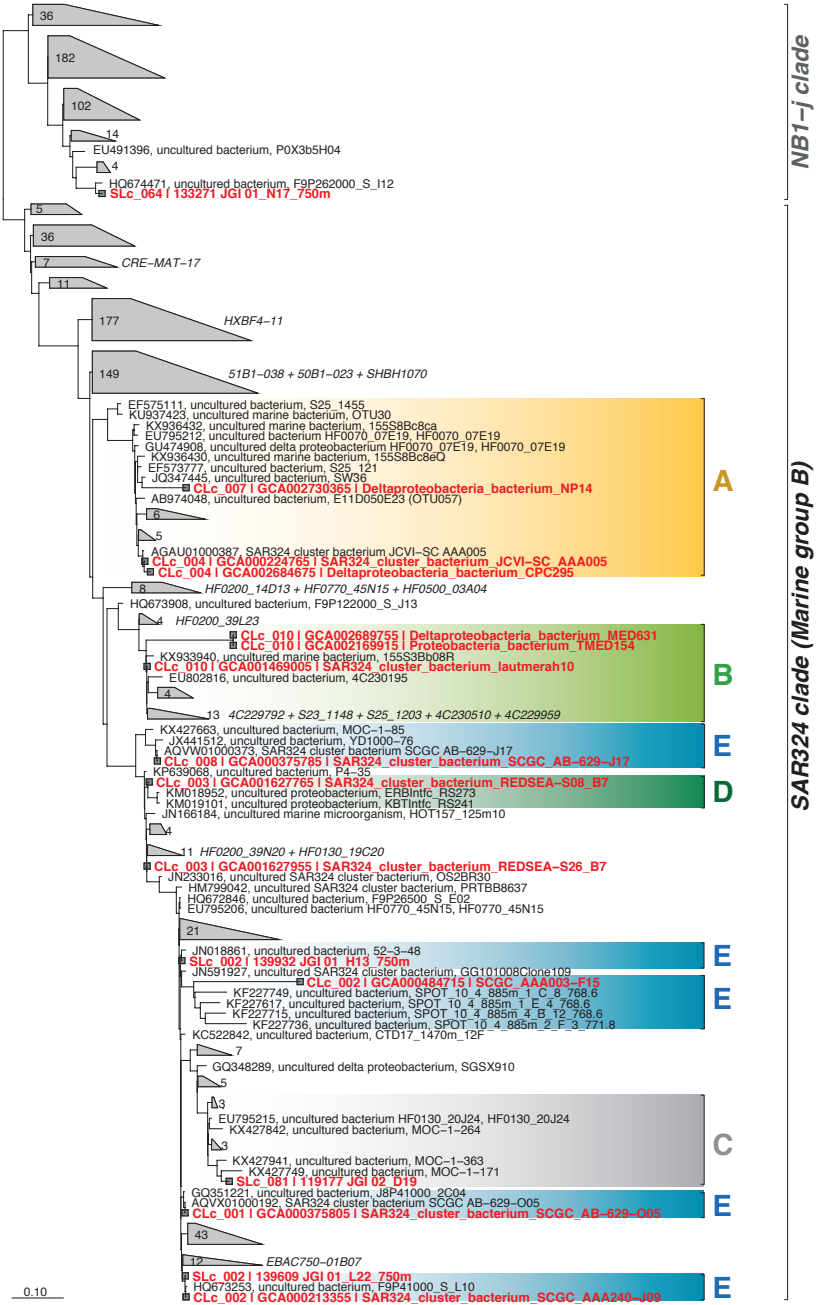
NB1–j clade
SAR324 clade (Marine group B)
36
182
102
14
EU491396, uncultured bacterium, P0X3b5H04
4
HQ674471, uncultured bacterium, F9P262000_S_I12
SLc_064 | 133271 JGI 01_N17_750m
5
36
7
CRE–MAT–17
11
177
HXBF4–11
149
51B1–038 + 50B1–023 + SHBH1070
EF575111, uncultured bacterium, S25_1455
KU937423, uncultured marine bacterium, OTU30
KX936432, uncultured marine bacterium, 155S8Bc8ca
EU795212, uncultured bacterium HF0070_07E19, HF0070_07E19
GU474908, uncultured delta proteobacterium HF0070_07E19, HF0070_07E19
KX936430, uncultured marine bacterium, 155S8Bc8eQ
EF573777, uncultured bacterium, S25_121
JQ347445, uncultured bacterium, SW36
CLc_007 | GCA002730365 | Deltaproteobacteria_bacterium_NP14
AB974048, uncultured bacterium, E11D050E23 (OTU057)
6
5
AGAU01000387, SAR324 cluster bacterium JCVI–SC AAA005
CLc_004 | GCA000224765 | SAR324_cluster_bacterium_JCVI–SC_AAA005
CLc_004 | GCA002684675 | Deltaproteobacteria_bacterium_CPC295
A
8
HF0200_14D13 + HF0770_45N15 + HF0500_03A04
HQ673908, uncultured bacterium, F9P122000_S_J13
4
HF0200_39L23
CLc_010 | GCA002689755 | Deltaproteobacteria_bacterium_MED631
CLc_010 | GCA002169915 | Proteobacteria_bacterium_TMED154
KX933940, uncultured marine bacterium, 155S3Bb08R
CLc_010 | GCA001469005 | SAR324_cluster_bacterium_lautmerah10
EU802816, uncultured bacterium, 4C230195
4
13
4C229792 + S23_1148 + S25_1203 + 4C230510 + 4C229959
B
KX427663, uncultured bacterium, MOC–1–85
JX441512, uncultured bacterium, YD1000–76
AQVW01000373, SAR324 cluster bacterium SCGC AB–629–J17
CLc_008 | GCA000375785 | SAR324_cluster_bacterium_SCGC_AB–629–J17
E
KP639068, uncultured bacterium, P4–35
CLc_003 | GCA001627765 | SAR324_cluster_bacterium_REDSEA–S08_B7
KM018952, uncultured proteobacterium, ERBIntfc_RS273
KM019101, uncultured proteobacterium, KBTIntfc_RS241
JN166184, uncultured marine microorganism, HOT157_125m10
D
4
11
HF0200_39N20 + HF0130_19C20
CLc_003 | GCA001627955 | SAR324_cluster_bacterium_REDSEA–S26_B7
JN233016, uncultured SAR324 cluster bacterium, OS2BR30
HM799042, uncultured SAR324 cluster bacterium, PRTBB8637
HQ672846, uncultured bacterium, F9P26500_S_E02
EU795206, uncultured bacterium HF0770_45N15, HF0770_45N15
21
JN018861, uncultured bacterium, 52–3–48
SLc_002 | 139932 JGI 01_H13_750m
E
JN591927, uncultured SAR324 cluster bacterium, GG101008Clone109
CLc_002 | GCA000484715 | SCGC_AAA003–F15
KF227749, uncultured bacterium, SPOT_10_4_885m_1_C_8_768.6
KF227617, uncultured bacterium, SPOT_10_4_885m_1_E_4_768.6
KF227715, uncultured bacterium, SPOT_10_4_885m_4_B_12_768.6
KF227736, uncultured bacterium, SPOT_10_4_885m_2_F_3_771.8
E
KC522842, uncultured bacterium, CTD17_1470m_12F
7
GQ348289, uncultured delta proteobacterium, SGSX910
5
3
EU795215, uncultured bacterium HF0130_20J24, HF0130_20J24
KX427842, uncultured bacterium, MOC–1–264
3
KX427941, uncultured bacterium, MOC–1–363
KX427749, uncultured bacterium, MOC–1–171
SLc_081 | 119177 JGI 02_D19
C
GQ351221, uncultured bacterium, J8P41000_2C04
AQVX01000192, SAR324 cluster bacterium SCGC AB–629–O05
CLc_001 | GCA000375805 | SAR324_cluster_bacterium_SCGC_AB–629–O05
E
43
12
EBAC750–01B07
SLc_002 | 139609 JGI 01_L22_750m
HQ673253, uncultured bacterium, F9P41000_S_L10
CLc_002 | GCA000213355 | SAR324_cluster_bacterium_SCGC_AAA240–J09
E
0.10