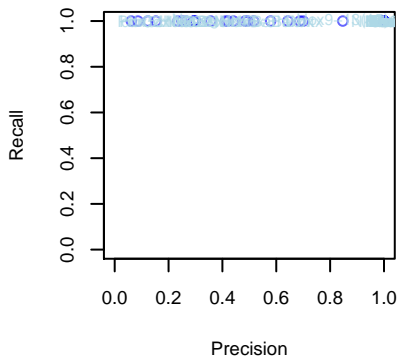
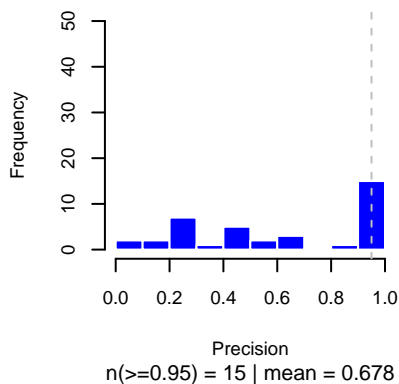


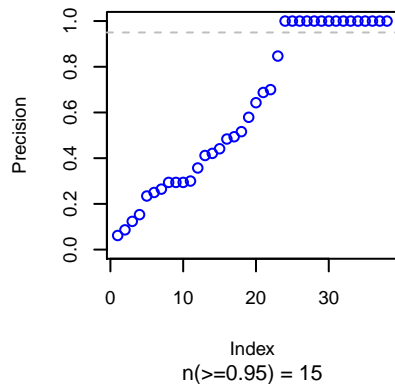
Precision & recall



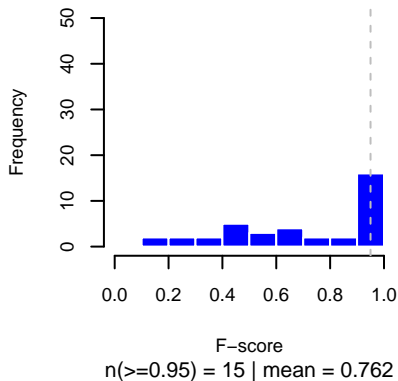
Precision



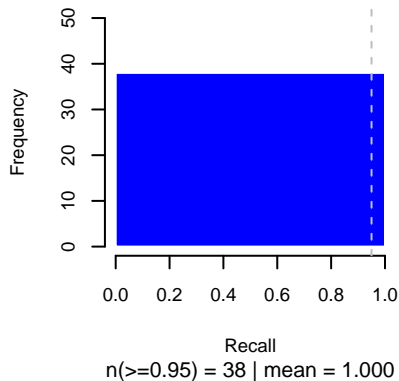
Precision



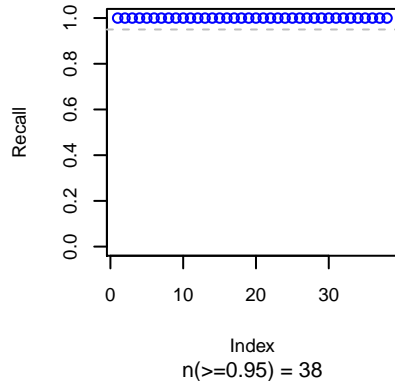
F-score



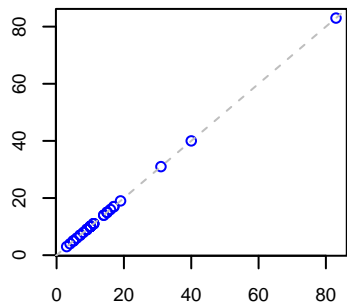
Recall



Recall

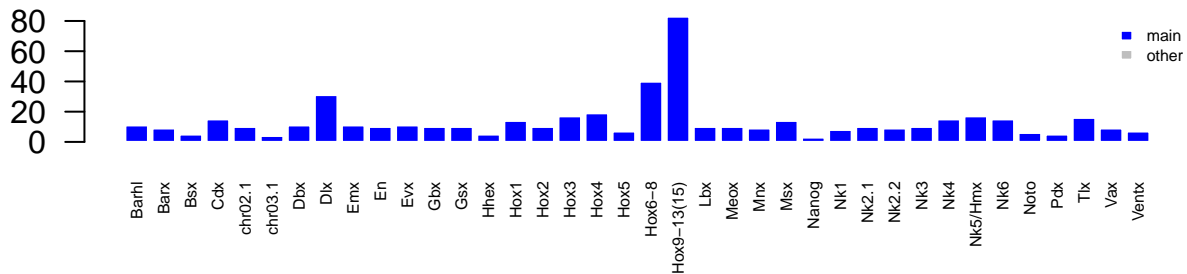


genes in best OG



n(best = all) = 28 (p=0.330 | r=1.000)
n(best = all) = 38 (p=0.630 | r=1.000)
n(best < all) = 0 (p=NaN | r=NaN)

number of reference genes in main and other OGs



num OGs per refOG

