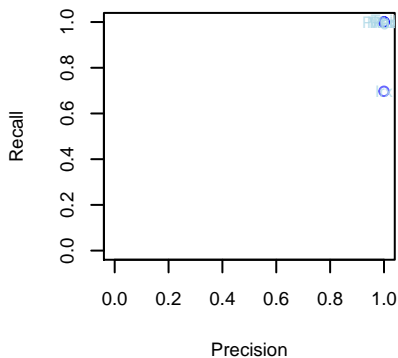
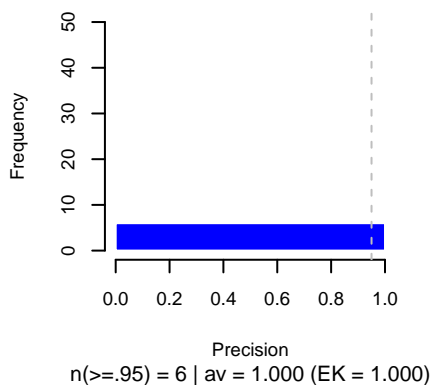


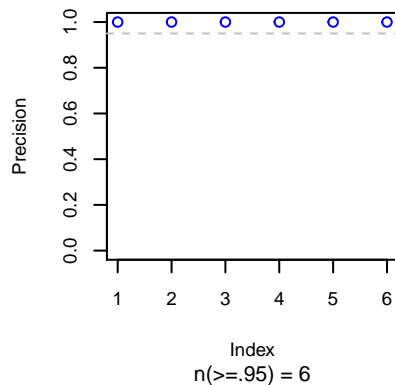
Precision & recall



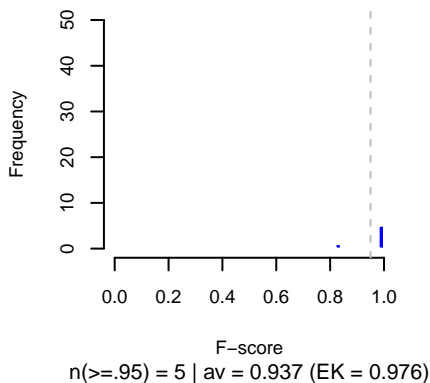
Precision



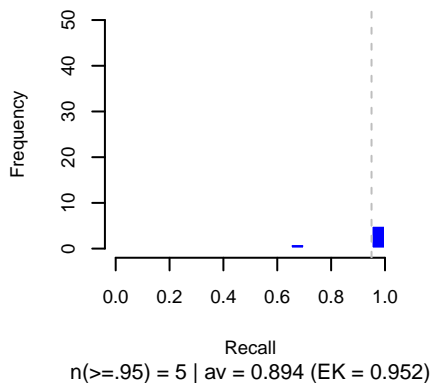
Precision



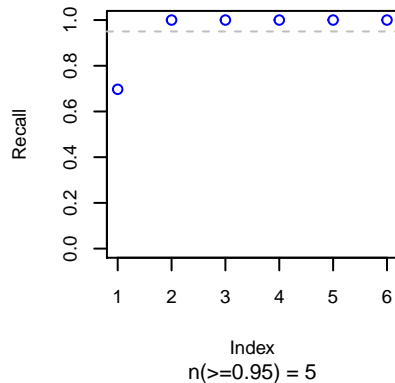
F-score



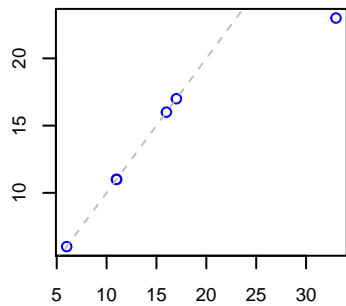
Recall



Recall

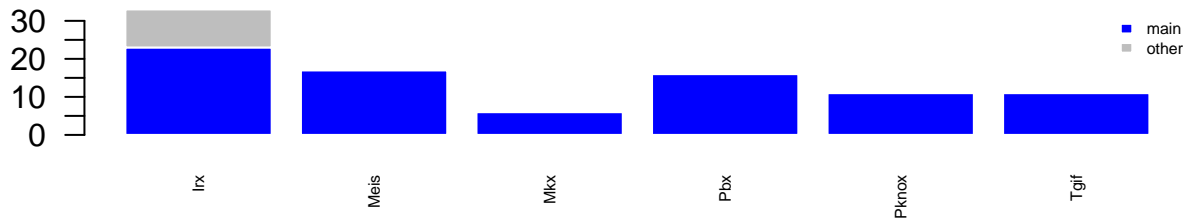


genes in best OG



n(best < all) = 5 (p=1.000 | r=1.000)
n(best = all) = 5 (p=1.000 | r=1.000)
n(best > all) = 1 (p=1.000 | r=0.697)

number of reference genes in main and other OGs



num OGs per refOG

