

Table 1. Sequences and species used in the study.

Species and strain	Designation of sequence in the respective figure	Length, nt	GenBank accession number
16S rRNA			
Archaea			
<i>Aeropyrum pernix</i>	Aeropyrum	1500	NC_000854
<i>Archaeoglobus fulgidus</i>	Archaeoglobus	1492	NC_000917
<i>Halobacterium</i> sp. NRC-1	Halobacterium	1473	NC_002607
<i>Methanobacterium thermoautotrophicum</i> deltaH	Methanobacterium	1479	NC_000916
<i>Methanococcus jannaschii</i>	Methanococcus	1475	NC_000909
<i>Pyrobaculum aerophilum</i>	Pyrobaculum	2211	NC_003364
<i>Pyrococcus horikoshii</i>	Pyrococcus	1495	NC_000961
<i>Sulfolobus solfataricus</i> P2	Sulfolobus	1496	NC_002754
Chloroflexi (green non-sulfur bacteria)			
<i>Chloroflexus aurantiacus</i>	Chloroflexus	1401	D38365
Proteobacteria (purple bacteria and relatives)			
<i>Agrobacterium tumefaciens</i> strain C58	Agrobacterium	1492	AE009201
<i>Burkholderia</i> sp. LMG 16307	Burkholderia	1472	AF215706
<i>Mesorhizobium loti</i> strain LMG 17826t2	Mesorhizobium	1426	AJ315352
<i>Pseudomonas fluorescens</i>	Pseudomonas	1527	D84013
<i>Rhodobacter sphaeroides</i> strain CGA009	Rhodobacter	1388	D16424
<i>Rhodopseudomonas palustris</i>	Rhodopseudomonas	1429	AY084079
<i>Rhodospirillum rubrum</i>	Rhodospirillum	1406	D30778
<i>Sinorhizobium meliloti</i>	Sinorhizobium	1484	15963753
<i>Xanthomonas campestris</i> pv. <i>campestris</i> str. ATCC 33913	Xanthomonas	1547	NC_003902
Thermotogae			
<i>Thermotoga maritima</i>	Thermotoga	1559	NC_000853
Cyanobacteria (blue-green algae)			
<i>Anabaena</i> sp. PCC7120	Anabaena	1489	17131110
<i>Calothrix parietina</i>	Calothrix	1110	AF334696
<i>Chlorogloeopsis</i> sp.	Chlorogloeopsis	1461	X68780
<i>Chroococciopsis thermalis</i>	Chroococciopsis	1443	AB039005
<i>Cyanothece</i> sp. PCC8801	Cyanothece	1446	AF296873
<i>Cylindrospermum</i> sp. PCC7417	Cylindrospermum	1444	AJ133163
<i>Geitlerinema</i> sp. PCC7105	Geitlerinema	1435	AB039010
<i>Gloeobacter violaceus</i>	Gloeobacter	1407	AF132790
<i>Nodularia spumigena</i> PCC73104	Nodularia	1517	AF268023

<i>Nostoc punctiforme</i> PCC73102	Nostoc	1410	AF027655
<i>Prochlorococcus marinus</i> MED4	Prochlorococcus	1147	AF001466
<i>Scytonema hyalinum</i>	Scytonema	1111	AF334700
<i>Synechococcus</i> sp. PCC7942	Synechococcus	1411	AF132930
<i>Synechocystis</i> sp. PCC6803	Synechocystis	1489	1653715
<i>Thermosynechococcus elongatus</i>	Elongatus	1491	AP005376
<i>Xenococcus</i> sp. PCC7305	Xenococcus	1408	AF132783
kaiC			
Archaea			
<i>Aeropyrum pernix</i>	Aeropyrum	756	NC_000854
<i>Archaeoglobus fulgidus</i>	Archaeoglobus 1 Archaeoglobus 2	1332 681	NC_000917
<i>Halobacterium</i> sp. NRC-1	Halobacterium	657	NC_002607
<i>Methanobacterium thermoautotrophicum</i> deltaH	<i>Methanobacterium</i>	1413	NC_000916
<i>Methanococcus jannaschii</i>	Methanococcus	729	NC_000909
<i>Pyrobaculum aerophilum</i>	Pyrobaculum 1 Pyrobaculum 2	846 846	NC_003364
<i>Pyrococcus horikoshii</i>	Pyrococcus	1452	NC_000961
<i>Sulfolobus solfataricus</i> P2	Sulfolobus 1 Sulfolobus 2	843 789	NC_002754
Chloroflexi (green non-sulfur bacteria)			
<i>Chloroflexus aurantiacus</i>	<i>Chloroflexus</i> 1 Chloroflexus 2	1536 1449	JGI*
Proteobacteria (purple bacteria and relatives)			
<i>Agrobacterium tumefaciens</i> C58 plasmid AT	Agrobacterium 1 Agrobacterium 2	1515 1509	NC_003064
<i>Burkholderia fungorum</i>	Burkholderia	1518	NC_003371
<i>Mesorhizobium loti</i> plasmid pMLb	Mesorhizobium 1	1530	NC_002682
<i>Mesorhizobium loti</i>	Mesorhizobium 2	1488	NC_002678
<i>Pseudomonas fluorescens</i>	Pseudomonas	1518	JGI
<i>Rhodobacter sphaeroides</i> CGA009	<i>Rhodobacter</i>	1713	JGI
<i>Rhodopseudomonas palustris</i> CGA009	<i>Rhodopseudomonas</i>	1650 [†]	JGI
<i>Rhodospirillum rubrum</i> ATCC 11170	<i>Rhodospirillum</i> 1	1683	JGI
<i>Sinorhizobium meliloti</i> plasmid pSymA	Sinorhizobium	1446	NC_003037
<i>Xanthomonas campestris</i> pv. <i>campestris</i> ATCC 33913	Xanthomonas	1542	AE012267
Thermotogae			
<i>Thermotoga maritima</i>	Thermotoga	762	NC_000853
Cyanobacteria (blue-green algae)			

<i>Anabaena</i> sp. PCC7120	<i>Anabaena</i>	1560	AP003591
<i>Calothrix parietina</i>	<i>Calothrix</i>	594 [†]	AF239754
<i>Chlorogloeopsis fritschii</i>	<i>Chlorogloeopsis</i>	594 [†]	AF222603
<i>Chroococciopsis thermalis</i>	<i>Chroococciopsis</i>	594 [†]	AF239752
<i>Cyanothece</i> sp. PCC 8801	<i>Cyanothece</i>	1566	AF442204
<i>Cylindrospermum</i> sp. PCC7417	<i>Cylindrospermum</i>	807 [†]	AF222605
<i>Geitlerinema</i> sp. PCC7105	<i>Geitlerinema</i>	594 [†]	AF222604
<i>Gloeobacter violaceus</i>	<i>Gloeobacter</i>	594 [†]	AF239755
<i>Nodularia spumigena</i> PCC73104	<i>Nodularia</i>	807 [†]	AF222602
<i>Nostoc punctiforme</i> ATCC 29133	<i>Nostoc</i>	1563	JGI
<i>Prochlorococcus marinus</i> MED4	<i>Prochlorococcus</i>	1530	JGI
<i>Scytonema</i> sp. PCC 7110	<i>Scytonema</i>	594 [†]	AF222600
<i>Synechococcus</i> sp. PCC7942	<i>Synechococcus</i>	1560	AB010691
<i>Synechocystis</i> sp. PCC6803	<i>Synechocystis 1</i>	1560	AB001339
	<i>Synechocystis 2</i>	1518	16329170
	<i>Synechocystis 3</i>	1707	D90910
<i>Thermosynechococcus elongatus</i>	<i>Elongatus</i>	1557	AP005370
<i>Xenococcus</i> sp. PCC7305	<i>Xenococcus</i>	594 [†]	AF222601
kaiB			
Archaea			
<i>Methanobacterium thermoautotrophicum</i> deltaH	<i>Methanobacterium</i>	255	NC_000916
Chloroflexi (green non-sulfur bacteria)			
<i>Chloroflexus aurantiacus</i>	<i>Chloroflexus 1</i>	306	JGI
	<i>Chloroflexus 2</i>	282	
Proteobacteria (purple bacteria and relatives)			
<i>Rhodobacter sphaeroides</i>	<i>Rhodobacter</i>	270	JGI
<i>Rhodopseudomonas palustris</i> CGA009	<i>Rhodopseudomonas</i>	273 [†]	JGI
<i>Rhodospirillum rubrum</i>	<i>Rhodospirillum 1</i>	291	JGI
	<i>Rhodospirillum 2</i>	308	
Cyanobacteria (blue-green algae)			
<i>Anabaena</i> sp. PCC7120	<i>Anabaena 1</i>	327	AP003591
	<i>Anabaena 2</i>	765	
<i>Cyanothece</i> sp. PCC8801	<i>Cyanothece</i>	315	AF442204
<i>Nostoc punctiforme</i>	<i>Nostoc 1</i>	315	JGI
	<i>Nostoc 2</i>	858	
<i>Prochlorococcus marinus</i> MED4	<i>Prochlorococcus</i>	324	JGI
<i>Synechococcus</i> sp. PCC7942	<i>Synechococcus</i>	309	AB010691
<i>Synechocystis</i> sp. PCC6803	<i>Synechocystis 1</i>	318	D90917
	<i>Synechocystis 2</i>	309	NC_000911
	<i>Synechocystis 3</i>	327	NC_000911

<i>Thermosynechococcus elongatus</i>	<i>Elongatus 1</i>	327	AP005370
	<i>Elongatus 2</i>	804	NC_004113
kaiA			
Cyanobacteria			
<i>Anabaena</i> sp. PCC7120		309	AP003591
<i>Nostoc punctiforme</i>		567	JGI
<i>Synechococcus</i> sp. PCC7942		855	AB010691
<i>Synechocystis</i> sp. PCC6803		900	16329170
<i>Thermosynechococcus elongatus</i>		852	AP005370

* JGI, sequence obtained from Joint Genome Institute database at www.jgi.doe.gov/JGI_microbial/.

† Partial sequence. The italics indicate *kai* sequences residing in the cluster (not shown in the figures).