Supplementary Table 2: Details of 683 bacteria species

Sl	Accession	11 V		1	Sl	Acces		Tava Nama	(C : C^*)
No	No.	Taxa_Name	(G+C)%	I	No	No	).	Taxa_Name	(G+C)%
1 2	NC_000853 NC_000908	Thermotoga maritima MSB8  Mycoplasma genitalium G37	46.39 31.55			NC_01 NC_01		Mycoplasma arthritidis 158L3-1 Chloroherpeton thalassium ATCC 35110	31.34 46.14
3	NC_000908 NC_000911	Synechocystis sp. PCC 6803	48.59			NC_01		Prosthecochloris aestuarii DSM 271	50.84
4	NC_000911	Mycoplasma pneumoniae M129	40.66			NC_01		Pelodictyon phaeoclathratiforme BU-1	48.92
5	NC_000918	Aquifex aeolicus VF5	43.63		346	NC_01		Phenylobacterium zucineum HLK1	71.60
6	NC_000963	Rickettsia prowazekii str. Madrid E	30.42		347	NC_01		Geobacter bemidjiensis Bem	61.42
7	NC_001263	Deinococcus radiodurans R1	67.57		348	NC_01	1184	Vibrio fischeri MJ11	39.50
8	NC_002162	Ureaplasma parvum serovar 3 str. ATCC 700970	25.71		349	NC_01	1229	Borrelia duttonii Ly	27.81
9	NC_002488	Xylella fastidiosa 9a5c	53.79			NC_01		Borrelia recurrentis A1	27.70
10	NC_002528	Buchnera aphidicola str. APS (Acyrthosiphon pisum)	27.41		351	NC_01		Coprothermobacter proteolyticus DSM 5265	44.91
11	NC_002570	Bacillus halodurans C-125	44.32		352	NC_01		Thermodesulfovibrio yellowstonii DSM 11347	34.16
12 13	NC_002620 NC_002663	Chlamydia muridarum Nigg Pasteurella multocida subsp. multocida str. Pm70	40.69 41.04		354	NC_01 NC_01		Dictyoglomus thermophilum H-6-12 Aliivibrio salmonicida LFI1238	33.81 39.69
14	NC_002678	Mesorhizobium loti MAFF303099	63.47			NC_01		Oligotropha carboxidovorans OM5	63.05
15	NC_002755	Mycobacterium tuberculosis CDC1551	65.77			NC_01		Rhodospirillum centenum SW	70.68
16	NC_002771	Mycoplasma pulmonis UAB CTIP	27.29					Candidatus Azobacteroides pseudotrichonymphae genomovar. CFP2	
17	NC_002927	Bordetella bronchiseptica RB50	68.49		358	NC_01	1566	Shewanella piezotolerans WP3	44.17
18	NC_002928	Bordetella parapertussis 12822	68.43		359	NC_01	1567	Anoxybacillus flavithermus WK1	42.02
19	NC_002929	Bordetella pertussis Tohama I	68.13		360	NC_01		Acinetobacter baumannii AB0057	40.28
20	NC_002932	Chlorobium tepidum TLS	57.66		361	NC_01		Bifidobacterium longum subsp. infantis ATCC 15697 = JCM 1222	60.70
21	NC_002935	Corynebacterium diphtheriae NCTC 13129	54.08			NC_01		Thermosipho africanus TCF52B	30.73
22	NC_002936 NC_002937	Dehalococcoides ethenogenes 195	49.62 63.41			NC_01 NC_01		Dictyoglomus turgidum DSM 6724	33.99 68.77
24	NC_002937 NC_002939	Desulfovibrio vulgaris str. Hildenborough  Geobacter sulfurreducens PCA	61.58			NC_01		Thauera sp. MZ1T  Methylocella silvestris BL2	63.72
25	NC_002939 NC_002940	Haemophilus ducreyi 35000HP	38.74		366	NC_01		Borrelia burgdorferi ZS7	28.87
26	NC_002967	Treponema denticola ATCC 35405	38.27		367	NC_01		Escherichia fergusonii ATCC 35469	50.89
27	NC_002976	Staphylococcus epidermidis RP62A	32.94		368	NC_01		Vibrio splendidus LGP32	44.96
28	NC_002977	Methylococcus capsulatus str. Bath	63.96		369	NC_01	1761	Acidithiobacillus ferrooxidans ATCC 23270	59.32
29	NC_003047	Sinorhizobium meliloti 1021	63.49		370	NC_01		Desulfatibacillum alkenivorans AK-01	55.52
30	NC_003103	Rickettsia conorii str. Malish 7	32.85		371	NC_01		Haemophilus parasuis SH0165	40.78
31	NC_003155	Streptomyces avermitilis MA-4680	71.14			NC_01		Arthrobacter chlorophenolicus A6	66.78
32	NC_003212	Listeria innocua Clip11262	37.83		373	NC_01		Anaeromyxobacter dehalogenans 2CP-1	74.68
33 34	NC_003272 NC_003295	Nostoc sp. PCC 7120 Ralstonia solanacearum GMI1000	42.44 67.62			NC_01 NC_01		Methylobacterium nodulans ORS 2060 Clostridium cellulolyticum H10	69.61 38.16
35	NC_003255	Chlamydophila caviae GPIC	39.82			NC_01		Halothermothrix orenii H 168	38.89
36	NC_003869	Thermoanaerobacter tengcongensis MB4	37.80		377	NC_01		Thioalkalivibrio sulfidophilus HL-EbGr7	66.15
37	NC_003888	Streptomyces coelicolor A3(2)	72.45		378	NC_01		Thermotoga neapolitana DSM 4359	47.08
38	NC_003909	Bacillus cereus ATCC 10987	36.08		379	NC_01	1979	Geobacter daltonii FRC-32	54.29
39	NC_003910	Colwellia psychrerythraea 34H	38.82		380	NC_01	1989	Agrobacterium vitis S4	58.74
40	NC_003911	Ruegeria pomeroyi DSS-3	64.77		381			Acidovorax ebreus TPSY	67.46
41	NC_003912	Campylobacter jejuni RM1221	30.59		382	NC_01		Streptococcus uberis 0140J	37.06
42	NC_003919	Xanthomonas axonopodis pv. citri str. 306	65.12		383	NC_01		Caldicellulosiruptor bescii DSM 6725	35.58
43	NC_004113 NC_004116	Thermosynechococcus elongatus BP-1 Streptococcus agalactiae 2603V/R	54.54 36.06		384 385	NC_01 NC_01		Campylobacter lari RM2100 Desulfobacterium autotrophicum HRM2	29.93 49.71
45	NC_004110 NC_004129	Pseudomonas protegens Pf-5	64.10			NC_01		Nautilia profundicola AmH	33.57
46	NC_004193	Oceanobacillus iheyensis HTE831	36.08			NC_01		Staphylococcus carnosus subsp. carnosus TM300	35.26
47	NC_004310	Brucella suis 1330	58.21		388	NC_01	2416	Wolbachia sp. wRi	35.40
48	NC 004344	Wigglesworthia glossinidia endosymbiont of Glossina	23.65		389	NC_01	2438	Sulfurihydrogenibium azorense Az-Fu1	32.77
49	NC_004347	brevipalpis Shewanella oneidensis MR-1	46.99			NC_01		Persephonella marina EX-H1	37.38
50	NC 004350	Streptococcus mutans UA159	37.47		391	NC_01		Brucella melitensis ATCC 23457	58.26
51	NC_004369	Corynebacterium efficiens YS-314	63.79		392	NC_01		Streptococcus equi subsp. equi 4047	42.00
52	NC_004463	Bradyrhizobium diazoefficiens USDA 110	64.66		393	NC_01		Acidobacterium capsulatum ATCC 51196	61.10
53	NC_004547	Pectobacterium atrosepticum SCRI1043	52.19		394	NC_01	2489	Gemmatimonas aurantiaca T-27	64.49
54	NC_004552	Chlamydophila abortus S26/3	40.49		395	NC_01	2490	Rhodococcus erythropolis PR4	62.79
55	NC_004567	Lactobacillus plantarum WCFS1	45.53			NC_01		Brevibacillus brevis NBRC 100599	47.88
56	NC_004572	Tropheryma whipplei str. Twist	46.46			NC_01		Rhodococcus opacus B4	68.40
57	NC_004578	Pseudomonas syringae pv. tomato str. DC3000	59.31			NC_01		Deinococcus deserti VCD115	64.12
58 59	NC_004603 NC_004663	Vibrio parahaemolyticus RIMD 2210633  Bacteroides thetaiotaomicron VPI-5482	46.21 43.88			NC_01 NC_01		Azotobacter vinelandii DJ Clostridium botulinum A2 str. Kyoto	66.48 29.13
60	NC_004663 NC_004757	Nitrosomonas europaea ATCC 19718	51.57		400	NC_01		Bacillus anthracis str. CDC 684	36.02
61	NC_004829	Mycoplasma gallisepticum str. R(low)	32.02			NC_01		Sinorhizobium fredii NGR234	63.72
62	NC_004842	Anaplasma marginale str. St. Maries	49.80			NC_01		Corynebacterium aurimucosum ATCC 700975	61.43
63	NC_004917	Helicobacter hepaticus ATCC 51449	36.33		404	NC_01	2633	Rickettsia africae ESF-5	32.92
64	NC_005027	Rhodopirellula baltica SH 1	55.46		405	NC_01	2660	Pseudomonas fluorescens SBW25	61.35
65	NC_005043	Chlamydophila pneumoniae TW-183	41.28			NC_01		Beutenbergia cavernae DSM 12333	73.18
66	NC_005085	Chromobacterium violaceum ATCC 12472	65.71			NC_01		Exiguobacterium sp. AT1b	49.10
67	NC_005125	Gloeobacter violaceus PCC 7421	62.86		408	NC_01		Tolumonas auensis DSM 9187	49.80
68	NC_005126	Photorhabdus luminescens subsp. laumondii TTO1	44.16 47.16		409	NC_01		Corynebacterium kroppenstedtii DSM 44385	58.09
69 70	NC_005139 NC_005362	Vibrio vulnificus YJ016 Lactobacillus johnsonii NCC 533	47.16 34.96		410	NC_01 NC_01		Burkholderia glumae BGR1 Rickettsia peacockii str. Rustic	68.81 33.16
71	NC_005364	Mycoplasma mycoides subsp. mycoides SC str. PG1	24.09			NC_01		Kosmotoga olearia TBF 19.5.1	41.55
	NC_005823	Leptospira interrogans serovar Copenhageni str. Fiocruz L1-						•	
72		130	30.08			NC_01		Desulfovibrio magneticus RS-1	63.47
73	NC_005835	Thermus thermophilus HB27	69.60			NC_01		Bifidobacterium animalis subsp. lactis BI-04	61.40
74 75	NC_005861 NC_005955	Candidatus Protochlamydia amoebophila UWE25  Bartonella quintana str. Toulouse	36.10 40.40			NC_01 NC_01		Bartonella grahamii as4aup Desulfovibrio salexigens DSM 2638	40.02 48.02
76	NC_005955 NC_005956	Bartonella quintana str. 1 oulouse  Bartonella henselae str. Houston-1	40.40			NC_01		Streptococcus dysgalactiae subsp. equisimilis GGS_124	40.18
77	NC_005956 NC_005957	Bacillus thuringiensis serovar konkukian str. 97-27	36.00			NC_01		Dickeya zeae Ech1591	55.80
78	NC_005966	Acinetobacter sp. ADP1	41.50			NC_01		Pectobacterium carotovorum subsp. carotovorum PC1	53.20
79	NC_006055	Mesoplasma florum L1	27.05		420	NC_01		Geobacter sp. M21	61.70
80	NC_006087	Leifsonia xyli subsp. xyli str. CTCB07	68.39		421	NC_01	2968	Methylotenera mobilis JLW8	46.44
81	NC_006142	Rickettsia typhi str. Wilmington	30.37			NC_01		Methylovorus glucosetrophus SIP3-4	55.64
82	NC_006156	Borrelia garinii PBi	28.53			NC_01		Hirschia baltica ATCC 49814	45.86
83	NC_006177	Symbiobacterium thermophilum IAM 14863	69.29			NC_01		Teredinibacter turnerae T7901	51.70
84 85	NC_006270 NC_006300	Bacillus licheniformis DSM 13 = ATCC 14580 Mannheimia succiniciproducens MBEL55E	47.18			NC_01 NC_01		Dyadobacter fermentans DSM 18053	52.58 42.77
86	NC_006300 NC_006360	Mycoplasma hyopneumoniae 232	43.41 29.39			NC_01		Pedobacter heparinus DSM 2366 Actinosynnema mirum DSM 43827	74.02
87	NC 006361	Nocardia farcinica IFM 10152	71.16		428	NC_01		Brucella microti CCM 4915	58.24
88	NC_006370	Photobacterium profundum SS9	42.83			NC_01		Acidimicrobium ferrooxidans DSM 10331	68.37
89	NC_006449	Streptococcus thermophilus CNRZ1066	39.90			NC_01		Catenulispora acidiphila DSM 44928	70.24
90	NC_006510	Geobacillus kaustophilus HTA426	52.94		431	NC_01	3132	Chitinophaga pinensis DSM 2588	46.21

No	Accession No.	Taxa_Name	(G+C)%		No	Accession No.	Taxa_Name	(G+C)%
91	NC_006512	Idiomarina loihiensis L2TR	47.49			NC_013162	Capnocytophaga ochracea DSM 7271	40.57
92	NC_006513	Aromatoleum aromaticum EbN1	65.48			NC_013166	Kangiella koreensis DSM 16069	44.29
93	NC_006582	Bacillus clausii KSM-K16	45.33			NC_013171	Anaerococcus prevotii DSM 20548	36.53
94 95	NC_006677 NC_006832	Gluconobacter oxydans 621H	61.71 30.58			NC_013173 NC_013174	Desulfomicrobium baculatum DSM 4028  Jonesia denitrificans DSM 20603	59.60 58.53
96	NC_006908	Ehrlichia ruminantium str. Welgevonden  Mycoplasma mobile 163K	25.35			NC_013174 NC_013192	Leptotrichia buccalis C-1013-b	30.79
97	NC_006932	Brucella abortus bv. 1 str. 9-941	58.25		438	NC_013194	Candidatus Accumulibacter phosphatis clade IIA str. UW-1	64.32
98	NC_006958	Corynebacterium glutamicum ATCC 13032	54.78			NC_013198	Lactobacillus rhamnosus GG	47.45
99	NC_007109	Rickettsia felis URRWXCal2	32.64		440	NC_013203	Atopobium parvulum DSM 20469	46.31
100	NC_007146	Haemophilus influenzae 86-028NP	38.79		441	NC_013204	Eggerthella lenta DSM 2243	64.85
101	NC_007164	Corynebacterium jeikeium K411	62.25			NC_013205	Alicyclobacillus acidocaldarius subsp. acidocaldarius DSM 446	62.93
102	NC_007168	Staphylococcus haemolyticus JCSC1435	33.42			NC_013209	Acetobacter pasteurianus IFO 3283-01	53.77
103 104	NC_007204	Psychrobacter arcticus 273-4	44.59			NC_013216 NC_013222	Desulfotomaculum acetoxidans DSM 771	42.75
	NC_007292 NC_007294	Candidatus Blochmannia pennsylvanicus str. BPEN  Mycoplasma synoviae 53	32.06 28.77		445	NC_013222 NC_013223	Robiginitalea biformata HTCC2501  Desulfohalobium retbaense DSM 5692	56.07 58.06
105	NC_007298	Dechloromonas aromatica RCB	59.87			NC_013235	Nakamurella multipartita DSM 44233	71.22
		Staphylococcus saprophyticus subsp. saprophyticus ATCC						
107	NC_007350	15305	34.02		448	NC_013282	Cronobacter turicensis z3032	58.48
108	NC_007354	Ehrlichia canis str. Jake	30.99		449	NC_013355	Zymomonas mobilis subsp. mobilis NCIMB 11163	48.21
109	NC_007356	Dehalococcoides sp. CBDB1	47.85		450	NC_013385	Ammonifex degensii KC4	59.93
	NC_007384	Shigella sonnei Ss046	52.34			NC_013410	Fibrobacter succinogenes subsp. succinogenes S85	48.89
111 112	NC_007406 NC_007413	Nitrobacter winogradskyi Nb-255  Anabaena variabilis ATCC 29413	63.02 42.55			NC_013411 NC_013421	Geobacillus sp. Y412MC61 Pectobacterium wasabiae WPP163	53.37 51.78
	NC_007413	Pseudoalteromonas haloplanktis TAC125	41.00			NC_013421 NC_013422	Halothiobacillus neapolitanus c2	55.49
114	NC_007498	Pelobacter carbinolicus DSM 2380	56.01			NC_013440	Haliangium ochraceum DSM 14365	69.92
	NC_007503	Carboxydothermus hydrogenoformans Z-2901	42.40			NC_013441	Gordonia bronchialis DSM 43247	67.42
	NC_007508	Xanthomonas campestris pv. vesicatoria str. 85-10	65.09			NC_013454	Blattabacterium sp. (Blattella germanica) str. Bge	27.25
	NC_007510	Burkholderia lata	66.90			NC_013501	Rhodothermus marinus DSM 4252	65.24
118	NC_007514	Chlorobium chlorochromatii CaD3	45.06		459	NC_013510	Thermomonospora curvata DSM 43183	72.22
	NC_007517	Geobacter metallireducens GS-15	60.35			NC_013511	Mycoplasma hominis ATCC 23114	27.38
120	NC_007519	Desulfovibrio alaskensis G20	58.24		461	NC_013512	Sulfurospirillum deleyianum DSM 6946	39.35
121	NC_007520	Thiomicrospira crunogena XCL-2	43.97			NC_013515	Streptobacillus moniliformis DSM 12112	26.20
	NC_007575	Sulfurimonas denitrificans DSM 1251	34.78			NC_013517	Sebaldella termitidis ATCC 33386	34.69
123	NC_007576 NC_007604	Lactobacillus sakei subsp. sakei 23K	41.91 56.05			NC_013520 NC_013522	Veillonella parvula DSM 2008 Thermanaerovibrio acidaminovorans DSM 6589	39.67 64.45
	NC_007606	Synechococcus elongatus PCC 7942 Shigella dysenteriae Sd197	52.36			NC_013522 NC_013530	Xylanimonas cellulosilytica DSM 15894	72.74
	NC_007614	Nitrosospira multiformis ATCC 25196	54.85			NC_013530	Anaplasma centrale str. Israel	50.03
127	NC_007626	Magnetospirillum magneticum AMB-1	66.13		468	NC_013642	Thermotoga naphthophila RKU-10	46.34
128	NC_007633	Mycoplasma capricolum subsp. capricolum ATCC 27343	24.00		469	NC_013656	Lactococcus lactis subsp. lactis KF147	35.80
129	NC_007643	Rhodospirillum rubrum ATCC 11170	66.10		470	NC_013714	Bifidobacterium dentium Bd1	59.32
130	NC_007645	Hahella chejuensis KCTC 2396	54.78		471	NC_013716	Citrobacter rodentium ICC168	55.92
131	NC_007651	Burkholderia thailandensis E264	68.08		472	NC_013720	Pirellula staleyi DSM 6068	57.91
	NC_007712	Sodalis glossinidius str. 'morsitans'	56.19			NC_013722	Xanthomonas albilineans GPE PC73	63.47
	NC_007722	Erythrobacter litoralis HTCC2594	63.43			NC_013729	Kribbella flavida DSM 17836	70.85
	NC_007794	Novosphingobium aromaticivorans DSM 12444	65.55 42.56			NC_013730	Spirosoma linguale DSM 74 Conexibacter woesei DSM 14684	51.32 72.93
	NC_007797 NC_007798	Anaplasma phagocytophilum str. HZ Neorickettsia sennetsu str. Miyayama	41.41			NC_013739 NC_013740	Acidaminococcus fermentans DSM 20731	57.15
	NC_007799	Ehrlichia chaffeensis str. Arkansas	31.42			NC 013757	Geodermatophilus obscurus DSM 43160	74.13
	NC_007802	Jannaschia sp. CCS1	62.69			NC_013766	Listeria monocytogenes 08-5578	38.43
139	NC_007880	Francisella tularensis subsp. holarctica LVS	32.87		480	NC_013791	Bacillus pseudofirmus OF4	40.71
140	NC_007899	Chlamydophila felis Fe/C-56	39.95		481	NC_013799	Hydrogenobacter thermophilus TK-6	44.13
141	NC_007907	Desulfitobacterium hafniense Y51	48.48		482	NC_013851	Allochromatium vinosum DSM 180	64.94
142	NC_007908	Rhodoferax ferrireducens T118	60.62		483	NC_013853	Streptococcus mitis B6	40.74
143	NC_007929	Lactobacillus salivarius UCC118	33.16		484	NC_013889	Thioalkalivibrio sp. K90mix	66.29
	NC_007948	Polaromonas sp. JS666	63.16			NC_013892	Xenorhabdus bovienii SS-2004	46.32
145 146	NC_007951 NC_007954	Burkholderia xenovorans LB400	63.24 46.12			NC_013894 NC_013921	Thermocrinis albus DSM 14484	47.01 34.27
146	NC_007934 NC_007964	Shewanella denitrificans OS217	62.65		488	NC_013921 NC_013929	Thermoanaerobacter italicus Ab9	71.76
	NC_007964 NC_007969	Nitrobacter hamburgensis X14 Psychrobacter cryohalolentis K5	44.03		489	NC_013929 NC_013939	Streptomyces scabiei 87.22 Deferribacter desulfuricans SSM1	31.08
149	NC_007973	Cupriavidus metallidurans CH34	64.35		490	NC_013943	Denitrovibrio acetiphilus DSM 12809	43.20
150	NC_007984	Baumannia cicadellinicola str. Hc (Homalodisca coagulata)	34.32		491	NC_013946	Meiothermus ruber DSM 1279	64.12
	NC_008025	Deinococcus geothermalis DSM 11300	67.00			NC_013947	Stackebrandtia nassauensis DSM 44728	68.66
152	NC_008027	Pseudomonas entomophila L48	64.93		493	NC_013948	Mycoplasma agalactiae	30.09
153	NC_008044	Ruegeria sp. TM1040	60.80		494	NC_013949	Helicobacter mustelae 12198	42.89
	NC_008048	Sphingopyxis alaskensis RB2256	65.87			NC_013956	Pantoea ananatis LMG 20103	54.64
155	NC_008095	Myxococcus xanthus DK 1622	69.13		496	NC_013959	Sideroxydans lithotrophicus ES-1	58.19
	NC_008150	Yersinia pestis Antiqua	48.94			NC_013960	Nitrosococcus halophilus Nc 4	52.52
157	NC_008209	Roseobacter denitrificans OCh 114	59.48		498	NC_013961	Erwinia amylovora CFBP1430	54.68
158 159	NC_008228 NC_008229	Pseudoalteromonas atlantica T6c Helicobacter acinonychis str. Sheeba	45.42 38.79		499 500	NC_014006 NC_014008	Sphingobium japonicum UT26S  Coraliomargarita akajimensis DSM 45221	65.35 53.93
	NC_008229 NC_008254	Chelativorans sp. BNC1	61.56			NC_014008 NC_014011	Aminobacterium colombiense DSM 12261	45.75
161	NC_008255	Cytophaga hutchinsonii ATCC 33406	39.61			NC_014011 NC_014012	Shewanella violacea DSS12	45.73
	NC_008260	Alcanivorax borkumensis SK2	55.24			NC_014019	Bacillus megaterium QM B1551	39.17
	NC_008261	Clostridium perfringens ATCC 13124	29.41			NC_014032	Salinibacter ruber M8	66.73
164	NC_008268	Rhodococcus jostii RHA1	67.92			NC_014033	Prevotella ruminicola 23	48.93
	NC_008277	Borrelia afzelii PKo	28.48			NC_014034	Rhodobacter capsulatus SB 1003	66.88
	NC_008278	Frankia alni ACN14a	73.03		507	NC_014039	Propionibacterium acnes SK137	60.42
167	NC_008313	Ralstonia eutropha H16	67.06		508	NC_014098	Kyrpidia tusciae DSM 2912	59.56
	NC_008319	Synechococcus sp. CC9311	53.44		509	NC_014106	Lactobacillus crispatus ST1	37.57
169	NC_008343	Granulibacter bethesdensis CGDNIH1	59.62			NC_014109	Candidatus Riesia pediculicola USDA	29.50
170 171	NC_008345 NC_008346	Shewanella frigidimarina NCIMB 400	42.49 45.68		511 512	NC_014147 NC_014150	Moraxella catarrhalis RH4	42.94 29.00
	NC_008346 NC_008358	Syntrophomonas wolfei subsp. wolfei str. Goettingen Hyphomonas neptunium ATCC 15444	45.68 62.34			NC_014150 NC_014151	Brachyspira murdochii DSM 12563 Cellulomonas flavigena DSM 20109	74.31
172	NC_008497	Lactobacillus brevis ATCC 367	47.11			NC_014151 NC_014152	Thermincola potens JR	46.64
174	NC_008508	Leptospira borgpetersenii serovar Hardjo-bovis str. L550	41.26			NC_014153	Thiomonas intermedia K12	64.47
	NC_008530	Lactobacillus gasseri ATCC 33323	35.54			NC_014158	Tsukamurella paurometabola DSM 20162	68.79
	NC_008531	Leuconostoc mesenteroides subsp. mesenteroides ATCC	38.31			NC_014165		72.48
		8293					Thermobispora bispora DSM 43833	
	NC_008541	Arthrobacter sp. FB24	65.96			NC_014166	Arcobacter nitrofigilis DSM 7299	28.62
	NC_008554	Syntrophobacter fumaroxidans MPOB	60.20			NC_014168 NC_014207	Segniliparus rotundus DSM 44985	67.22
179	NC_008555 NC_008570	Listeria welshimeri serovar 6b str. SLCC5334  Aeromonas hydrophila subsp. hydrophila ATCC 7966	36.75 62.81			NC_014207 NC_014212	Methylotenera versatilis 301 Meiothermus silvanus DSM 9946	43.53 63.01
	NC_008570 NC_008571	Gramella forsetii KT0803	37.27				Propionibacterium freudenreichii subsp. shermanii CIRM-BIA1	67.42
	0003/1	Omnorm 15.50th K1 0005	21.21			01721J	r	U1.42

No.     Proceed   Procee		Accession	Taxa_Name	(G+C)%	Sl	Accession	Taxa_Name	(G+C)%
N. P. 1960   Programmer (1967)   Programmer (1968)   Programmer			Shewanella sp. ANA-3	49.11			Desulfurivibrio alkaliphilus AHT2	61.16
Second	183	NC_008593	Clostridium novyi NT	29.13	524	NC_014218	Arcanobacterium haemolyticum DSM 20595	53.68
18   18   18   18   18   18   18   18			·					
18   1.00   1								
18. P. 10. Sep. 10. Contained formits analysis and cardiolysis and process of the proce						_	-	
March   Marc	188				529		•	
198   198	189	NC_008610		35.36	530	NC_014230	Croceibacter atlanticus HTCC2559	34.33
18   S. C.	190	NC_008611		65.74	531	NC_014246	Mobiluncus curtisii ATCC 43063	55.89
No.   Septime   Septime	191				532			40.40
19.   19.	192		*	49.20			Acinetobacter oleivorans DR1	39.77
18   1.0							_	
No.   1967-100   1968-100   1969-100   196								
18   No.   19.00529   Montanetame properties   19.00529   19.005								
19.   N. 1966/16   Memmether reported YM 25   Sep   No. 1948   Cheesiam inguigable DSM 1538   1.28	197	NC_008711	Arthrobacter aurescens TC1	62.76	538	NC_014319	Leuconostoc gasicomitatum LMG 18811	37.29
10.   10.			·					
18.   18.		_	•					
10.   1.0.								
10. N. C. 200781         Rounders hombitional KOSPN         94.4         58. N. 01878         Choodines non-bombinem working 18.4         43.5           50. N. C. 200786         Commission and microscope in the contemp strategy of the processing and process are part of the commission								
18.   18.	203	NC_008782	Acidovorax sp. JS42	66.82	544	NC_014366	gamma proteobacterium HdN1	53.69
No. 1987   Vermissperincher seinass EPIU-2   0.529   5.7 No. 1987   S. 1988   S. 19								
M. D. (2008/08)         Humbachodospinaluplainish(s) (4.38)         4.38 (4.38)         4.39 (8.9.) (1.394)         Challes of Galional Conferencemena S2 (3.34)         5.3.4           D. (2008)         N. (2008)         Prechimentoroccus mateins sr. MIT (19.0)         4.28 (4.38)         5.9.         8.9. (19.14)         Challes of Galional Longeniteric membas S2 (3.34)         5.3.4           M. (2009)         Controller of Mittal (19.0)         4.04         4.3.         8.0.         8.0.         1.0.         Challes (19.0)         1.0.         4.2.         7.0.         1.0.         1.0.         7.0.         7.0.         1.0.         4.0.         7.0.         1.0.         4.0.							* *	
NS         NS         Value (mission of March March)         43.9         NS         Collision (mission of March March)         3.35         NS         Collision (mission of March)         4.35         NS         NS         NS         Collision (mission of March)         4.35         NS         NS         NS         NS         Collision (mission of March)         4.35         NS			ı			_	_	
12.   12.		_						
11.1         12.0         P. 1.00007000         Melandication periadomical H1100         68.71         533         N. 1.9482         H1.500070         4.5000000         6.7.7         4.5000000         6.7.7         4.5000000         6.7.7         5.5000000         7.5000000         6.7.7         5.5000000         7.50000000         7.50000000         6.7.7         5.50000000         7.500000000000000         6.7.7         6.7.7         7.5000000000000000000000000000000000000	209	NC_008820	Prochlorococcus marinus str. MIT 9303	51.25	550	NC_014410	Thermoanaerobacterium thermosaccharolyticum DSM 571	34.58
12.1 S. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1.								
13.1         N. (200908)         Chostrations difficile (20)         551         P. (5) (45)         Beallian unpolisipatione (1984)         5.62         P. (5)         P. (7)         P. (5)         P. (7)         P. (7			_					
14.1                 C. 1009/32                 Section Sect							-	
150   N. 0.00943   Suckampolopous cyrllines NRU, 238   71.65   57   N. 0.014679   BilliAbocaccinum Inflorms NT   45.72   58   N. 0.014679   Bellia storophases NP2   41.72								
217 N. P. (200728)         Desiration mechanism microsis M. 1         4.12 bit M. (200728)         S. (200728)         R. (200727)         Chlorobitum pilorum PVK-CK (% 6.27 kg.)         5.59 N. (200484)         C. (200484)         S. (200484)         C. (200			Herminiimonas arsenicoxydans				[Clostridium] sticklandii	
18   N. 1909373   Chlorobium placoviletionical DSM 205   53-64   559   N. 2014605   Caldiscillulosimpor hydrobiants (5   5-25)   5-25   N. 2014605   Caldiscillulosimpor hydrobiants (5   5-25)   N. 2014605   Caldiscillulosimpor hydrobiants (5   5-								
219         N. 200338         Mycordestering alyour PR-CCK         68.27         550         N. 201465         C. Cabificellulosingupor hydrothermatis 108         36.2           210         N. 200339         Mycordesteric reconstrict subsp. symbolicus QLW         45.5         P. 50         N. 201465         Cabificellulosingupor nomenenis OL         35.8           212         N. 200338         Salinispora tropics CRS-4         0.80         4.56         N. 201409         Robothers cryption of the control of the							*	
200   N.C.   1009430   Accordances altenonicals surbay, surbanecials A449   9.78   9.78   9.78   N.C.   1009450   Polymicohacter creates usbep, assignment only accordance of the No.   1.000450   N.C.   1.000			*					
22 NC,00938   Salimpront tropics CNP-14-00   69,00   69,00   70,00046   Enterobacter pt. 618   54,10   54,00047   Caldicellusion reproductive pt. 618   54,10   54,00047   Caldicellusion recording percentage (NP)   54,00047   Caldicellusion patch resistant politocoline UVI 01   54,50   55,00048   Severatella patterfacieres CN-32   54,50   50,00048   Severatella patterfacieres CN-32   54,50   50,00048   Severatella patterfacieres CN-32   54,50   50,00048   Severatella patterfacieres CN-32   54,50   50,00044   Parobacteria politocoline UVI 01   54,94   50,00048   Severatella patterfacieres CN-32   54,50   50,00048   Severatella patterfacieres CN-32   54,50   50,00048   Severatella patterfaciere Archae VCS/TB3   41,85   50,00048   Severatella patterfaciere Archae VCS/TB3   41,85   50,00048   Candidate Vericerroprocinc charatil IA   32,85   50,00048   Candidate Vericerroprocinc charatil IA   32,85   50,00048   Candidate Vericerroprocinc charatil IA   52,85   50,00048   Severatella patterfaciere Archae VCS/TB3   50,00048   Candidate Vericerroprocinc charatil IA   52,95   50,00048   Severatella patterfacier Archae VCS/TB3   50,00048   Candidate Vericerroprocinc charatil IA   52,95   50,00048   Severatella patterfacier Archae VCS/TB3   50,00048   Severatella patterface Archae VCS/TB3   50,00048   Severatella patterfa	220	NC_009348		59.78	561			33.74
222         NC.009380         Salinipora tropiace CNP4-40         69.80         49.80         SC.016450         R.Bodococcas oqui 10SS         99.25           223         NC.00947         Caldicellulosingroor saccharopticas DNR 8903         35.61         55.6         NC.01472         Caldicellulosingroor knowledge profession SNR 3903         35.61           255         NC.009441         Ehroducterium johanosine UW 101         34.94         35.95         NC.009442         Larcharophila profession SNR 1903         34.94         38.92           276         NC.009445         Decharibacter randoms VCS1703A         4.51         35.95         NC.016433         Mcdylovorus sp. MP88         52.29           278         NC.009445         Candidatas Vesiconysoscias odutail HA         45.01         NC.016403         Psecadulationomic promotine Braid         45.05           279         NC.009445         Candidatas Vesiconysoscias odutail HA         45.01         NC.016403         Psecadulationomic promotine Braid         45.02           281         NC.009448         Bodyloval promotine St. 19         45.02         NC.016403         Psecadulationomic promotine Braid         45.02           287         NC.009505         Brancial covis ATCC 25840         58.27         NC.016410         Tayloridate patient promotine Braid         17.22	221	NC_009379		45.26	562	NC_014657	Caldicellulosiruptor owensensis OL	35.83
23 N. (200436)         Enterwolvector p. (6.88)         54.10         54.10         Caldicellulosingore knowlessy (2002)         35.5           24 N. (200434)         Shawmand patterdicates CN 32         45.50         55.8         NC. (1944)         Lacchhaellulasingore knizsy (2014)         35.9           27 N. (20044)         Shawmand patterdicates CN 32         45.50         NC. (1944)         Sterptococcus and 65271133         4.18         56.8         NC. (1944)         Lacchhaellulas amplorums (2014)         35.9           27 N. (20044)         Sterptococcus and 65271133         4.18         4.18         56.8         NC. (1945)         Mechylorum sp. (1945)         35.9           28 N. (20044)         Chrobaelter molosus VCST/ROA         4.51         1.50         NC. (1946)         1.40         4.50         Nc. (1946)         Nc.	222	NC 009380		69.80	563	NC 014659	Rhodococcus equi 103S	69.27
225         NC,009418         Shewanella purefacienc N-22         45.0         Jest         NC,019472         Lacoboxality Methylosoms, MPR08         55.2           227         NC,009442         Streptococcus sin OSZYBE3         41.8         1.5         NC,019452         Methylosoms, MPR08         52.2           228         NC,009446         Chronicolomic control columni HA         43.01         1.5         NC,019452         Nessental calantine Q200.0         3.23           230         NC,009446         Chronicolomic columni HA         3.23         1.7         NC,018410         Pecudoalteromans sys.MM913         41.00           231         NC,009445         Accidiphilium control in Extr. 1         46.5         1.7         NC,018410         Helicobacter falis ATCC 49179         44.80           232         NC,009445         Braciphinobism pi, ITVA1         65.0         373         NC,018421         Mycoplasmi tementam M61         27.21           233         NC,009454         Lejocolal permophilia RNL1         46.2         9.7         NC,018421         Mycoplasmi tementam M61         27.21           245         NC,009513         Bracella ovis ATCC 28440         88.7         378         NC,018421         Mycoplasmi tementam M64         27.21           257         NC,009512								
226         NC.009444         Elmobracerium johnnosina CW101         34.94         578         NC.014373         Methodosens and MPG88         5.22           278         NC.009446         Dichelobacter modosus VCS1703A         4.50         1         56         NC.014512         Nectorium Landianus Vesicomyoscuc ostantial A         3.25         3.75         NC.014803         Pesadosterium componential POS         23.96           28         NC.009444         Cardidianus Vesicomyoscucio ostantial NATA         32.85         4         571         NC.014803         Pesadosterium morphomos y SMP913         4.489           23         NC.009444         Acidiphillum cyprum IF-5         68.51         571         NC.014912         Mycopilama Emericania MCEP         27.2           23         NC.009486         Bradyinzobium cyprum IF-5         68.51         572         NC.014921         Mycopilama Emericania MCEP         27.2           23         NC.009494         Legiomella preumophila RCO         35.9         4         757         NC.014921         Mycopilama Emericania MSM1271         7.13           23         NC.009513         Phycophocater sp. PR961         461         478         NC.014921         Mycopilama Emericania MSM1271         7.2           23         NC.009614         Psychocater sp. PR961	224	NC_009437	Caldicellulosiruptor saccharolyticus DSM 8903	35.63	565	NC_014721	Caldicellulosiruptor kristjanssonii 177R1B	36.57
227         NC. 09442         Skreptococcus sais 05/YH133         41.85         568         NC. 01472         Mc. 009460         Dischloshed rendosinu VCEP103A         4.51         568         NC. 014732         Nc. 009460         Candidatus Vesicomyroscius okutani IIA         32.85         7.00         Nc. 014732         Nc. 014782         Nc. 014810         Peaudoalteromonias sp. Ne9913         41.00           320         NC. 009486         Clavilacure micligamentsis subspt. michigamentsis NCPPW         72.84         571         Nc. 014810         Heliciochacter elist ATCC 49179         44.89           221         NC. 009486         Bradyntizobium sp. BTA11         65.50         473         Nc. 014921         Teylorella equiagmenimenta M64         27.21           233         NC. 009486         Thermotoga peterophila RKU-1         46.26         573         Nc. 014921         Mecubiant miceri brown bisermina M64         27.21           235         NC. 009512         Bracella ovis ATCC 25840         58.27         756         Nc. 014921         Mecubiant miceri brown bisermina WSM1271         70.32           237         NC. 009523         Rose offictous sp. RS         70.73         Nc. 014921         Nc. 014921         Paramotosia assumensis II-1         52.24           238         NC. 009613         Besterotides virgianus ATCC 5882		_	_			_		
28. N. 2.009446   N. 2.009450   Candidates Vesicomynocius columnit A							* *	
29 NC_009465   Clavibacter michigamensis subsp. michigamensis NCPPB   7.34   4.80			*				• •	
23   N.C.,00984						_		
231         NC,009484         Acadiphilium cypumn IF-S         68.53         572         NC,014912         Tayloretta equigentialis MCE9         37.78           223         NC,009486         Thermotoga petrophila RRU-1         46.26         573         NC,014923         Mesonizobium ciceri biovar binermalas WSM1271         63.37           234         NC,009494         Legionella peneumophila str. Corby         39.09         575         NC,014923         Mesonizobium ciceri biovar binermalas WSM1271         63.37           235         NC,009524         Beacle nois ATCC 25840         85.27         675         NC,014924         Pseudoamahomonas suwomensis 11-1         52.24           256         NC,009524         Rosoiflecus ap RS-1         60.93         577         NC,014931         Variovorax pandosus EFS         67.05           237         NC,009513         Flavobacterium psychrophilum IIPO286         33.24         45.78         NC,014938         Deinencecus maricopensis DSM 21211         70.23           240         NC,009613         Bacteroides vidgamas ATCC 8482         43.18         580         NC,014958         Deinencecus maricopensis DSM 21211         70.23           241         NC,009615         Parabacterioides vidgamas ATCC 8482         43.18         580         NC,014944         Themosecus accordance and accordanc	230	NC_009480		72.84	571	NC_014810	Helicobacter felis ATCC 49179	44.89
22 NC_009485   Bradyrhizobium p. BTAil   65.50   573 NC_014921   Mycoplasma fermentuns M64   27.21   233 NC_009486   Thermotoga pertophila RRU-1   46.26   574 NC_014923   Mesonizobium cicert biovar bisernals wSM1271   63.37 NC_009593   Bruella ovis ATCC 25840   58.27   576 NC_014924   Pseudoannona suvomensis II-1   70.46	231	NC 009484		68.53	572	NC 014914	Taylorella equigenitalis MCE9	37.78
234         NC_009494         Leginnella preumophila str. Corby         39.09         575         NC_014926         Peeudoxambomonas suwomensis 11-1         70.46           235         NC_009523         Roseiflexus sp. RS-1         60.93         577         NC_014926         Thermovibrio ammonificam HB-1         52.24           237         NC_009523         Psychrobacter sp. PRwf-1         46.41         578         NC_014932         Baronacella charidgener 73         37.38           238         NC_009613         Flavobacterium psychrophilum IIPO286         33.24         NC_009616         Psychrobacter sp. PRwf-1         46.41         578         NC_014952         Baronacella charidgener 73         37.38           240         NC_009615         Parabacteroides stitasonis ATCC 8503         46.01         581         NC_014960         Anaerolinea thermophila UN-1         5.37           241         NC_009616         Thermosipho melanesinsins Bi429         31.23         382         NC_015161         Thermus scotodecus 8A-01         65.12           242         NC_009616         Thermosipho melanesinsins Bi429         31.23         582         NC_015164         Welcobacterium restaceum St.B.037         70.39           242         NC_009636         Sinnohizobium medicase WSM419         62.14         458         NC_01		_						
225         NC_009505         Brucella ovis ATCC 2840         58.27         576         NC_01923         Thermovibrio ammonificans HB-1         32.24           236         NC_009524         Psychrobacters psychrobacters psychrobacterium psychrophilum JPU256         32.4         578         NC_0194932         Banoella clarifagiea 73         37.38           238         NC_009613         Flavobacterium psychrophilum JPU256         32.4         579         NC_019498         Deinococcus maricopensis DSM 21211         70.23           240         NC_009615         Parabacteriudes distasonis ATCC 8482         43.18         8         NC_019498         Deinococcus maricopensis DSM 21211         70.23           241         NC_009615         Parabacteriudes distasonis ATCC 8492         43.18         8         NC_019409         Anaerolinea thermophila UNI-1         54.37           241         NC_009616         Thermosipho melanesiensis BI429         31.23         582         NC_019521         Mc_nobacterium states must MB037         70.39           242         NC_009633         Alkaliphilus metalliredigens QYMF         37.43         588         NC_015162         Sphaerochaetat globos atr. Buddy         49.41           245         NC_009665         Sinothizobium melicae wSM419         62.14         585         NC_015162         Sphaeroch	233	NC_009486	Thermotoga petrophila RKU-1	46.26	574	NC_014923	Mesorhizobium ciceri biovar biserrulae WSM1271	63.37
25         NC_009523         Roseiflexus sp. Rs-1         60.93         577         NC_019324         Variovorax paradoxus EPS         67.05           277         NC_009524         Psychrobacter sp. PRwF-1         46.41         578         NC_014932         Bartonella clarridgeiae 73         37.38           238         NC_009613         Flavobacterium psychrophilum IPI02/86         33.24         579         NC_014958         Delicoccessus maricopeans ESM2 1211         70.23           239         NC_009614         Bacteroides vulgatus ATCC 8482         43.18         850         NC_014969         Anaerolinea thermophila UN-1         45.37           241         NC_009616         Thermosipho melianesiensis BH29         31.23         581         NC_014974         Thermus scooductus SA-01         65.12           242         NC_009616         Clostridium beligreins CVIII         37.43         581         NC_015125         Microbacterium testaceum StLB037         70.39           242         NC_009636         Sinorhizobium medicae WSM419         62.14         585         NC_015152         Microbacterium testaceum StLB037         70.39           245         NC_009655         Acinobactillus unicolomosa peruginosa PA7         67.11         588         NC_015156         Odorbacteria perulachius SDM 28017         47.55								
237         NC_009524         Psychrobacterum psychrophilum JIP0286         33.24         S78         NC_014932         Bartonella clarridgeine 73         37.38           238         NC_009614         Bacteriodes valgaus ATCE 8482         43.18         580         NC_014960         Anaerolinea thermophila UNI-1         54.37           240         NC_009615         Parnhacteroides distasonis ATCE 6803         46.01         581         NC_014932         Delmococcus maricopensis DSM 21211         70.23           242         NC_009616         Thermosipho melanesiensis BI429         31.23         282         NC_015061         Ranhealte psychocolicus AO         65.12           242         NC_009617         Closifidium beiginenis NCMB 8052         30.91         583         NC_015161         Thermosipho melanesiensis BI429         31.23           243         NC_009633         Alkaliphilus mediliredigens QVMF         37.43         584         NC_015162         Microbacterium testaceum StLB037         70.39           245         NC_009653         Actinobacillus succinogenes 1302         45.93         86         NC_015162         Odoribacter spharchicus DSM 220712         44.35           246         NC_009665         Actinobacillus succinogenes 1302         45.93         587         NC_015161         Deinococcus parcholicus DSM 220712<		_						
238         NC_009613         Flavobacterium psychrophilum JIP0286         33.24         579         NC_014960         Deinococcus maricopensis DSM 21211         70.23           239         NC_009615         Basteroides vulgatus ATCC 8482         43.18         580         NC_014960         Anneroince thermophila UNI-1         54.37           241         NC_009616         Thermosipho melanesiensis B1429         31.23         582         NC_019061         Rahnella sp. Y9602         53.52           242         NC_009613         Alkaliphilas mealliredigens QYMF         37.43         588         NC_015125         Microbacterium testaceum SLB.037         70.39           242         NC_009633         Alkaliphilas mealliredigens QYMF         37.43         588         NC_015125         Microbacterium testaceum SLB.037         70.39           243         NC_009636         Sinorhizobium medicae WSM419         62.14         585         NC_015162         Odorbacter splanchicus DSM 20712         44.35           245         NC_009655         Actinobacillus sucionegnes 1307         45.93         587         NC_015161         Odorbacter splanchicus DSM 20712         44.35           248         NC_009665         Pseudomonacterium sp. Marseille         55.15         588         NC_015161         Deinococcus proteolyticus MRP         6			_				-	
240         NC_009615         Parabacteroides distasonis ATCC 8503         46.01         581         NC_019616         Thermosipho melanesiensis Bla29         31.23         S82         NC_015051         Rahnella sp. Y9602         53.52           242         NC_009637         Clostidium beigiernicki NCIMB 8052         30.91         583         NC_015125         Microbacterium testaceum SLB037         70.39           243         NC_009633         Alkaliphilus metalliredigens QYMF         37.43         584         NC_015125         Shapencchaeta globosa str. Buddy         49.41           243         NC_009654         Samo pNWYL1         43.23         586         NC_015162         Sphapenchaeta globosa str. Buddy         49.41           246         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus protoolyticus MRP         66.90           247         NC_009665         Pseudomonas aeruginosa PA7         67.11         588         NC_015161         Bacteroides salanitronis DSM 18170         47.83           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015164         Bacteroides salanitronis DSM 18170         47.83           251         NC_009663         Sulfurovum sp. NBC37-1         44.72         591 </td <td></td> <td></td> <td>• •</td> <td></td> <td></td> <td></td> <td>-</td> <td></td>			• •				-	
241         NC_009616         Thermosipho melanesiensis BI429         31.23         582         NC_0195125         Microbacterium testaceum StLB037         70.39           242         NC_009633         Alkaliphilus metalliredigens (VMF         37.43         588         NC_015144         Weckseld virosa DSM 16922         36.88           244         NC_009636         Sinorhizobium medicae WSM419         62.14         585         NC_015160         Odoribacter: splanchnicus DSM 20712         44.35           245         NC_009654         Marinomonas sp. MWYL1         43.32         586         NC_015160         Odoribacter: splanchnicus DSM 20712         44.35           247         NC_009656         Actionbacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus proteolyticus MRP         66.90           247         NC_009659         Janthinobacterium sp. Marseille         55.15         589         NC_015167         Cellulophaga lytica DSM 7489         32.65           248         NC_009663         Sulfarrovum sp. NBC37-1         44.72         591         NC_015166         Acidiphilium multivorum Allu301         68.22           250         NC_009664         Kinecocccus radiotolerans RSR30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus Allum Allum	239	NC_009614	Bacteroides vulgatus ATCC 8482	43.18	580	NC_014960	Anaerolinea thermophila UNI-1	54.37
242         NC_009617         Clostridium beijerinckii NCIMB 8052         30.91         583         NC_015125         Microbacterium testaceum Stl.B037         70.39           243         NC_009636         Alkaliphilus metalliredigene QYMF         37.43         584         NC_015144         Weeksella virosa DSM 16922         36.68           244         NC_009656         Sinorhizzobium medicae WSM419         62.14         585         NC_015160         Odoribacter splanchnicus DSM 220712         44.35           246         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus proteolyticus MRP         66.90           249         NC_009655         Pseudomonas aeruginosa PA7         67.11         588         NC_015161         Bacteroides salanitronis DSM 18170         47.83           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Acidiphillium multivorum AlU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.18           251         NC_009664         Kinecococcus machitolerium Aluxori Machitolerium Aluxori Machitolerium Aluxori Machitolerium Aluxori Machitolerium Aluxori Machitolerium Aluxori M								
243         NC_009633         Alkaliphilus metalliredigens QYMF         37.43         584         NC_015142         Weeksella virosa DSM 16922         36.68           244         NC_009636         Sinorhizobium medicae WSM419         62.14         585         NC_015152         Sphaencchaeta globosa str. Buddy         49.41           245         NC_009654         Marinomonas p. MWYL1         43.32         586         NC_015160         Odoribacter splanchnicus DSM 220712         44.35           246         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus protodyticus MRP         66.90           247         NC_009656         Pseudomonas aeruginosa PA7         67.11         588         NC_015161         Bacteroides salanitronis DSM 18170         47.83           248         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015166         Actidiphilium multivorum AlU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015214         Lactobacillus acidophilus 30SC         38.88           251         NC_009664         Kinecoccus radiotolerans SK830216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus subsp. gallolyticus subsp			_					
244         NC_009636         Sinorhizobium medicae WSM419         62.14         585         NC_015152         Sphaerochaeta globosa str. Buddy         49.41           245         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015160         Odoribacter splanchnicus DSM 220712         44.35           247         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus proteolyticus MRP         66.90           248         NC_009659         Janthinobacterium sp. Marseille         55.15         588         NC_015164         Bacteroides salanitronis DSM 18170         47.83           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Acidiphilium multivorum AltJ301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015186         Acidiphilium multivorum AltJ301         68.22           251         NC_009664         Kineococcus radiotolerams SRS30216         74.50         592         NC_015215         Streptococcus gailolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_01527         Sphingobacterium sp.								
246         NC_009655         Actinobacillus succinogenes 130Z         45.93         587         NC_015161         Deinococcus proteolyticus MRP         66.90           247         NC_009656         Pseudomonas aeruginosa PA7         67.11         588         NC_015164         Bacteroides salanitronis DSM 18170         47.83           248         NC_0096659         Jamhinobacterium sp. Marseille         55.15         589         NC_015167         Cellulophaga lytica DSM 7489         32.65           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Accidiphilium multivorum AIU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           251         NC_009664         Kinecococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         599         NC_015216         Marinomonas mediterrane MMB-1         44.87           255         NC_009716         Clostridium kluyveri DSM 555         32.78         596         NC_015211         Prevotell								
247         NC_009656         Pseudomonas aeruginosa PA7         67.11         588         NC_015164         Bacteroides salanitronis DSM 18170         47.83           248         NC_009659         Janthinobacterium sp. Marseille         55.15         589         NC_015167         Cellulophaga lytica DSM 7489         32.65           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Acidiphilium multivorum AlU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015215         Lactobacillus acidophilus 30SC         38.88           251         NC_009664         Kineococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_015276         Marinomonas mediterranea MMB-1         44.87           253         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus gallolyticus atCro BAA-2069         38.19           255         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         595         NC_015311         Prevolen denticola F02			Marinomonas sp. MWYL1			NC_015160	Odoribacter splanchnicus DSM 220712	
248         NC_009659         Janthinobacterium p. Marseille         55.15         589         NC_015167         Cellulophaga lytica DSM 7489         32.65           249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Acidiphilium multivorum AlU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015215         Inchobacillus acidophilus 30SC         38.88           251         NC_009664         Kineococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_015276         Marinomonas mediteranea MMB-1         44.87           253         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015217         Sphingobacterium sp. 21         41.87           254         NC_009716         Clostridium kluyveri DSM 555         32.78         595         NC_015311         Prevotella denticola F0289         51.79           255         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015311         Hippea maritima DSM 10411         37.48								
249         NC_009662         Nitratiruptor sp. SB155-2         39.81         590         NC_015186         Acidiphilium multivorum AIU301         68.22           250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015214         Lactobacillus acidophilus 30SC         38.88           251         NC_009664         Kineococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_015277         Sphingobacterium sp. 21         44.87           253         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus oralis Uo5         41.89           255         NC_009706         Clostridium kluyveri DSM 555         32.78         596         NC_015211         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           258         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         3			-					
250         NC_009663         Sulfurovum sp. NBC37-1         44.72         591         NC_015214         Lactobacillus acidophilus 30SC         38.88           251         NC_009664         Kineococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_015276         Marinomonas mediterranea MMB-1         44.87           253         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus oralis Uo5         41.87           255         NC_009706         Clostridium kluyveri DSM 555         32.78         596         NC_015311         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015385         Treponema succinifactiens DSM 14884 </td <td></td> <td></td> <td>_</td> <td></td> <td></td> <td></td> <td></td> <td></td>			_					
251         NC_009664         Kineococcus radiotolerans SRS30216         74.50         592         NC_015215         Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069         38.19           252         NC_009667         Ochrobactrum anthropi ATCC 49188         57.07         593         NC_015276         Marinomonas mediterranea MMB-1         44.87           253         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus oralis Uo5         41.89           255         NC_009716         Clostridium kluyveri DSM 555         32.78         596         NC_015311         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacter ium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009720         Xanthobacter autotrophicus Py         67.91         600         NC_015387         Marinithermus hydr							-	
253         NC_009674         Bacillus cytotoxicus NVH 391-98         36.39         594         NC_015277         Sphingobacterium sp. 21         41.87           254         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus oralis Uo5         41.89           255         NC_009706         Clostridium kluyveri DSM 555         32.78         596         NC_015311         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015385         Treponema succinifaciens DSM 2489         39.55           259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009787         Roseiflexus castenholzii DSM 13941         60.94         602         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25	251		Kineococcus radiotolerans SRS30216				Streptococcus gallolyticus subsp. gallolyticus ATCC BAA-2069	
254         NC_009675         Anaeromyxobacter sp. Fw109-5         73.54         595         NC_015291         Streptococcus oralis Uo5         41.89           255         NC_009706         Clostridium kluyveri DSM 555         32.78         596         NC_015311         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015385         Treponema succinifaciens DSM 2489         39.55           259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009727         Coxiella burnetii Dugway 51108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomona mendocina NK-01 <t< td=""><td></td><td></td><td>_</td><td></td><td></td><td></td><td></td><td></td></t<>			_					
255         NC_009706         Clostridium kluyveri DSM 555         32.78         596         NC_015311         Prevotella denticola F0289         51.79           256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015385         Treponema succinifaciens DSM 2489         39.55           259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009727         Coxiella burnetii Dugway 51108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           265         NC_009828         Streptococcus gordonii str. Challis substr. CHII         41.40         604         NC_015422         Alicycliphilus deni								
256         NC_009714         Campylobacter hominis ATCC BAA-381         32.97         597         NC_015312         Pseudonocardia dioxanivorans CB1190         73.41           257         NC_009718         Fervidobacterium nodosum Rt17-B1         35.23         598         NC_015318         Hippea maritima DSM 10411         37.48           258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015385         Treponema succinifaciens DSM 2489         39.55           259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009727         Coxiella burnetii Dugway 51108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009767         Roseiflexus castenholzii DSM 13941         60.94         602         NC_015391         Carnobacterium sp. 17-4         35.91           262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           265         NC_009828         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus de								
258         NC_009719         Parvibaculum lavamentivorans DS-1         62.74         599         NC_015385         Treponema succinifaciens DSM 2489         39.55           259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009727         Coxiella burnetii Dugway 5J108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009767         Roseiflexus castenholzii DSM 13941         60.94         602         NC_015391         Carnobacterium sp. 17-4         35.91           262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           NC_009785         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015422         Alcycliphilus denitrificans K601         68.72           265         NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929								
259         NC_009720         Xanthobacter autotrophicus Py2         67.91         600         NC_015387         Marinithermus hydrothermalis DSM 14884         68.53           260         NC_009727         Coxiella burnetii Dugway 5J108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009767         Roseiflexus castenholzii DSM 13941         60.94         602         NC_015391         Carnobacterium sp. 17-4         35.91           262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           263         NC_009785         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015424         Aeromonas veronii B565         60.05           NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015434         Verrucosispora maris AB-18-032         71.28								
260         NC_009727         Coxiella burnetii Dugway 5J108-111         42.83         601         NC_015388         Desulfobacca acetoxidans DSM 11109         54.25           261         NC_009767         Roseiflexus castenholzii DSM 13941         60.94         602         NC_015391         Carnobacterium sp. 17-4         35.91           262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           263         NC_009785         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015424         Aeromonas veronii B565         60.05           NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61								
261         NC_009767         Roseiflexus castenholzii DSM 13941         60.94         60.2         NC_015391         Carnobacterium sp. 17-4         35.91           262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           263         NC_009785         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015424         Aeromonas veronii B565         60.05           265         NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58 <td></td> <td>_</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>		_						
262         NC_009783         Vibrio campbellii ATCC BAA-1116         46.33         603         NC_015410         Pseudomonas mendocina NK-01         63.26           263         NC_009785         Streptococcus gordonii str. Challis substr. CHI         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015424         Aeromonas veronii B565         60.05           265         NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71			* *					
263         NC_009785         Streptococcus gordonii str. Challis substr. CH1         41.40         604         NC_015422         Alicycliphilus denitrificans K601         68.72           264         NC_009802         Campylobacter concisus 13826         40.20         605         NC_015424         Aeromonas veronii B565         60.05           265         NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88								
265         NC_009828         Thermotoga lettingae TMO         38.71         606         NC_015428         Lactobacillus buchneri NRRL B-30929         45.31           266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88		NC_009785	_			NC_015422		
266         NC_009831         Shewanella sediminis HAW-EB3         47.31         607         NC_015434         Verrucosispora maris AB-18-032         71.28           267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusillimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88								
267         NC_009850         Arcobacter butzleri RM4018         27.19         608         NC_015458         Pusilimonas sp. T7-7         57.61           268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88		_						
268         NC_009879         Rickettsia canadensis str. McKiel         31.85         609         NC_015460         Gallibacterium anatis UMN179         40.58           269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88								
269         NC_009881         Rickettsia akari str. Hartford         32.79         610         NC_015470         Chlamydia psittaci 6BC         39.71           270         NC_009900         Rickettsia massiliae MTU5         32.98         611         NC_015496         Krokinobacter sp. 4H-3-7-5         37.88							•	
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271 NC_009901         Shewanella pealeana ATCC 700345         45.72         612 NC_015497         Glaciecola sp. 4H-3-7+YE-5         44.92								
	271	NC_009901	Shewanella pealeana ATCC 700345	45.72	612	NC_015497	Glaciecola sp. 4H-3-7+YE-5	44.92

SI	Accession	Taxa_Name	(G+C)%
No 272	No. NC 009921	Frankia sp. EAN1pec	71.59
273	NC_009922	Alkaliphilus oremlandii OhILAs	37.09
274	NC_009937	Azorhizobium caulinodans ORS 571	67.66
275	NC_009943	Desulfococcus oleovorans Hxd3	57.24
276	NC_009952	Dinoroseobacter shibae DFL 12	66.35
277	NC_009953	Salinispora arenicola CNS-205	69.86
278	NC_009997	Shewanella baltica OS195	47.37
279	NC_010001	Clostridium phytofermentans ISDg	36.29
280	NC_010002	Delftia acidovorans SPH-1	67.25
281	NC_010003	Petrotoga mobilis SJ95	34.20
	NC_010084	Burkholderia multivorans ATCC 17616	67.31
283 284	NC_010103	Brucella canis ATCC 23365 Neisseria meningitidis 053442	58.23
285	NC_010120 NC_010125	Gluconacetobacter diazotrophicus PAl 5	53.18 67.06
	NC_010123	Bartonella tribocorum CIP 105476	40.73
287	NC_010162	Sorangium cellulosum So ce56	71.59
288	NC_010170	Bordetella petrii DSM 12804	65.98
289	NC_010184	Bacillus weihenstephanensis KBAB4	36.11
290	NC_010263	Rickettsia rickettsii str. Iowa	32.86
291	NC_010320	Thermoanaerobacter sp. X514	34.67
292	NC_010321	Thermoanaerobacter pseudethanolicus ATCC 33223	34.64
293	NC_010322	Pseudomonas putida GB-1	62.69
294	NC_010334	Shewanella halifaxensis HAW-EB4	45.70
295	NC_010336	Francisella philomiragia subsp. philomiragia ATCC 25017	33.07
296	NC_010337	Heliobacterium modesticaldum Ice1	58.01
297	NC_010338	Caulobacter sp. K31	68.13
298 299	NC_010376 NC_010397	Finegoldia magna ATCC 29328  Mycobacterium abscessus	32.56 64.50
300	NC_010397 NC_010424	Mycobacterium abscessus  Candidatus Desulforudis audaxviator MP104C	62.05
301	NC_010424	Yersinia pseudotuberculosis YPIII	48.75
	NC_010403	Leuconostoc citreum KM20	39.52
303	NC_010483	Thermotoga sp. RQ2	46.34
304	NC_010505	Methylobacterium radiotolerans JCM 2831	71.82
305	NC_010506	Shewanella woodyi ATCC 51908	44.69
306	NC_010511	Methylobacterium sp. 4-46	72.09
307	NC_010519	Haemophilus somnus 2336	37.97
308	NC_010524	Leptothrix cholodnii SP-6	69.30
309	NC_010528	Cupriavidus taiwanensis LMG 19424	68.15
310	NC_010545	Corynebacterium urealyticum DSM 7109	64.85
311	NC_010554	Proteus mirabilis HI4320 Exiguobacterium sibiricum 255-15	40.15
312	NC_010556 NC_010571	Exiguobacterium sibiricum 255-15 Opitutus terrae PB90-1	48.50 65.47
314	NC_010571 NC_010572	Streptomyces griseus subsp. griseus NBRC 13350	72.41
315	NC_010572 NC_010581	Beijerinckia indica subsp. indica ATCC 9039	58.26
316	NC_010602	Leptospira biflexa serovar Patoc strain 'Patoc 1 (Paris)'	38.96
317	NC_010610	Lactobacillus fermentum IFO 3956	53.02
318	NC_010612	Mycobacterium marinum M	66.08
319	NC_010617	Kocuria rhizophila DC2201	71.15
320	NC_010622	Burkholderia phymatum STM815	63.39
321	NC_010644	Elusimicrobium minutum Pei191	40.69
	NC_010645	Bordetella avium 197N	62.10
323	NC_010658	Shigella boydii CDC 3083-94	52.38
324	NC_010673	Borrelia hermsii HS1	30.07
325	NC_010681	Burkholderia phytofirmans PsJN	63.10
326 327	NC_010682	Ralstonia pickettii 12J Erwinia tasmaniensis Et1/99	64.01
328	NC_010694 NC_010717	Erwinia tasmaniensis Et1/99 Xanthomonas oryzae pv. oryzae PXO99A	54.91 64.28
329	NC_010717 NC_010725	Methylobacterium populi BJ001	70.01
330	NC_010723	Sulfurihydrogenibium sp. YO3AOP1	32.03
331	NC_010793	Orientia tsutsugamushi str. Ikeda	31.54
332	NC_010794	Methylacidiphilum infernorum V4	45.85
333	NC_010803	Chlorobium limicola DSM 245	52.23
334	NC_010814	Geobacter lovleyi SZ	55.35
335	NC_010939	Actinobacillus pleuropneumoniae serovar 7 str. AP76	42.16
336	NC_010943	Stenotrophomonas maltophilia K279a	66.71
337	NC_010981	Wolbachia endosymbiont of Culex quinquefasciatus Pel	34.62
	NC_010994	Rhizobium etli CIAT 652	62.44
338			
339	NC_010995	Cellvibrio japonicus Ueda107	52.65
	NC_010995 NC_010999 NC_011004	Cellvibrio japonicus Ueda107 Lactobacillus casei BL23 Rhodopseudomonas palustris TIE-1	52.65 47.00 65.40

Sl	Accession	Taxa_Name	(G+C)%
No	No.		
	NC_015499	Thermodesulfobium narugense DSM 14796	34.30
	NC_015500	Treponema brennaborense DSM 12168	52.08 53.51
	NC_015501 NC_015510	Porphyromonas asaccharolytica DSM 20707 Haliscomenobacter hydrossis DSM 1100	48.00
	NC_015519	Tepidanaerobacter acetatoxydans Re1	38.31
	NC_015556	Pseudomonas fulva 12-X	64.26
	NC_015558	Streptococcus parauberis KCTC 11537	36.40
620	NC_015559	Marinomonas posidonica IVIA-Po-181	44.96
621	NC_015563	Delftia sp. Cs1-4	67.54
622	NC_015564	Amycolicicoccus subflavus DQS3-9A1	62.52
	NC_015565	Desulfotomaculum carboxydivorans CO-1-SRB	47.70
	NC_015566	Serratia sp. AS12	57.32
	NC_015567	Serratia plymuthica AS9	57.31
	NC_015571 NC_015572	Porphyromonas gingivalis TDC60 Methylomonas methanica MC09	49.36 52.01
	NC_015573	Desulfotomaculum kuznetsovii DSM 6115	55.90
	NC_015581	Thioalkalimicrobium cyclicum ALM1	47.51
	NC_015588	Isoptericola variabilis 225	74.05
631	NC_015589	Desulfotomaculum ruminis DSM 2154	48.74
632	NC_015600	Streptococcus pasteurianus ATCC 43144	38.13
633	NC_015601	Erysipelothrix rhusiopathiae str. Fujisawa	36.73
634	NC_015602	Lactobacillus kefiranofaciens ZW3	38.38
635	NC_015634	Bacillus coagulans 2-6	48.15
	NC_015635	Microlunatus phosphovorus NM-1	67.64
	NC_015638	Lacinutrix sp. 5H-3-7-4	31.35
	NC_015660	Geobacillus thermoglucosidasius C56-YS93	44.84
	NC_015663 NC_015671	Enterobacter aerogenes KCTC 2190	56.12 73.84
641	NC_015671 NC_015672	[Cellvibrio] gilvus ATCC 13127 Flexistipes sinusarabici DSM 4947	38.76
	NC_015675	Mesorhizobium opportunistum WSM2075	63.59
	NC_015677	Ramlibacter tataouinensis TTB310	70.36
	NC_015681	Thermodesulfatator indicus DSM 15286	42.61
645	NC_015682	Thermodesulfobacterium geofontis OPF15	30.67
646	NC_015683	Corynebacterium ulcerans BR-AD22	54.13
647	NC_015687	Clostridium acetobutylicum DSM 1731	31.49
	NC_015696	Francisella sp. TX077308	33.38
	NC_015702	Parachlamydia acanthamoebae UV-7	39.72
	NC_015703	Runella slithyformis DSM 19594	47.70
651 652	NC_015711	Myxococcus fulvus HW-1 Simkania negevensis Z	70.98 42.45
	NC_015713 NC_015725	Mycoplasma bovis Hubei-1	29.75
	NC_015726	Cupriavidus necator N-1	67.16
	NC_015731	Nitrosomonas sp. Is79A3	46.63
	NC_015732	Treponema caldaria DSM 7334	45.94
657	NC_015735	Candidatus Moranella endobia PCIT	45.25
658	NC_015740	Pseudomonas stutzeri ATCC 17588 = LMG 11199	64.52
659	NC_015758	Mycobacterium africanum GM041182	65.87
660	NC_015844	Zobellia galactanivorans	43.55
661	NC_015856	Collimonas fungivorans Ter331	60.91
662	NC_015857	Brucella pinnipedialis B2/94	58.16 67.83
663 664	NC_015859 NC_015866	Corynebacterium variabile DSM 44702 Rickettsia heilongjiangensis 054	67.83 32.85
665	NC_015866 NC_015873	Megasphaera elsdenii DSM 20460	54.05
666	NC_015875	Streptococcus pseudopneumoniae IS7493	40.71
667	NC_015913	Candidatus Arthromitus sp. SFB-mouse-Japan	28.34
	NC_015914	Cyclobacterium marinum DSM 745	39.13
669	NC_015930	Lactococcus garvieae ATCC 49156	39.70
670	NC_015942	Acidithiobacillus ferrivorans SS3	57.27
671	NC_015945	Muricauda ruestringensis DSM 13258	42.09
672	NC_015946	Mycoplasma putrefaciens KS1	27.42
673	NC_015953	Streptomyces sp. SirexAA-E	71.99
674	NC_015964	Haemophilus parainfluenzae T3T1	40.18
675	NC_015968	Enterobacter asburiae LF7a Lactobacillus ruminis ATCC 27782	54.88
676 677	NC_015975 NC_015976	Sphingobium sp. SYK-6	44.40 66.00
678	NC_015976 NC_015978	Lactobacillus sanfranciscensis TMW 1.1304	34.87
679	NC_016001	Flavobacterium branchiophilum FL-15	33.91
	NC_016026	Micavibrio aeruginosavorus ARL-13	55.40
681	NC_016041	Glaciecola nitratireducens FR1064	43.00
682	NC_016048	Oscillibacter valericigenes Sjm18-20	54.67
683	NC_016052	Tetragenococcus halophilus NBRC 12172	36.64