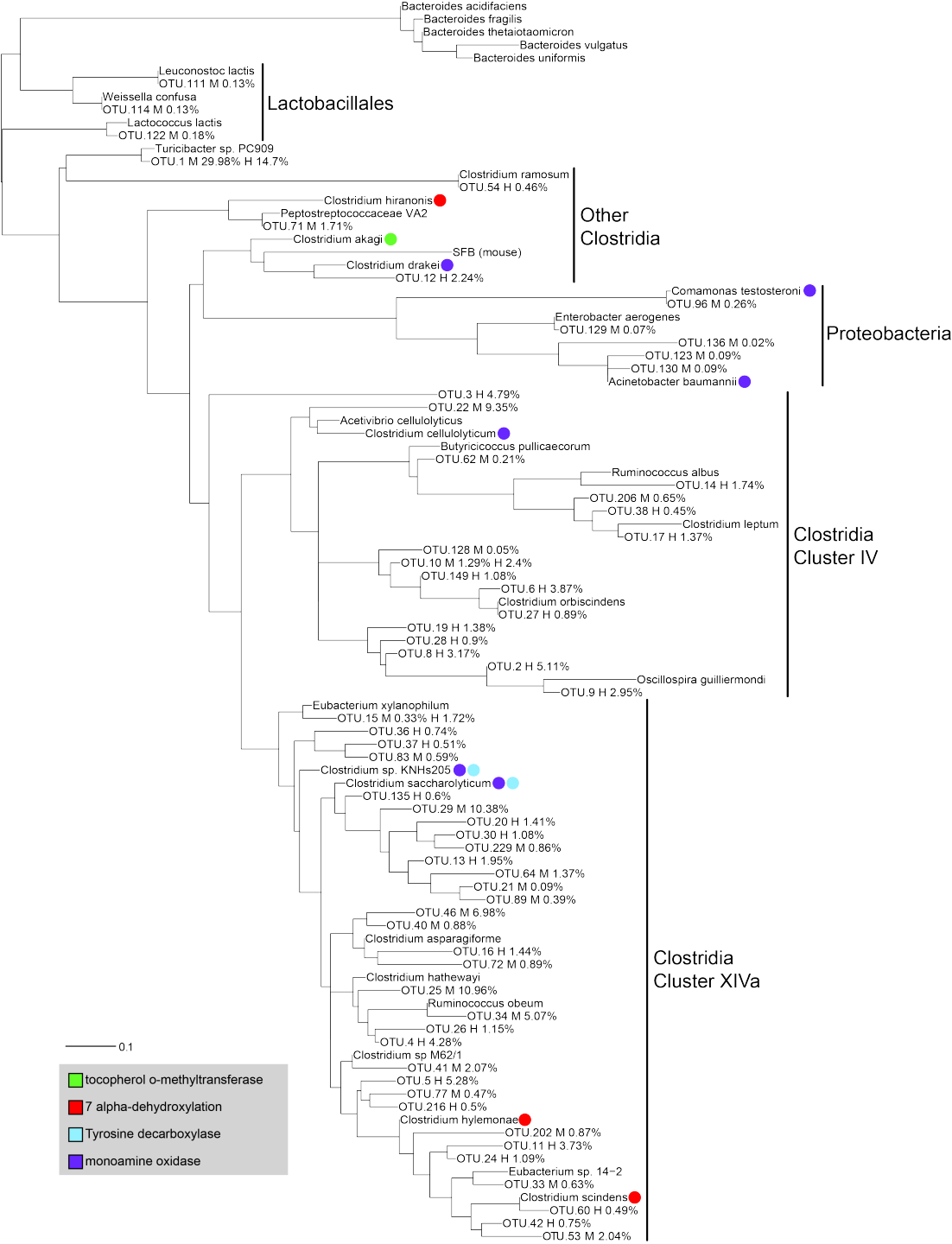
Bacteroides acidifaciens
Bacteroides fragilis
Bacteroides thetaiotaomicron
Bacteroides vulgatus
Bacteroides uniformis
Leuconostoc lactis
OTU.111 M 0.13%
Weissella confusa
OTU.114 M 0.13%
Lactococcus lactis
OTU.122 M 0.18%
Lactobacillales
Turicibacter sp. PC909
OTU.1 M 29.98% H 14.7%
Clostridium ramosum
OTU.54 H 0.46%
Clostridium hiranonis
Peptostreptococcaceae VA2
OTU.71 M 1.71%
Clostridium akagi
SFB (mouse)
Clostridium drakei
OTU.12 H 2.24%
Other
Clostridia
Comamonas testosteroni
OTU.96 M 0.26%
Enterobacter aerogenes
OTU.129 M 0.07%
OTU.136 M 0.02%
OTU.123 M 0.09%
OTU.130 M 0.09%
Acinetobacter baumannii
Proteobacteria
OTU.3 H 4.79%
OTU.22 M 9.35%
Acetivibrio cellulolyticus
Clostridium cellulolyticum
Butyricicoccus pullicaecorum
OTU.62 M 0.21%
Ruminococcus albus
OTU.14 H 1.74%
OTU.206 M 0.65%
OTU.38 H 0.45%
Clostridium leptum
OTU.17 H 1.37%
OTU.128 M 0.05%
OTU.10 M 1.29% H 2.4%
OTU.149 H 1.08%
OTU.6 H 3.87%
Clostridium orbiscindens
OTU.27 H 0.89%
OTU.19 H 1.38%
OTU.28 H 0.9%
OTU.8 H 3.17%
OTU.2 H 5.11%
Oscillospira guilliermondi
OTU.9 H 2.95%
Clostridia
Cluster IV
Eubacterium xylanophilum
OTU.15 M 0.33% H 1.72%
OTU.36 H 0.74%
OTU.37 H 0.51%
OTU.83 M 0.59%
Clostridium sp. KNHs205
Clostridium saccharolyticum
OTU.135 H 0.6%
OTU.29 M 10.38%
OTU.20 H 1.41%
OTU.30 H 1.08%
OTU.229 M 0.86%
OTU.13 H 1.95%
OTU.64 M 1.37%
OTU.21 M 0.09%
OTU.89 M 0.39%
OTU.46 M 6.98%
OTU.40 M 0.88%
Clostridium asparagiforme
OTU.16 H 1.44%
OTU.72 M 0.89%
Clostridia
Cluster XIVa
Clostridium hathewayi
OTU.25 M 10.96%
Ruminococcus obeum
OTU.34 M 5.07%
OTU.26 H 1.15%
OTU.4 H 4.28%
Clostridium sp M62/1
OTU.41 M 2.07%
OTU.5 H 5.28%
OTU.77 M 0.47%
OTU.216 H 0.5%
Clostridium hylemonae
OTU.202 M 0.87%
OTU.11 H 3.73%
OTU.24 H 1.09%
Eubacterium sp. 14−2
OTU.33 M 0.63%
Clostridium scindens
OTU.60 H 0.49%
OTU.42 H 0.75%
OTU.53 M 2.04%
0.1
tocopherol o-methyltransferase
7 alpha-dehydroxylation
Tyrosine decarboxylase
monoamine oxidase