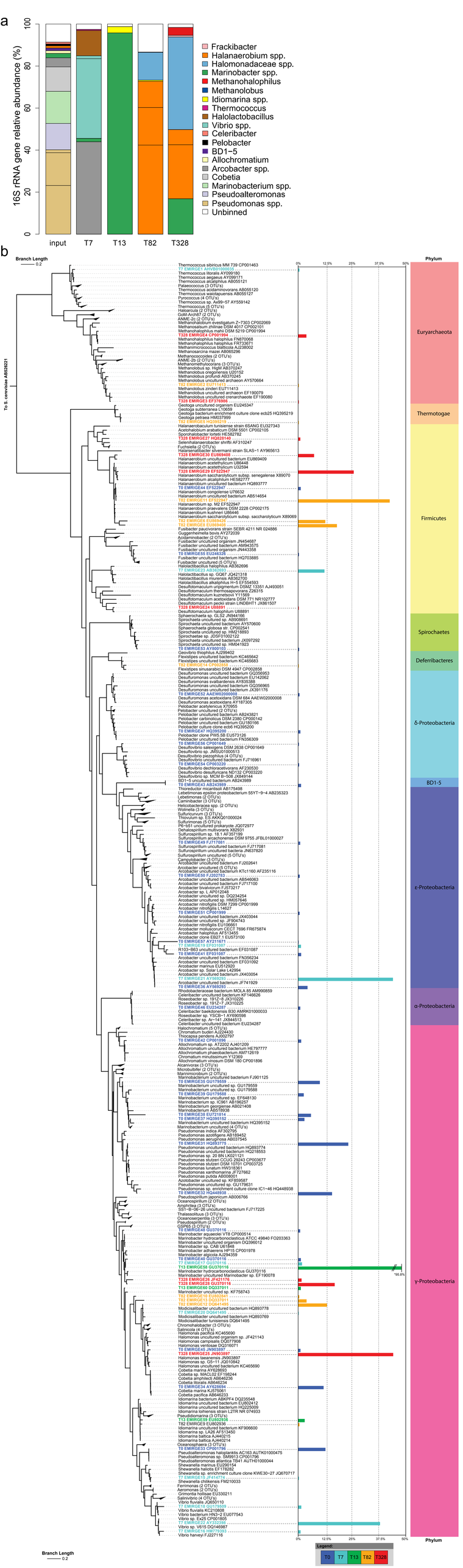
a
16S rRNA gene relative abundance (%)
0
20
40
60
80
100
input
T7
T13
T82
T328
Frackibacter
Halanaerobium spp.
Halomonadaceae spp.
Marinobacter spp.
Methanohalophilus
Methanolobus
Idiomarina spp.
Thermococcus
Halolactobacillus
Vibrio spp.
Celeribacter
Pelobacter
BD1−5
Allochromatium
Arcobacter spp.
Cobetia
Marinobacterium spp.
Pseudoalteromonas
Pseudomonas spp.
Unbinned
b
Branch Length
0.2
To S. cerevisiae AB526221
Phylum
Euryarchaeota
Thermotogae
Firmicutes
Spirochaetes
Deferribacteres
δ-Proteobacteria
BD1-5
ε-Proteobacteria
α-Proteobacteria
γ-Proteobacteria
0%
12.5%
25%
37.5%
50%
*95.8%
Legend:
T0
T7
T13
T82
T328