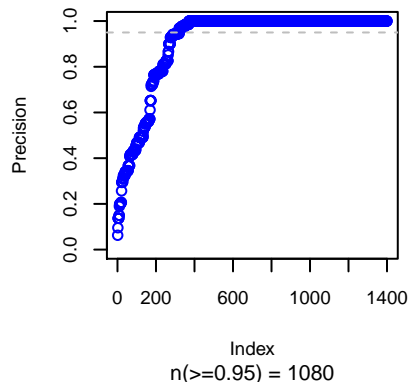
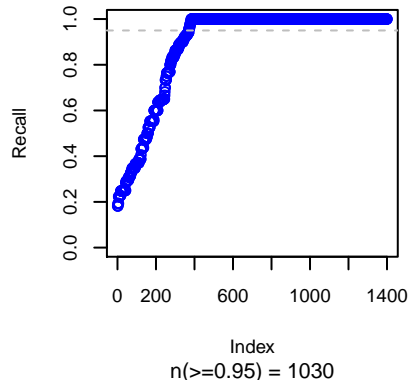


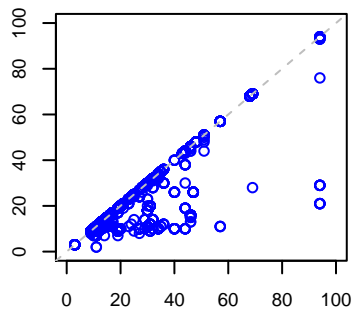
Precision



Recall



genes in best OG



n(best > #OG all) = 170 (p=0.997)
n(best = all) = 1018 (p=0.884 | r=1.000)
n(best < all) = 382 (p=0.978 | r=0.587)

number of reference genes in main and other OGs



num OGs per refOG

