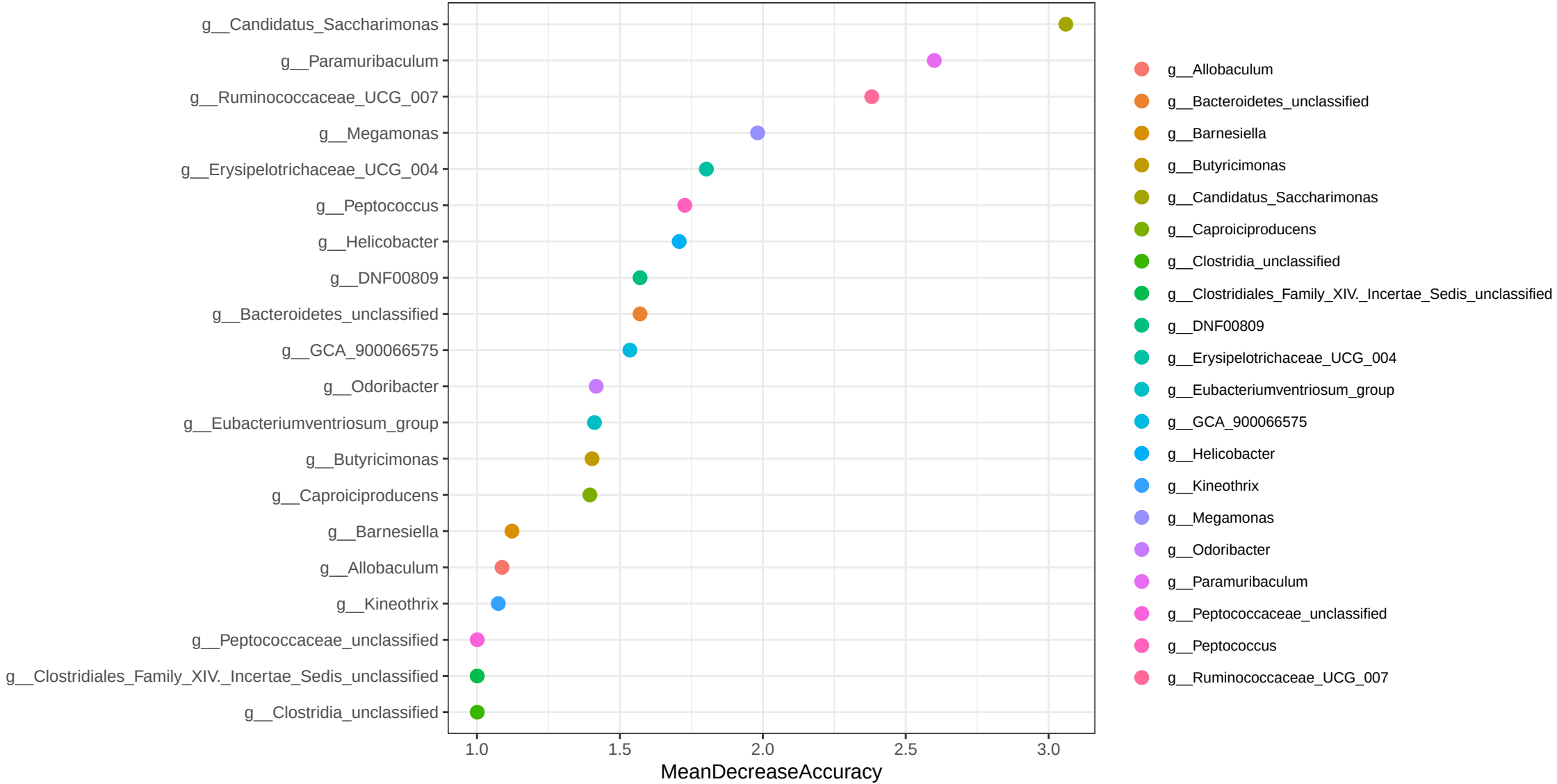
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g__Megamonas
g__Erysipelotrichaceae_UCG_004
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g__Helicobacter
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g__Odoribacter
g__Eubacteriumventriosum_group
g__Butyricimonas
g__Caproiciproducens
g__Barnesiella
g__Allobaculum
g__Kineothrix
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g__Clostridiales_Family_XIV._Incertae_Sedis_unclassified
g__Clostridia_unclassified
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1.5
2.0
2.5
3.0
MeanDecreaseAccuracy
g__Allobaculum
g__Bacteroidetes_unclassified
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g__Butyricimonas
g__Candidatus_Saccharimonas
g__Caproiciproducens
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