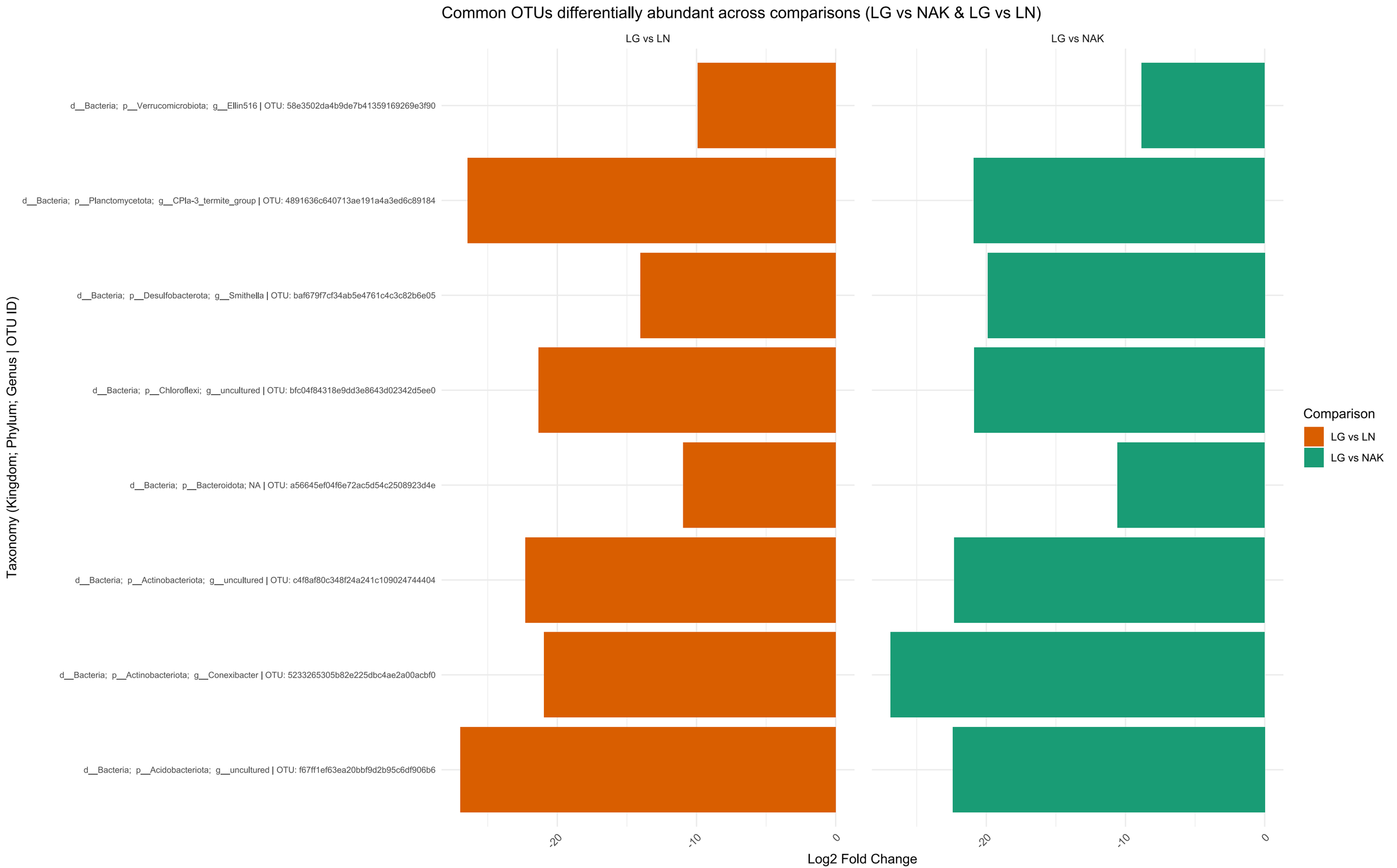
Common OTUs differentially abundant across comparisons (LG vs NAK & LG vs LN)
LG vs LN
LG vs NAK
Taxonomy (Kingdom; Phylum; Genus | OTU ID)
d__Bacteria; p__Verrucomicrobiota; g__Ellin516 | OTU: 58e3502da4b9de7b41359169269e3f90
d__Bacteria; p__Planctomycetota; g__CPla-3_termite_group | OTU: 4891636c640713ae191a4a3ed6c89184
d__Bacteria; p__Desulfobacterota; g__Smithella | OTU: baf679f7cf34ab5e4761c4c3c82b6e05
d__Bacteria; p__Chloroflexi; g__uncultured | OTU: bfc04f84318e9dd3e8643d02342d5ee0
d__Bacteria; p__Bacteroidota; NA | OTU: a56645ef04f6e72ac5d54c2508923d4e
d__Bacteria; p__Actinobacteriota; g__uncultured | OTU: c4f8af80c348f24a241c109024744404
d__Bacteria; p__Actinobacteriota; g__Conexibacter | OTU: 5233265305b82e225dbc4ae2a00acbf0
d__Bacteria; p__Acidobacteriota; g__uncultured | OTU: f67ff1ef63ea20bbf9d2b95c6df906b6
-20
-10
0
Log2 Fold Change
Comparison
LG vs LN
LG vs NAK