Bulk community
Active community
Relative abundance (%)
100
80
60
40
20
0
0.05
0.09
0.13
0.17
0.21
0.25
0.29
DFAA turnover
other bacteria
MB11C04 marine group
Stenotrophomonas
SAR116 clade
Sulfitobacter
Roseobacter NAC11-6
Roseobacter CHAB-I-5
Planktomarina
Synechococcus
Polaribacter
Owenweeksia
NS9 marine group
NS7 marine group
NS5 marine group
NS4 marine group
NS2b marine group
Marinoscillum
Formosa
Fluviicola