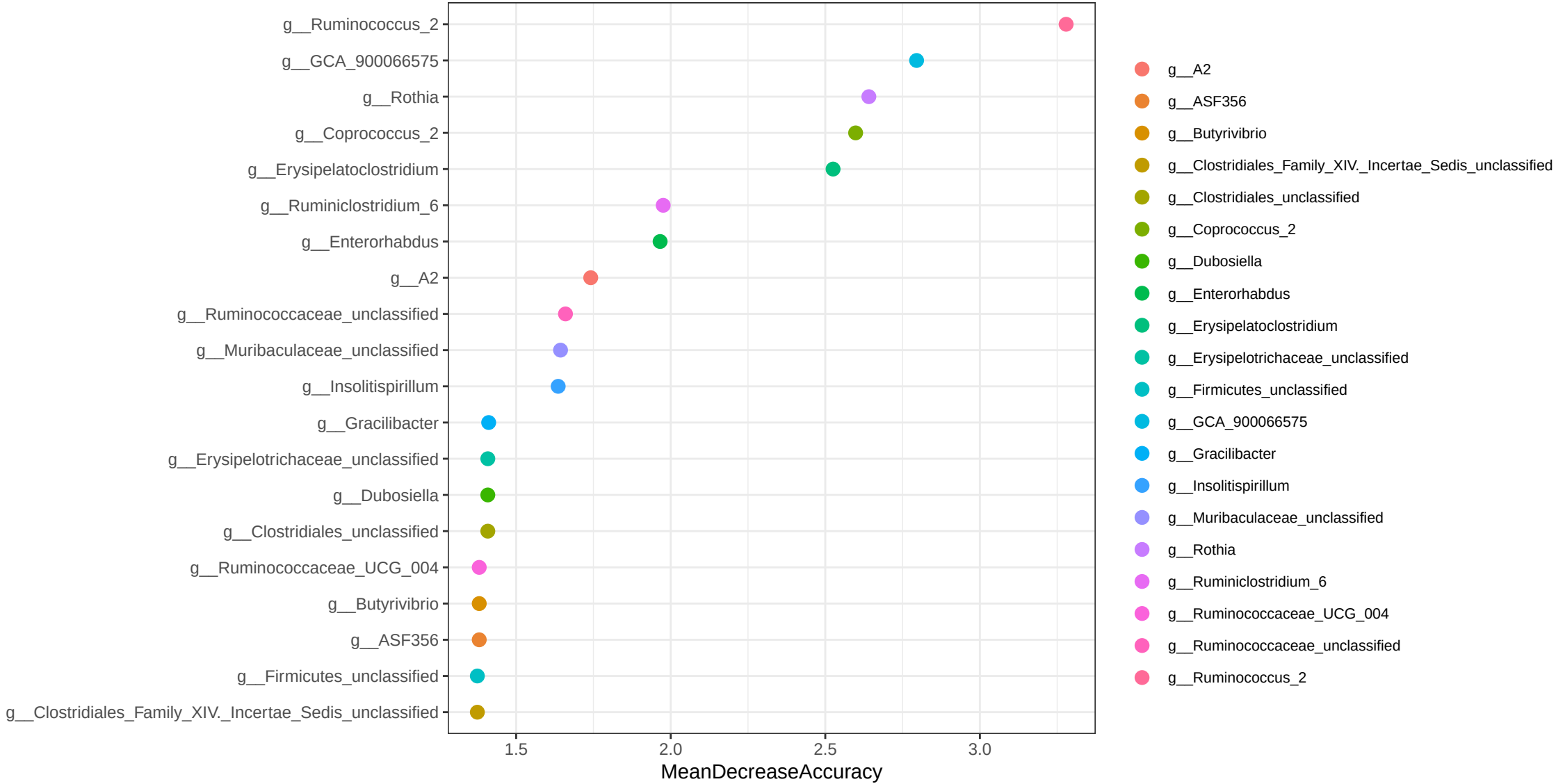
g__Ruminococcus_2
g__GCA_900066575
g__Rothia
g__Coprococcus_2
g__Erysipelatoclostridium
g__Ruminiclostridium_6
g__Enterorhabdus
g__A2
g__Ruminococcaceae_unclassified
g__Muribaculaceae_unclassified
g__Insolitispirillum
g__Gracilibacter
g__Erysipelotrichaceae_unclassified
g__Dubosiella
g__Clostridiales_unclassified
g__Ruminococcaceae_UCG_004
g__Butyrivibrio
g__ASF356
g__Firmicutes_unclassified
g__Clostridiales_Family_XIV._Incertae_Sedis_unclassified
1.5
2.0
2.5
3.0
MeanDecreaseAccuracy
g__A2
g__ASF356
g__Butyrivibrio
g__Clostridiales_Family_XIV._Incertae_Sedis_unclassified
g__Clostridiales_unclassified
g__Coprococcus_2
g__Dubosiella
g__Enterorhabdus
g__Erysipelatoclostridium
g__Erysipelotrichaceae_unclassified
g__Firmicutes_unclassified
g__GCA_900066575
g__Gracilibacter
g__Insolitispirillum
g__Muribaculaceae_unclassified
g__Rothia
g__Ruminiclostridium_6
g__Ruminococcaceae_UCG_004
g__Ruminococcaceae_unclassified
g__Ruminococcus_2