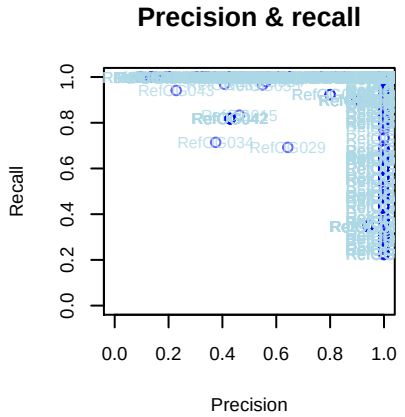
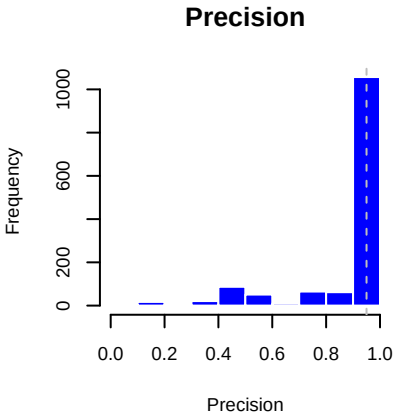
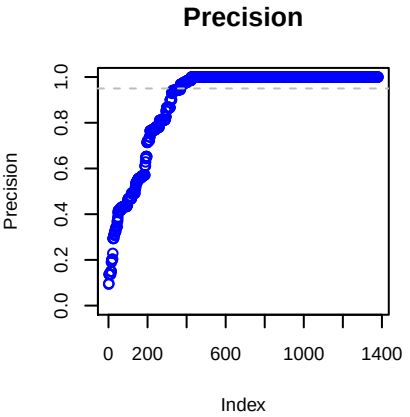
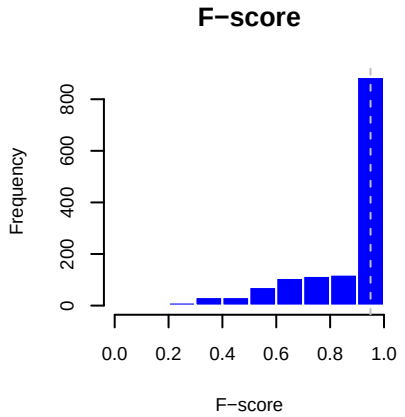




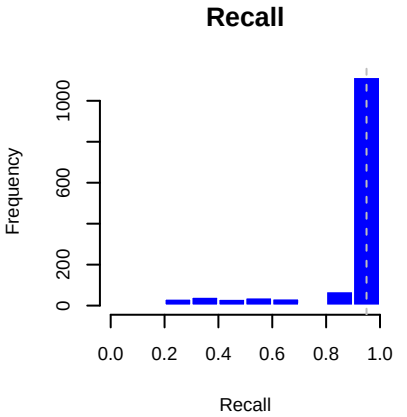
n(>=0.95) = 1012 | mean = 0.910



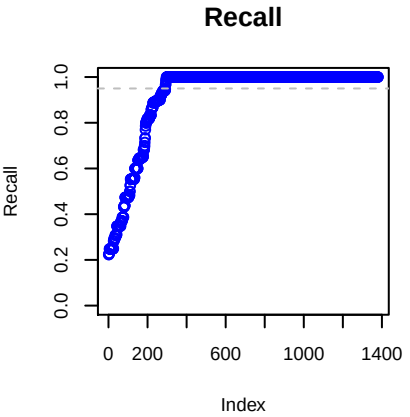
n(>=0.95) = 1012



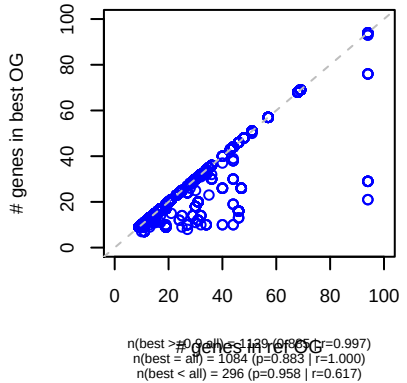
n(>=0.95) = 815 | mean = 0.838

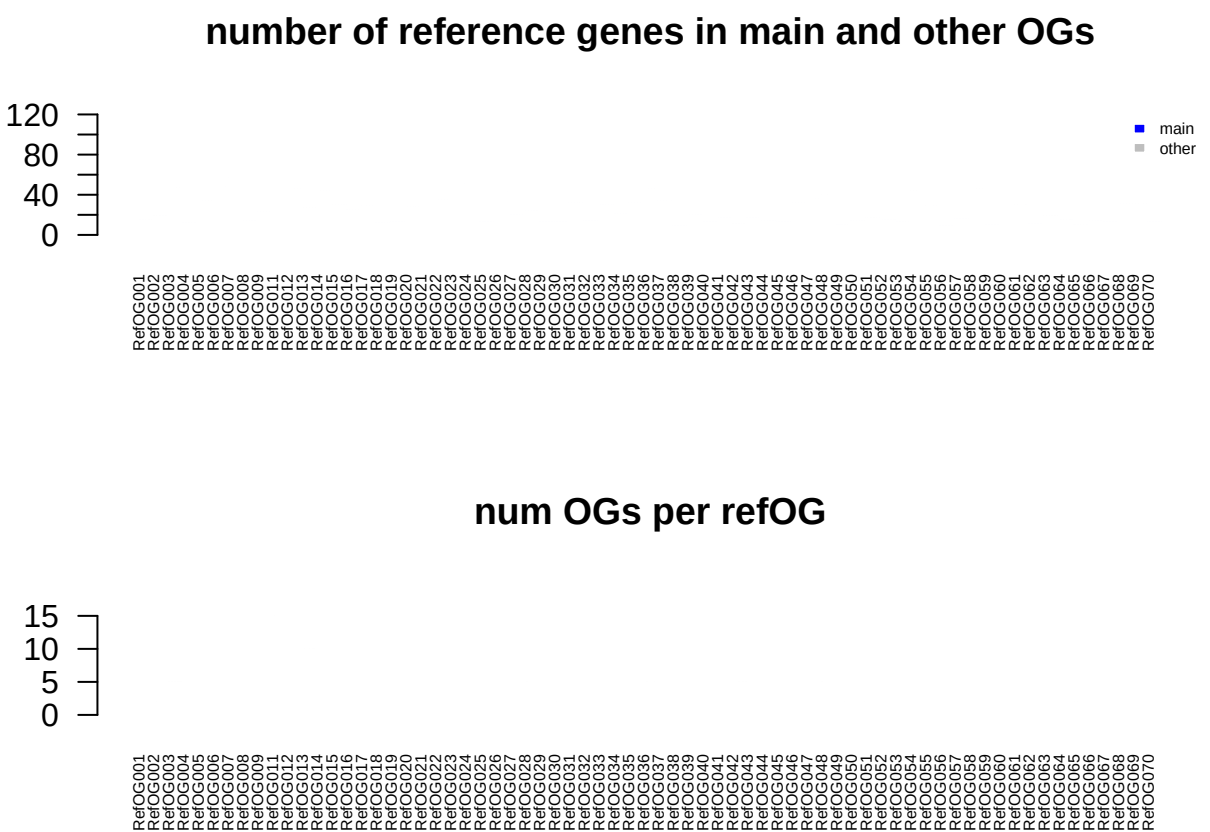


n(>=0.95) = 1089 | mean = 0.860



n(>=0.95) = 1089


genes in best OG
0
20
40
60
80
100
0
20
40
60
80
100
# genes in ref OG
n(best > all) = 1139 (p=0.995 | r=0.997)
n(best = all) = 1084 (p=0.883 | r=1.000)
n(best < all) = 296 (p=0.958 | r=0.617)


number of reference genes in main and other OGs
120
80
40
0
main
other
RefOG001
RefOG002
RefOG003
RefOG004
RefOG005
RefOG006
RefOG007
RefOG008
RefOG009
RefOG011
RefOG012
RefOG013
RefOG014
RefOG015
RefOG016
RefOG017
RefOG018
RefOG019
RefOG020
RefOG021
RefOG022
RefOG023
RefOG024
RefOG025
RefOG026
RefOG027
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RefOG056
RefOG057
RefOG058
RefOG059
RefOG060
RefOG061
RefOG062
RefOG063
RefOG064
RefOG065
RefOG066
RefOG067
RefOG068
RefOG069
RefOG070
num OGs per refOG
15
10
5
0
RefOG001
RefOG002
RefOG003
RefOG004
RefOG005
RefOG006
RefOG007
RefOG008
RefOG009
RefOG011
RefOG012
RefOG013
RefOG014
RefOG015
RefOG016
RefOG017
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