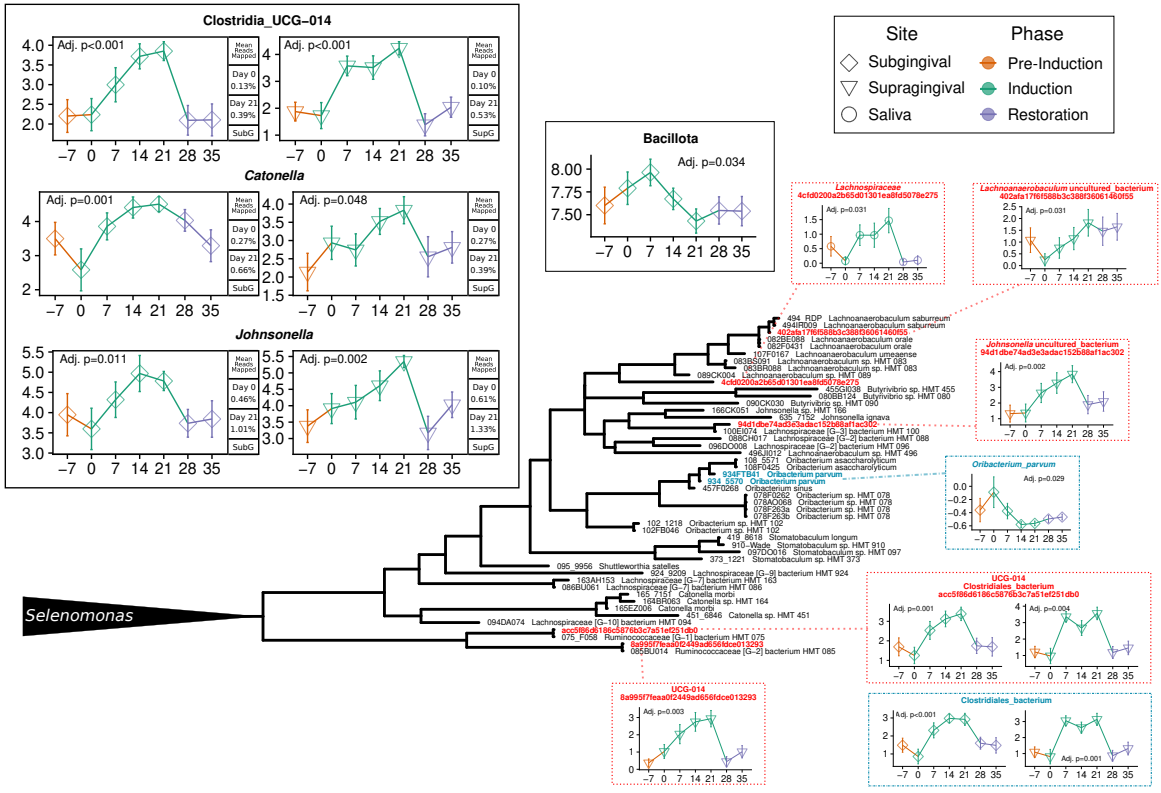
Site
Subgingival
Supragingival
Saliva
Phase
Pre-Induction
Induction
Restoration
Clostridia_UCG_014
Adj. p<0.001
Adj. p<0.001
Mean Reads Mapped
Day 0 0.13%
Day 21 0.39%
SubG
Day 0 0.10%
Day 21 0.53%
SupG
Catonella
Adj. p=0.001
Adj. p=0.048
Day 0 0.27%
Day 21 0.66%
Day 0 0.27%
Day 21 0.39%
Johnsonella
Adj. p=0.011
Adj. p=0.002
Day 0 0.46%
Day 21 1.01%
Day 0 0.61%
Day 21 1.33%
Bacillota
Adj. p=0.034
Lachnospiraceae 4cfd0200a2b65d01301ea8fd5078e275
Adj. p=0.031
Lachnoanaerobaculum uncultured_bacterium 402ab176f588b3c388f3606146bf55
Adj. p=0.031
Johnsonella uncultured_bacterium 94d1dbe74ad363adac152b68af1ac302
Adj. p=0.002
Oribacterium_parvum
Adj. p=0.029
494_RDP Lachnoanaerobaculum saburreum
494IR009 Lachnoanaerobaculum saburreum
402ab176f588b3c388f3606146bf55
082E088 Lachnoanaerobaculum orale
082F041 Lachnoanaerobaculum orale
107F0167 Lachnoanaerobaculum umeaense
083BS091 Lachnoanaerobaculum sp. HMT 083
083BR088 Lachnoanaerobaculum sp. HMT 083
089CK004 Lachnoanaerobaculum sp. HMT 089
4cfd0200a2b65d01301ea8fd5078e275
455S038 Butyrivibrio sp. HMT 455
080BB124 Butyrivibrio sp. HMT 080
090CK030 Butyrivibrio sp. HMT 090
166CK051 Johnsonella sp. HMT 166
635_7152 Johnsonella ignava
94d1dbe74ad363adac152b68af1ac302
100E074 Lachnospiraceae [G–3] bacterium HMT 100
088CH017 Lachnospiraceae [G–2] bacterium HMT 088
096CO008 Lachnospiraceae [G–2] bacterium HMT 096
496JI012 Lachnoanaerobaculum sp. HMT 496
108_5571 Oribacterium asaccharolyticum
108F0425 Oribacterium asaccharolyticum
934FT841 Oribacterium parvum
934_5970 Oribacterium parvum
457F0268 Oribacterium sinus
078F0262 Oribacterium sp. HMT 078
078AD068 Oribacterium sp. HMT 078
078F263a Oribacterium sp. HMT 078
078F263b Oribacterium sp. HMT 078
102_1218 Oribacterium sp. HMT 102
102FB048 Oribacterium sp. HMT 102
419_8618 Stomatobaculum longum
910–Wade Stomatobaculum sp. HMT 910
097DO016 Stomatobaculum sp. HMT 097
373_1221 Stomatobaculum sp. HMT 373
095_9956 Shuttleworthia satelles
924_9209 Lachnospiraceae [G–9] bacterium HMT 924
163AH153 Lachnospiraceae [G–7] bacterium HMT 163
086BU061 Lachnospiraceae [G–7] bacterium HMT 086
165_7151 Catonella morbi
164BR063 Catonella sp. HMT 164
165EZ006 Catonella morbi
451_6846 Catonella sp. HMT 451
094DA074 Lachnospiraceae [G–10] bacterium HMT 094
075_F058 Ruminococcaceae [G–1] bacterium HMT 075
8a995f7feaa0f249ad656fdce013293
085BU014 Ruminococcaceae [G–2] bacterium HMT 085
Selenomonas
UCG-014 Clostridiales_bacterium acc5f86d6186c5876b3c7a51ef251db0
Adj. p=0.001
Adj. p=0.004
UCG-014 8a995f7feaa0f249ad656fdce013293
Adj. p=0.003
Clostridiales_bacterium
Adj. p<0.001
Adj. p=0.001