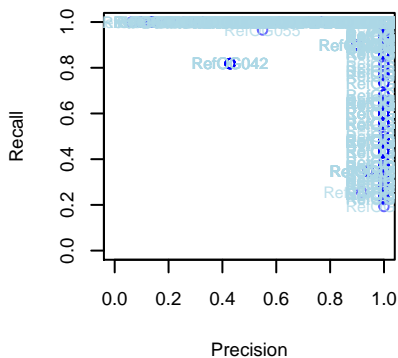
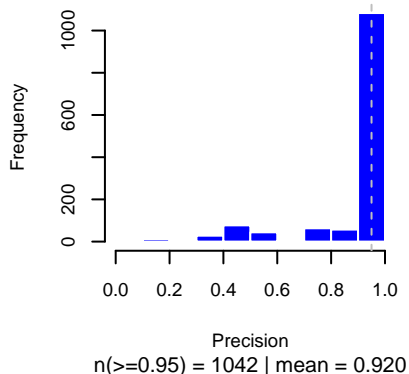


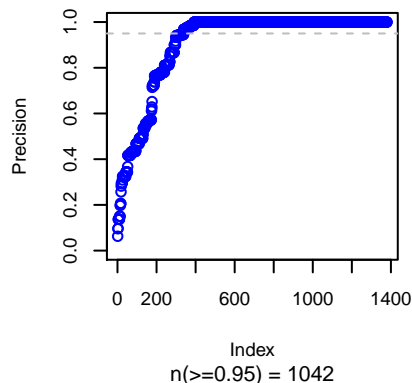
### Precision & recall



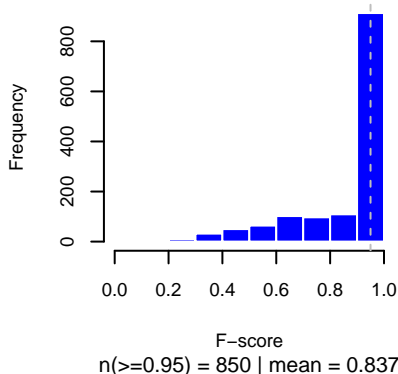
### Precision



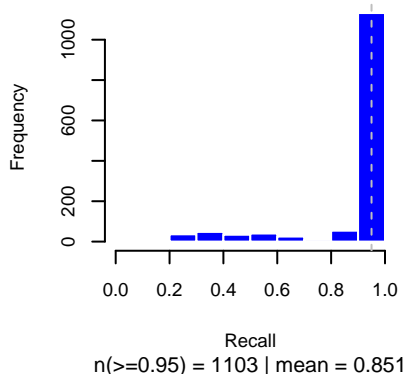
### Precision



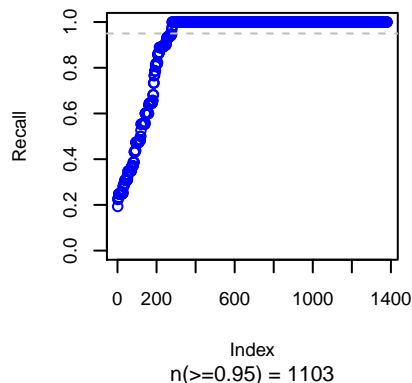
### F-score



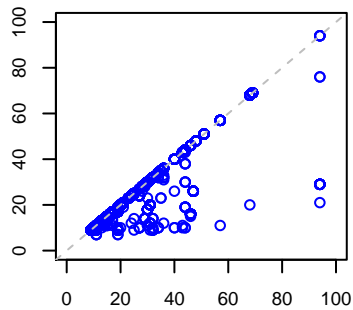
### Recall



### Recall



# genes in best OG



n(best > #OG all) = 140 (p=0.998)  
n(best = all) = 1101 (p=0.889 | r=1.000)  
n(best < all) = 279 (p=0.976 | r=0.591)

## number of reference genes in main and other OGs



## num OGs per refOG

