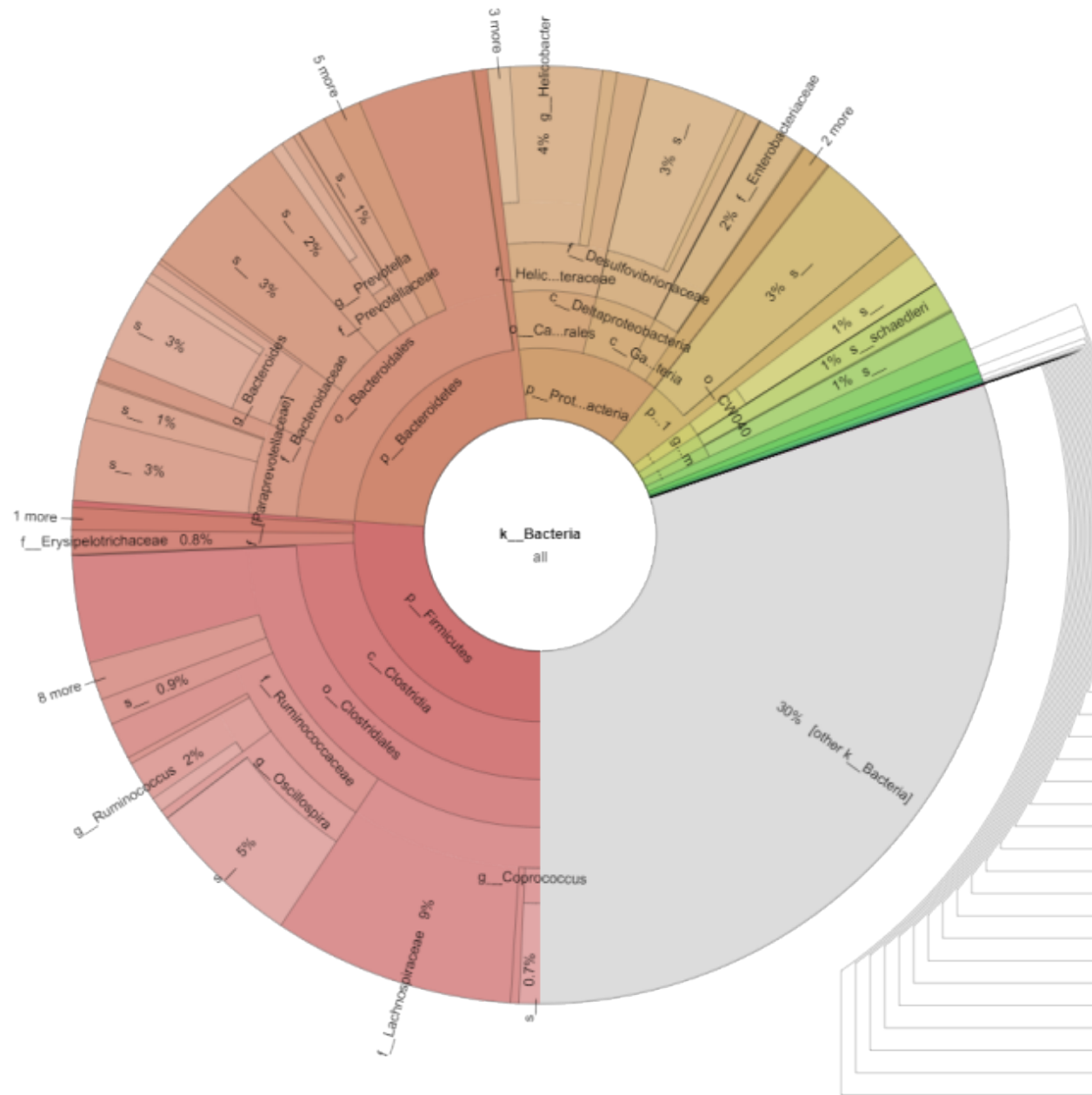
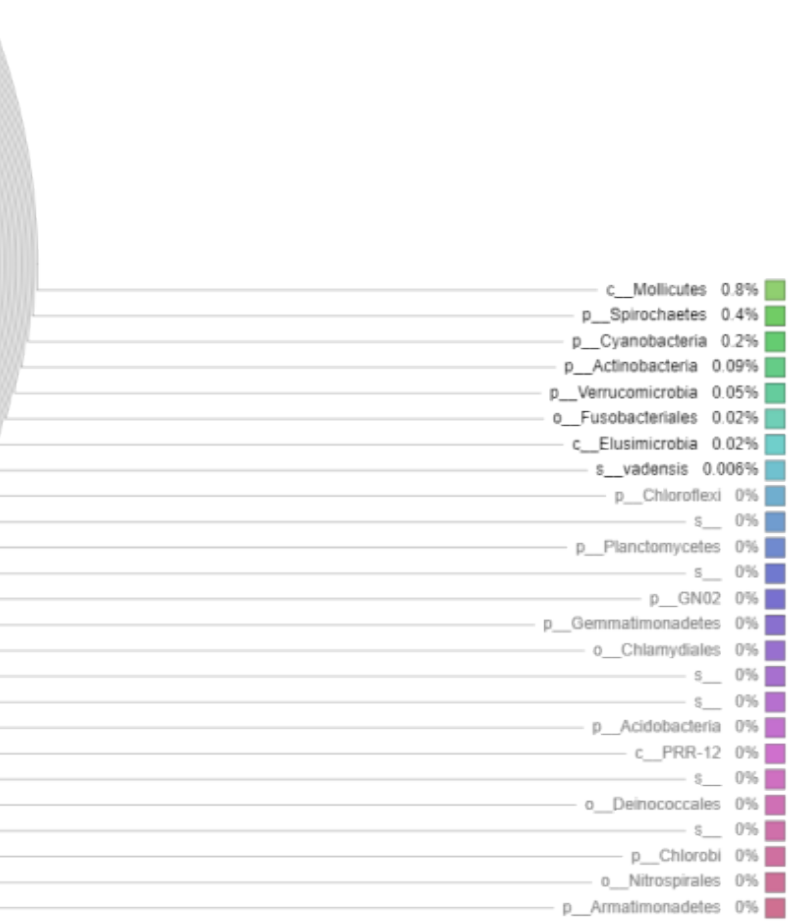


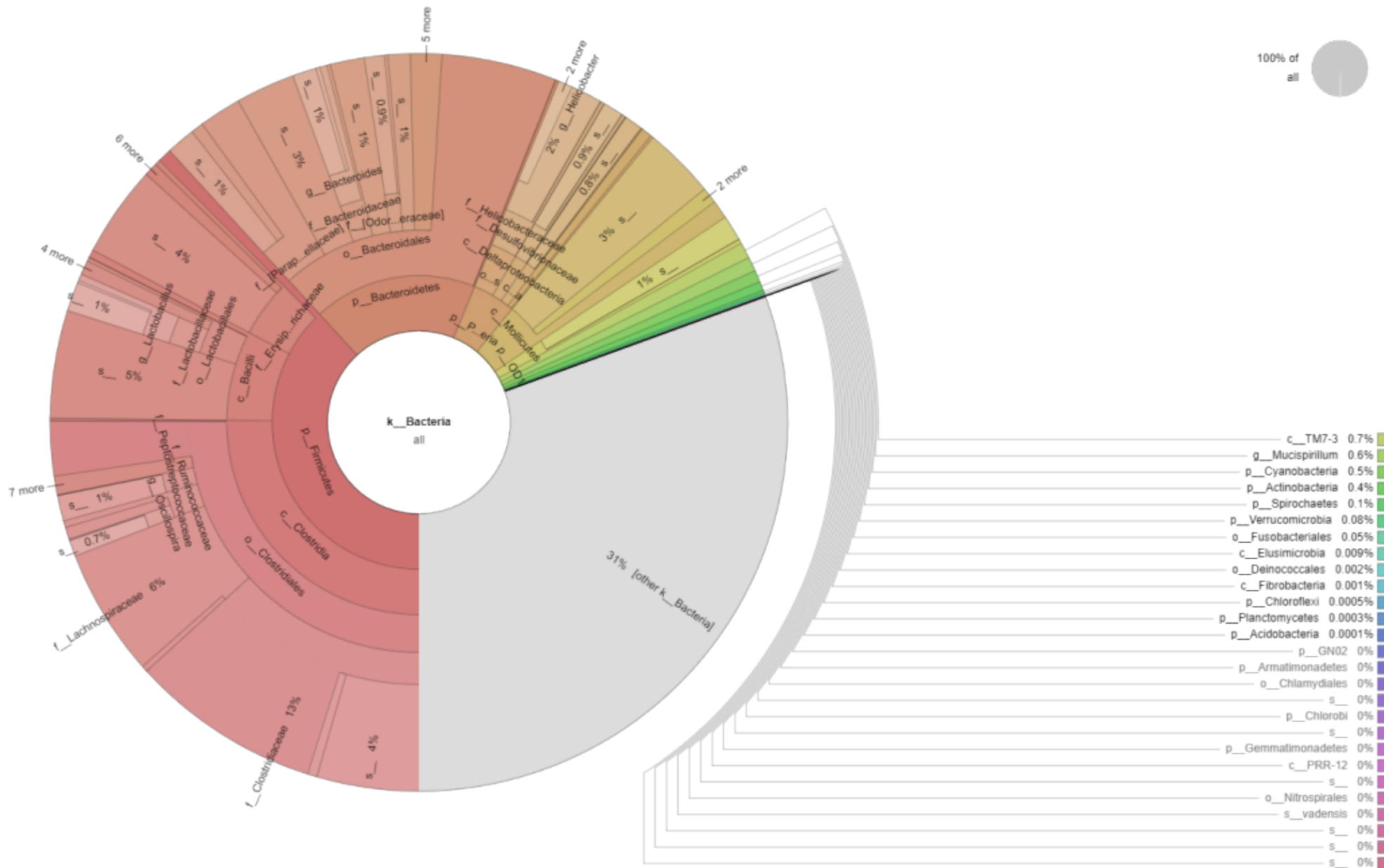
99% of all

- p_Actinobacteria 0.7%
- p_Verrucomicrobia 0.3%
- p_Cyanobacteria 0.2%
- c_Fibrobacteria 0.1%
- p_Acidobacteria 0.1%
- g_Mucispirillum 0.1%
- p_Spirochaetes 0.04%
- p_Gemmatimonadetes 0.04%
- o_Fusobacteriales 0.03%
- p_Planctomycetes 0.02%
- o_Nitrospirales 0.007%
- c_PRR-12 0.004%
- s_vadensis 0.003%
- c_Elusimicrobia 0.002%
- p_Armatimonadetes 0.002%
- p_Chloroflexi 0.002%
- o_Deinococcales 0.001%
- s_ 0.0009%
- s_ 0.0007%
- s_ 0.0003%
- p_Chlorobi 0.0003%
- s_ 0%
- p_GN02 0%
- o_Chlamydiales 0%
- s_ 0%
- s_ 0%



100% of all





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99% of all

p_Cyanobacteria	0.9%
c_Fibrobacteria	0.4%
p_Verrucomicrobia	0.1%
o_Fusobacteriales	0.07%
g_Mucispirillum	0.05%
p_Spirochaetes	0.02%
c_Elusimicrobia	0.007%
o_Deinococcales	0.004%
p_Acidobacteria	0.004%
p_Chloroflexi	0.003%
p_Chlorobi	0.003%
p_GN02	0.003%
p_Planctomycetes	0.002%
p_Gemmatimonadetes	0.002%
s_	0.002%
s_	0.0008%
o_Chlamydiales	0.0005%
o_Nitrospirales	0.0004%
s_	0.0003%
s_vadensis	0.0002%
s_	0.0001%
s_	0%
p_Armatimonadetes	0%
s_	0%
c_PRR-12	0%



98% of all

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