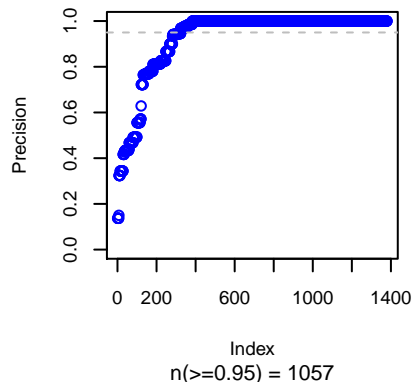
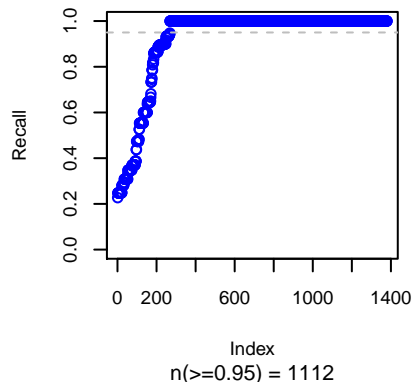


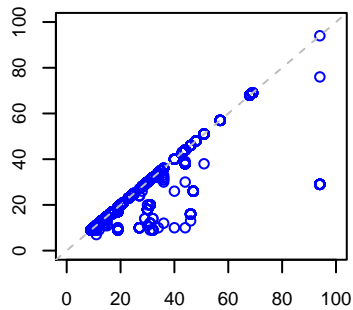
Precision



Recall



genes in best OG



n(best > # genes in ref OG) = 150 (p=0.997)
n(best = all) = 1112 (p=0.913 | r=1.000)
n(best < all) = 268 (p=0.980 | r=0.593)

number of reference genes in main and other OGs



num OGs per refOG

