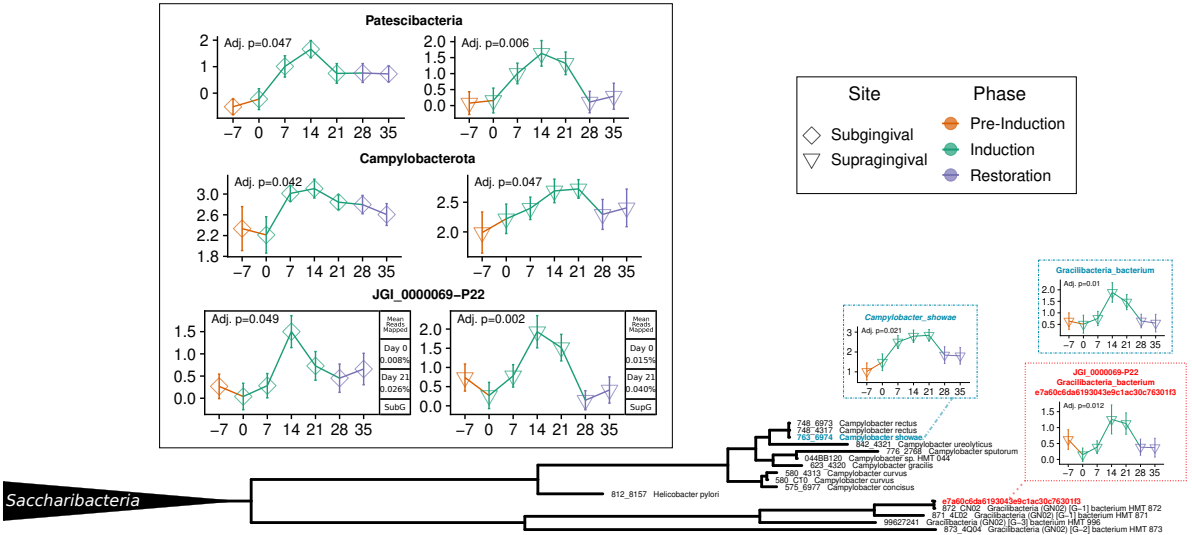
Patescibacteria
Adj. p=0.047
Adj. p=0.006
Campylobacterota
Adj. p=0.042
Adj. p=0.047
JGI_0000069–P22
Adj. p=0.049
Adj. p=0.002
Mean Reads Mapped
Day 0 0.008%
Day 21 0.026%
SubG
Day 0 0.015%
Day 21 0.040%
SupG
Site
Subgingival
Supragingival
Phase
Pre-Induction
Induction
Restoration
Campylobacter_showae
Adj. p=0.021
Gracilibacteria_bacterium
Adj. p=0.01
JGI_0000069-P22
Gracilibacteria_bacterium
e7a60c6da6193043e9c1ac30c76301f3
Adj. p=0.012
748_6973 Campylobacter rectus
748_4317 Campylobacter rectus
763_6974 Campylobacter showae
842_4321 Campylobacter ureolyticus
776_2768 Campylobacter sputorum
044BB120 Campylobacter sp. HMT 044
623_4320 Campylobacter gracilis
580_4313 Campylobacter curvus
580_C10 Campylobacter curvus
575_6977 Campylobacter concisus
812_8157 Helicobacter pylori
e7a60c6da6193043e9c1ac30c76301f3
872_CN02 Gracilibacteria (GN02) [G-1] bacterium HMT 872
871_4L02 Gracilibacteria (GN02) [G-1] bacterium HMT 871
99627241 Gracilibacteria (GN02) [G-3] bacterium HMT 996
873_4Q04 Gracilibacteria (GN02) [G-2] bacterium HMT 873
Saccharibacteria