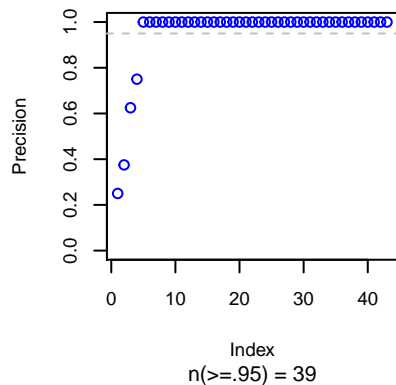
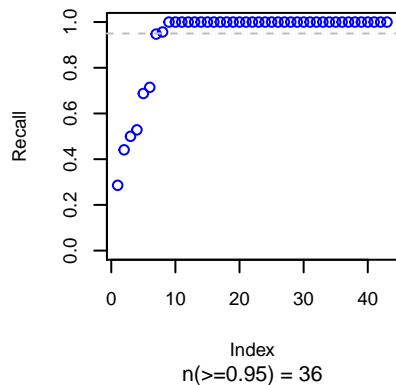


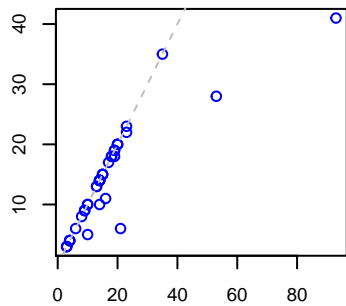
## Precision



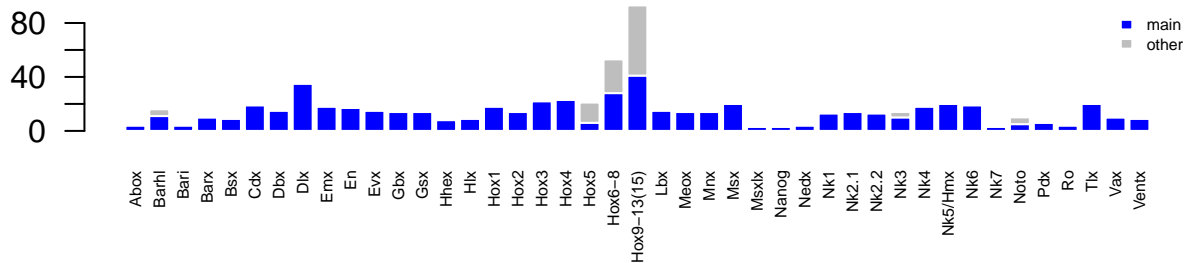
## Recall



# genes in best OG



## number of reference genes in main and other OGs



## num OGs per refOG

