

A dynamic interaction process between KaiA and KaiC is critical to the cyanobacterial circadian oscillator

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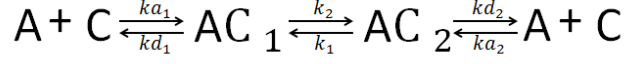
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Supplementary Text S1. The analysis of the KaiA-KaiC binding equilibrium

In this section we will prove that K_{Dfit} , which is the apparent value of the real equilibrium constants of two steps of the KaiA-KaiC binding processes, should be smaller than the apparent values corresponding to K_{Dapp1} and K_{Dapp2} in the following part. For simplicity, A stands for KaiA, and C stands for KaiC (C1 and C2 are two different conformational statuses).



According to the binding processes shown above and the mass action principle, we obtain that:

$$\frac{d[AC_1]}{dt} = ka_1[A][C] - kd_1[AC_1] + k_1[AC_2] - k_2[AC_1] \quad (1)$$

$$\frac{d[AC_2]}{dt} = ka_2[A][C] - kd_2[AC_2] + k_2[AC_1] - k_1[AC_2] \quad (2)$$

Moreover, we suppose that all steps above are rapid and pre-equilibrium processes compared to the phosphorylation processes. Thus, we have

$$\frac{d[AC_1]}{dt} = ka_1[A][C] - kd_1[AC_1] + k_1[AC_2] - k_2[AC_1] = 0 \quad (3)$$

$$\frac{d[AC_2]}{dt} = ka_2[A][C] - kd_2[AC_2] + k_2[AC_1] - k_1[AC_2] = 0 \quad (4)$$

Combining (3) with (4), we may conclude that

$$(ka_1 + ka_2)[A][C] = kd_1[AC_1] + kd_2[AC_2] \quad (5)$$

Suppose that $[AC_{\text{tot}}]$ is the total concentration of complex AC, i.e.

$$[AC_{\text{tot}}] = [AC_1] + [AC_2] \quad (6)$$

Thus, there is a number γ ($0 < \gamma < 1$) such that

$$[AC_1] = \gamma[AC_{\text{tot}}] \quad (7)$$

Combining (5), (6), and (7), we have

$$(ka_1 + ka_2)[A][C] = (\gamma kd_1 + (1 - \gamma)kd_2)[AC_{\text{tot}}] \quad (8)$$

$$(ka_1 + ka_2)[A][C] = \left(kd_1 + \frac{(1 - \gamma)}{\gamma} kd_2 \right) [AC_1] \quad (9)$$

$$(ka_1 + ka_2)[A][C] = \left(\frac{\gamma}{(1 - \gamma)} kd_1 + kd_2 \right) [AC_2] \quad (10)$$

Denoting $K_{\text{Dapp1}} = [A][C]/[AC_1]$, $K_{\text{Dapp2}} = [A][C]/[AC_2]$, and from (8), (9), (10), we obtain

$$\begin{aligned} K_{\text{Dfit}} &= \frac{[A][C]}{[AC_{\text{tot}}]} \\ &= \frac{(\gamma kd_1 + (1 - \gamma)kd_2)}{ka_1 + ka_2}, \end{aligned} \quad (11)$$

$$\begin{aligned}
K_{Dobs1} &= \frac{[A][C]}{[AC_1]} \\
&= \frac{(\gamma kd_1 + (1 - \gamma)kd_2)}{\gamma(ka_1 + ka_2)} ,
\end{aligned} \tag{12}$$

$$\begin{aligned}
K_{Dobs2} &= \frac{[A][C]}{[AC_2]} \\
&= \frac{(\gamma kd_1 + (1 - \gamma)kd_2)}{(1 - \gamma)(ka_1 + ka_2)} ,
\end{aligned} \tag{13}$$

According to (11), (12) and (13), it is obvious that

$$K_{Dfit} = \gamma K_{Dapp1} = (1 - \gamma) K_{Dapp2} \tag{14}$$

Supplementary Figure S1. The sequence analyses of the clock proteins. (A) The sequence alignment of 65 KaiA sequences aligned in MEGA v5.05. The alignments were rendered with STRAP. The secondary structure elements were shown according to the protein structure of *Synechococcus elongates* PCC 7942. (B) The sequence alignment of 65 KaiB sequences aligned in MEGA v5.05. (C) The sequence alignment of 65 KaiC sequences aligned in MEGA v5.05. (D) The phylogenetic tree generated from concatenated KaiA/KaiB/KaiC sequences. (E) The calculated Relative Evolution Rates (RERs) of KaiA residues. (F) The calculated RERs of KaiC residues.

(A)

		$\beta 1$	$\alpha 1$
		10	aaaa
Synechococcus e. PCC 7942	1		
Acaryochloris m. MBIC11017	1		
Acaryochloris sp. CCME 5410	1		
Anabaena v. ATCC 29413	1		
Arthrospira m. CS-328	1		
Arthrospira m. FACHB-438	1		
Arthrospira p. str. Paraca	1		
Crocospaera w. WH 0003	1		
Crocospaera w. WH 8501	1		
Cyanobium sp. PCC 7001	1		
Cyanotheca sp. ATCC 51142	1		
Cyanotheca sp. CCY0110	1		
Cyanotheca sp. PCC 7424	1		
Cyanotheca sp. PCC 7425	1		
Cyanotheca sp. PCC 7822	1		
Cyanotheca sp. PCC 8801	1		
Cyanotheca sp. PCC 8802	1		
Cylindrospermopsis r. CS-505	1		
Fischerella sp. JSC-11	1		
Leptolyngbya b. IAM M-101	1		
Lyngbya sp. PCC 8106	1		
Microcoleus c. PCC 7420	1		
Microcoleus v. FGP-2	1		
Microcystis a. NIES-843	1		
Microcystis a. PCC 7806	1		
Microcystis a. PCC 7820	1		
Microcystis a. PCC 7941	1		
Microcystis a. PCC 9443	1		
Microcystis a. PCC 9701	1		
Microcystis a. PCC 9717	1		
Microcystis a. PCC 9806	1		
Microcystis a. PCC 9807	1		
Microcystis sp. TI-4	1		
Moorea p. 3L	1		
Nodularia s. CCY9414	1		
Nostoc a. 0708	1		
Nostoc c.	1		
Nostoc p. PCC 73102	1		
Nostoc sp. PCC 7120	1		
Nostoc sp. PCC 9709	1		
Oscillatoria sp. PCC 6506	1		
Raphidiopsis b. D9	1		
Synechococcus e. PCC 6301	1		
Synechococcus sp. PCC 7335	1		
Synechococcus sp. BL107	1		
Synechococcus sp. CB0101	1		
Synechococcus sp. CB0205	1		
Synechococcus sp. CC9311	1		
Synechococcus sp. CC9605	1		
Synechococcus sp. CC9902	1		
Synechococcus sp. JA-2-3B	1		
Synechococcus sp. JA-3-3Ab	1		
Synechococcus sp. PCC 7002	1		
Synechococcus sp. RCC307	1		
Synechococcus sp. RS9916	1		
Synechococcus sp. RS9917	1		
Synechococcus sp. WH 5701	1		
Synechococcus sp. WH 7803	1		
Synechococcus sp. WH 7805	1		
Synechococcus sp. WH 8016	1		
Synechococcus sp. WH 8102	1		
Synechococcus sp. WH 8109	1		
Synechocystis sp. PCC 6803	1		
Thermosynechococcus e. BP-1	1		
Trichodesmium e. IMS101	1		
consensus			

logo



Synechococcus e. PCC 7942
 Acaryochloris m. MBIC11017
 Acaryochloris sp. CCME 5410
 Anabaena v. ATCC 29413
 Arthrospira m. CS-328
 Arthrospira m. FACHB-438
 Arthrospira p. str. Paraca
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 Cylandropermopsis r. CS-505
 Fischerella sp. JSC-11
 Leptolyngbya b. IAM M-101
 Lyngbya sp. PCC 8106
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 Synechococcus sp. WH 8109
 Synechocystis sp. PCC 6803
 Thermosynechococcus e. BP-1
 Trichodesmium e. IMS101
 consensus

α3 β4
 70 80 90
 64 RAVVQQLCFEGVVVPAIVY GDRDSEDPD.....EPAKE
 70 PKVVSHPHREAILLPVAVLQVEESVEKTSQPNSAD.....RDPQQD
 70 PKVVSHPHREAILLPVAVLQVEESVEKTSQPNSAD.....RDPQQD
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 55 QGLVNLQQLNLVSIPTLITISDEQSPVVQPSLTEP.....SEN
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 71 LPLFNQLYERGTLLPVVIVE..KGEDPS.....IALANVESPT
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 63 LPLFNQLYEQGTLLPVVIVE..DKVES.....SDNVEPPT
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 63 SPLFNQLYEQGTLLPVVILDSLDQSAS.....TQSENMETAT
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 1
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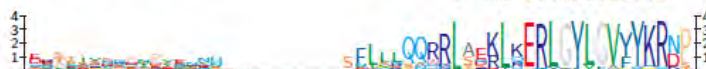
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 Synechococcus sp. WH 8102
 Synechococcus sp. WH 8109
 Synechocystis sp. PCC 6803
 Thermosynechococcus e. BP-1
 Trichodesmium e. IMS101
 consensus

140 150 160 170
 α5 α6
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 153 TLLANADKKTEIHRG...SLMKQQR...SEKLQERLGYLGVYKRN
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 1
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 128 SGSE...SASTPLS...WRPNRLQERLGYLGVYKRD
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 130 DSSSSSRVSN...SDRLQERLGYLGVYKRD
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 128 GGLEASDSHAGESDQ...SNWR...PHRLQERLGYLGVYKRD
 131 EDGS.APSSAES...AS...SAWKSSRLQERLGYLGVYKRD
 131 EDGSDANTQPSG...ES...SAWKSSRLQERLGYLGVYKRD
 131 EDGSAESDQVGGQPEG...SAWKSSRLQERLGYLGVYKRD
 134 DD...DGLASVDK...SRRLQERLGYLGVYKRD
 130 DESSSSSRVSN...SDRLQERLGYLGVYKRD
 136 NQTETIPAPIVDESSQ...SFLLLQQR...ADKLQERLGYLGVYKRN
 134 PHVLFRLP.ALKESS...NVDPQHR...SQRLQERLGYLGVYKRD
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 * * * * *

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		$\alpha 6$	$\alpha 7$	$\alpha 8$	$\alpha 9$		
		180	190	200	210	220	230
Synechococcus e. PCC 7942	176	DRFLRN	PAYESQKHQA	QTSYRE	IVSYFSPNSNL	NQSIDN	VNMAFFAD
Acaryochloris m. MBIC11017	191	QQFHKH	TEEBQTDY	VTIKRDYRD	ILIN	YFRQDSSVNQ	IEDDVTQIEFFAD
Acaryochloris sp. CCME 5410	191	QQFHKH	TEEBQADY	VTIKRDYRD	ILIN	YFRQDSSVNQ	IEDDVTQIEFFAD
Anabaena v. ATCC 29413	1		MTQEV	QQVQ	QQKSDYR	QILSYFTTDKALKEK	IDKFNAYFCAN
Arthrospira m. CS-328	174	KNFLRN	SKAQRKQL	EQKSYRQ	IVSYFANDKNL	NNKIDE	VNMAFFGD
Arthrospira m. FACHB-438	174	KNFLRN	SKAQRKQL	EQKSYRQ	IVSYFANDKNL	NNKIDE	VNMAFFGD
Arthrospira p. str. Paraca	174	KNFLRN	SKAQRKQL	EQKSYRQ	IVSYFANDKNL	NNKIDE	VNMAFFGD
Crocospaera w. WH 0003	164	QYFYRN	SPEKQAL	KEKMTNYRE	ILIN	YFGEDADINQ	AIDQVNQAFFAD
Crocospaera w. WH 8501	191	QYFYRN	SPEKQAL	KEKMTNYRE	ILIN	YFGEDADINQ	AIDQVNQAFFAD
Cyanobium sp. PCC 7001	183	SRFLRN	PLTQNE	RRSERSYR	DLLN	YFRNPSAANQ	ALESVNTAFFSD
Cyanothece sp. ATCC 51142	180	KYFYRN	SPEKREL	KEKMTANYRE	ILIN	YFDEEDSEVNQ	AIDQVNQAFFAD
Cyanothece sp. CCY0110	164	KYFYRN	SPEKREL	KEKMTANYRE	ILIN	YFDEEDSEVNQ	AIDQVNQAFFAD
Cyanothece sp. PCC 7424	197	QYFYRN	SQSEKKE	ESQSGSEYRE	IVSYFQDFFPINET	IDQ	VNRCFFVD
Cyanothece sp. PCC 7425	182	HLFLRH	PEPEKQDF	REFKLDYRE	IVSYFNKENSINP	KIDA	VTKAPFAD
Cyanothece sp. PCC 7822	202	EHFYRN	SSSKKE	IALSSDYRE	IVSYFQDFFPVNQ	AIDQ	VNRCFFAD
Cyanothece sp. PCC 8801	186	KDYFYRN	SPNEKKQL	LAQRAEYRE	ILIN	YFGDNIDVNQ	AIDQVNQAFFSD
Cyanothece sp. PCC 8802	186	KDYFYRN	SPNEKKQL	LAQRAEYRE	ILIN	YFGDNIDVNQ	AIDQVNQAFFSD
Cylindrospermopsis r. CS-505	94	NQQFQD	TDGGRQEL	KKKLDYGR	ILIN	YFSDQNLKTT	IDQESTLFCAN
Fischerella sp. JSC-11	10	QSFQ	TKAQKQAF	QELKSDYR	QILSYFTTDKTLKEK	IDKFNAYFCAN	IPV
Leptolyngbya b. IAM M-101	179	QALFLRN	PPQDQHE	LEALREYRA	ILIN	YFDETALNHR	IDNVMNAFFAD
Lyngbya sp. PCC 8106	170	KNFLRN	PPAQRQEL	EQKLYRQ	IVSYFSGDKALN	NNKIDE	VNMAFFAD
Microcoleus c. PCC 7420	197	QYFRYL	PQNEKSKL	EQKSDYR	IVSYFQSDTAINQ	IDN	VNEAFFAD
Microcoleus v. FGP-2	198	QILKLN	SGPEKQKF	DQKSSYR	DILIN	YFQDQTA	VNNCIDEVNLAPFAD
Microcystis a. NIES-843	172	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 7806	187	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 7820	172	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 7941	218	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 9443	191	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 9701	191	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 9717	191	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 9806	219	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis a. PCC 9807	191	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Microcystis sp. TI-4	191	QLFYRN	APEKKE	RELRADYRE	ILIN	YFQDQYPINQ	AIDQVNNVFFTD
Moorea p. 3L	198	RLFRN	SLNRDEL	LEHRSQYR	QIVSYFAPGDTL	NQDIDH	VEQVFFAD
Nodularia s. CCY9414	1		MTQVRQV	QQKSDYR	ILIN	YFTTDKTLKEE	IDKFNAYFCAN
Nostoc a. 0708	90	QQFQ	KSADQRQEL	LAQKSDYR	ILIN	YFTTDKTLKIA	IDKFNAYFCAN
Nostoc c.	87	QQVFQ	MTQVRQV	QQKSDYR	ILIN	YFTTDKTLKEE	IDKFNAYFCAN
Nostoc p. PCC 73102	87	QQHFNE	NPAQRQGL	ROKLDYSL	ILIN	YFTTDKTLKEE	IDKFNAYFCAN
Nostoc sp. PCC 7120	1		MTQEV	QQVQ	QQKSDYR	QILSYFTTDKALKEK	IDKFNAYFCAN
Nostoc sp. PCC 9709	87	QQHLQE	TPAQRQGL	ROKLDYSL	ILIN	YFTTDKTLKDK	IDKFNAYFCAN
Oscillatoria sp. PCC 6506	218	QYFRN	PSAQRQEL	EQKSNYRE	IVSYFSPDSTL	NNKIDE	VNMAFFAD
Raphidiopsis b. D9	1		TDGGRQED	KKKLDYGR	ILIN	YFSDENLKT	ITIDQESTLFCAN
Synechococcus e. PCC 6301	176	DRFLRN	PAYESQKHQA	QTSYRE	IVSYFSPNSNL	NQSIDN	VNMAFFAD
Synechococcus sp. PCC 7335	194	QNLFLRH	NTDRREEF	DKLRQDYRR	IVSYFADDESIN	QKID	VNMAFFAD
Synechococcus sp. BL107	160	SRFLGS	APDQRREL	ESLHRTYR	DLLSYFGDPAAANQ	ALES	VNTAFFSD
Synechococcus sp. CB0101	163	SLFLRN	PPDQREEL	RSKRSYR	DVLGYFRDPAAANQ	ALES	VHSAFFSD
Synechococcus sp. CB0205	161	SLFLRN	PEPHGE	RQSLQRGYR	DLLSYFKDPAAANQ	ALES	VHSAFFCD
Synechococcus sp. CC9311	201	SRFLAN	PPNPQREL	LQSLQRTYR	DLLSYFRDPAAANQ	ALES	VNTAFFGD
Synechococcus sp. CC9605	162	SRFLGS	PPPERRDL	LQSLQRTYR	DLLSYFRDPAAANQ	ALES	VNTAFFSD
Synechococcus sp. CC9902	160	SRFLNS	APDQRREL	ESLHRTYR	DLLSYFGDPAAANQ	ALES	VNTAFFSD
Synechococcus sp. JA-2-3B	222	ELFYRN	PPHQQAL	MRRRNLYQA	CVEDYFQSPETV	NARIDE	VALAFFAD
Synechococcus sp. JA-3-3Ab	192	ELFYRN	PPDQQAL	MRRRNLYQA	CVEDYFQSPETV	NARIDE	VALAFFAD
Synechococcus sp. PCC 7002	181	KYFYRS	SPEKQEF	REQFVADY	EDIVSYFSGDLPT	NQALDQ	VNQAFFAD
Synechococcus sp. RCC307	164	RLFLRN	PTDQRSEL	DLASERSYR	DVLSYFRDPPAANQ	ALES	VNTAFFCD
Synechococcus sp. RS9916	171	SRFLSR	PPGQEEL	DKSERTYR	DLLSYFRDPPAANQ	ALES	VNTAFFGD
Synechococcus sp. RS9917	171	SRFLAH	AAADQA	EKSLQRTYR	DLLSYFRDPPAANQ	ALES	VNTAFFAD
Synechococcus sp. WH 5701	167	SRFLRN	PSAQRKVL	RSLQRTYR	DLLSYFRDPPAANQ	ALES	VNTAFFSD
Synechococcus sp. WH 7803	168	SRFLAN	PPDQREL	RLRSLQRTYR	DLLSYFRDPPAANQ	ALES	VNTAFFGD
Synechococcus sp. WH 7805	169	SRFLAN	PPPEQREL	QSLQRTYR	DLLSYFRDPPAANQ	ALES	VNTAFFGD
Synechococcus sp. WH 8016	171	SRFLAN	PPHQQREL	QSLQRTYR	DLLSYFRDPPAANQ	ALES	VNTAFFGD
Synechococcus sp. WH 8102	164	SRFLGS	PTPERR	DLRSLQRTYR	DLLSYFRDPAASNQ	ALES	VNTAFFSD
Synechococcus sp. WH 8109	162	SRFLGS	PPGERRD	LQSLQRTYR	DLLSYFRDPAANQ	ALES	VNTAFFSD
Synechocystis sp. PCC 6803	181	SHFYRN	SPQKQ	QYEDSSQYRE	ILIN	YFSGDEGT	VNDLQVNQAFFAD
Thermosynechococcus e. BP-1	175	AFYFRN	SPAQRK	QDELRSIYR	TIVSYFNTDAK	VNERIDE	VSKAPFAD
Trichodesmium e. IMS101	217	TNFI	RNAPGRDKF	YDQKLEYR	HIVKYSQD	NSLNNNID	VNLCFFSD
consensus		*	*	*	*	*	*

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Synechococcus e. PCC 7942
 Acaryochloris m. MBIC11017
 Acaryochloris sp. CCME 5410
 Anabaena v. ATCC 29413
 Arthrospira m. CS-328
 Arthrospira m. FACHB-438
 Arthrospira p. str. Paraca
 Crocosphaera w. WH 0003
 Crocosphaera w. WH 8501
 Cyanobium sp. PCC 7001
 Cyanothece sp. ATCC 51142
 Cyanothece sp. CCY0110
 Cyanothece sp. PCC 7424
 Cyanothece sp. PCC 7425
 Cyanothece sp. PCC 7822
 Cyanothece sp. PCC 8801
 Cyanothece sp. PCC 8802
 Cylandrospermopsis r. CS-505
 Fischerella sp. JSC-11
 Leptolyngbya b. IAM M-101
 Lyngbya sp. PCC 8106
 Microcoleus c. PCC 7420
 Microcoleus v. FGP-2
 Microcystis a. NIES-843
 Microcystis a. PCC 7806
 Microcystis a. PCC 7820
 Microcystis a. PCC 7941
 Microcystis a. PCC 9443
 Microcystis a. PCC 9701
 Microcystis a. PCC 9717
 Microcystis a. PCC 9806
 Microcystis a. PCC 9807
 Microcystis sp. T1-4
 Moorea p. 3L
 Nodularia s. CCY9414
 Nostoc a. 0708
 Nostoc c.
 Nostoc p. PCC 73102
 Nostoc sp. PCC 7120
 Nostoc sp. PCC 9709
 Oscillatoria sp. PCC 6506
 Raphidiopsis b. D9
 Synechococcus e. PCC 6301
 Synechococcus sp. PCC 7335
 Synechococcus sp. BL107
 Synechococcus sp. CB0101
 Synechococcus sp. CB0205
 Synechococcus sp. CC9311
 Synechococcus sp. CC9605
 Synechococcus sp. CC9902
 Synechococcus sp. JA-2-3B
 Synechococcus sp. JA-3-3Ab
 Synechococcus sp. PCC 7002
 Synechococcus sp. RCC307
 Synechococcus sp. RS9916
 Synechococcus sp. RS9917
 Synechococcus sp. WH 5701
 Synechococcus sp. WH 7803
 Synechococcus sp. WH 7805
 Synechococcus sp. WH 8016
 Synechococcus sp. WH 8102
 Synechococcus sp. WH 8109
 Synechocystis sp. PCC 6803
 Thermosynechococcus e. BP-1
 Trichodesmium e. IMS101
 consensus

$\alpha 9$ $\alpha 10$

231 TKVVEIHMELDFAKKKLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRET.
 246 LKLEIHMELDFAKKKLEGRSEIILDYRLTLDTIAHLCEMYRRSIPKTR.
 246 LKLEIHMELDFAKKKLEGRSEIILDYRLTLDTIAHLCEMYRRSIPKTR.
 51 PEIIEIHMELDFFSKQLLEGRGDEITMDYRLTLDTIAHLCEMYRRSIPRES.
 229 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRESW
 229 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRESW
 229 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRESW
 219 SRVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRGDL
 246 SRVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRGDL
 238 TTTVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPPDVP
 235 SRVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRGDL
 219 SRVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRGDL
 252 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 237 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 257 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 241 SKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 241 SKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 149 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 65 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 234 AQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 225 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 252 SKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 253 TQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 227 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 242 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 227 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 273 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 246 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 246 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 246 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 274 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 246 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 246 SQLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 253 SKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 50 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 145 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 142 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 142 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 51 PEIIEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 142 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 273 TQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 50 PQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 231 TKVVEIHMELDFAKKKLEGRSEIILDYRLTLDTIAHLCEMYRRSIPRET.
 249 PKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 215 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 218 NKVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 216 NSVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 256 TQTVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 217 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 215 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 277 AQLVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 247 AQLVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 236 SYLVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 219 TRAVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 226 TRAVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 226 TRAVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 222 TKTVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 223 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 224 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 226 TQTVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 219 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 217 TRIVEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 236 SQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 230 SQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI
 272 TQVLEIHMELDFFSKQLLEGRSEIILDYRLTLDTIAHLCEMYRRSIPREDI



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Synechococcus e. PCC 7942	285	
Acaryochloris m. MBIC11017	300	
Acaryochloris sp. CCMEE 5410	300	
Anabaena v. ATCC 29413	90	
Arthrospira m. CS-328	284	GE.....
Arthrospira m. FACHB-438	284	GE.....
Arthrospira p. str. Paraca	284	GE.....
Crocospaera w. WH 0003	274	LFDLLNQMD.....
Crocospaera w. WH 8501	301	LFDLLNQMD.....
Cyanobium sp. PCC 7001	293	LIRFSAST.....
Cyanothece sp. ATCC 51142	290	LFDLLNQID.....
Cyanothece sp. CCY0110	274	LFDLLNQID.....
Cyanothece sp. PCC 7424	307	PYELLFRID.....
Cyanothece sp. PCC 7425	292	SRET.....
Cyanothece sp. PCC 7822	312	PYELLFRID.....
Cyanothece sp. PCC 8801	296	LFDLLYQID.....
Cyanothece sp. PCC 8802	296	LFDLLYQID.....
Cylindrospermopsis r. CS-505	203	
Fischerella sp. JSC-11	117	
Leptolyngbya b. IAM M-101	288	
Lyngbya sp. PCC 8106	279	
Microcoleus c. PCC 7420	306	
Microcoleus v. FGP-2	308	QGRM.....
Microcystis a. NIES-843	282	PYDLLLGID.....
Microcystis a. PCC 7806	297	PYDLLLGID.....
Microcystis a. PCC 7820	282	PYDLLLGID.....
Microcystis a. PCC 7941	328	PYDLSQMSG.....
Microcystis a. PCC 9443	301	PYDLLLGID.....
Microcystis a. PCC 9701	301	PYDLLLGID.....
Microcystis a. PCC 9717	301	PYDLLLGID.....
Microcystis a. PCC 9806	329	PYDLSQMSG.....
Microcystis a. PCC 9807	301	PYDLLLGID.....
Microcystis sp. T1-4	301	PDDLQMSG.....
Moorea p. 3L	307	
Nodularia s. CCY9414	102	
Nostoc a. 0708	199	
Nostoc c.	194	
Nostoc p. PCC 73102	194	
Nostoc sp. PCC 7120	103	
Nostoc sp. PCC 9709	194	
Oscillatoria sp. PCC 6506	327	
Raphidiopsis b. D9	104	
Synechococcus e. PCC 6301	285	
Synechococcus sp. PCC 7335	303	
Synechococcus sp. BL107	270	LPGEASNRPSVAVDQPMSEELL
Synechococcus sp. CB0101	273	LGAPQDPPEEASQSPSQEVI....
Synechococcus sp. CB0205	271	LAEPFPPGESVG.....
Synechococcus sp. CC9311	311	LVPAGSQRRQLQNSEVSL.....
Synechococcus sp. CC9605	272	LSGTASSRVRPMDQLDASEESS
Synechococcus sp. CC9902	270	LPSEASNRLSVAVDQPMSEELL
Synechococcus sp. JA-2-3B	332	RDP.....
Synechococcus sp. JA-3-3Ab	302	RDP.....
Synechococcus sp. PCC 7002	291	PFELLFRID.....
Synechococcus sp. RCC307	274	GAGARASAGSAVAQQEESPS...
Synechococcus sp. RS9916	281	LARSEQFGEEVVSS.....
Synechococcus sp. RS9917	281	LSASVASQPAGASAVTDQELSS.
Synechococcus sp. WH 5701	277	DKPSEGMVAKIITASQVQS....
Synechococcus sp. WH 7803	278	LAVPSEQRRSLMDSEVSS.....
Synechococcus sp. WH 7805	279	LATSAEYRQALMDSQVSP.....
Synechococcus sp. WH 8016	281	LVPSGSQRRQLQDSEVTL.....
Synechococcus sp. WH 8102	274	LSGLASGRHRREADLPDAPEVSS
Synechococcus sp. WH 8109	229	
Synechocystis sp. PCC 6803	291	PFDVYYQTD.....
Thermosynechococcus e. BP-1	284	
Trichodesmium e. IMS101	326	
consensus		

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(B)

[illegible]

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			$\beta 3$	$\alpha 2$	$\beta 4$		
		50	60	70	80	90	100
Synechococcus e. PCC 7942	48	KNPQLAEEDKILATPTLAK	LPLPVR	IIGDLS	DRE	VLIGLDLLYGE	IQSDDF
Acaryochloris m. MBIC11017	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ISGIM
Acaryochloris sp. CCME 5410	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ISGIM
Anabaena v. ATCC 29413	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETEDW
Arthrospira m. CS-328	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETEDDY
Arthrospira m. FACHB-438	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETEDDY
Arthrospira p. str. Paraca	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETEDDY
Crocospaera w. WH 0003	56	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Crocospaera w. WH 8501	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Cyanobium sp. PCC 7001	48	VNPQLAEEDKILATPTLTK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETDEGF
Cyanothece sp. ATCC 51142	51	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETKE
Cyanothece sp. CCY0110	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Cyanothece sp. PCC 7424	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETIQ
Cyanothece sp. PCC 7425	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETAE
Cyanothece sp. PCC 7822	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETQ
Cyanothece sp. PCC 8801	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Cyanothece sp. PCC 8802	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Cylindrospermopsis r. CS-505	49	KSPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Fischerella sp. JSC-11	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETDEGF
Leptolyngbya b. IAM M-101	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Lyngbya sp. PCC 8106	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETG
Microcoleus c. PCC 7420	53	KSPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETD
Microcoleus v. FGP-2	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETEDL
Microcystis a. NIES-843	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 7806	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 7820	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 7941	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 9443	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 9701	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 9717	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 9806	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis a. PCC 9807	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Microcystis sp. T1-4	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETRE
Moorea p. 3L	49	KSPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETD
Nodularia s. CCY9414	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Nostoc a. 0708	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Nostoc c.	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Nostoc p. PCC 73102	49	KSPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Nostoc sp. PCC 7120	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Nostoc sp. PCC 9709	49	KSPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Oscillatoria sp. PCC 6506	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Raphidiopsis b. D9	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus e. PCC 6301	48	KNPQLAEEDKILATPTLAK	LPLPVR	IIGDLS	DRE	VLIGLDLLYGE	IQSDDF
Synechococcus sp. PCC 7335	49	ESPLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. BL107	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. CB0101	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. CB0205	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. CC9311	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. CC9605	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. CC9902	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. JA-2-3B	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. JA-3-3Ab	48	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. PCC 7002	51	QNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. RCC307	48	KPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. RS9916	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. RS9917	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 5701	28	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 7803	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 7805	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 8016	28	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 8102	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechococcus sp. WH 8109	48	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Synechocystis sp. PCC 6803	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Thermosynechococcus e. BP-1	49	KNPQLAEEDKILATPTLAK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
Trichodesmium e. IMS101	49	KNPQLAEEDKILATPTLSK	LPPPVR	IIGDLS	DRE	VLIGLDLLYGE	ETSE
consensus		*****	*****	*****	*****	*****	*****

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Synechococcus e. PCC 7942	103	
Acaryochloris m. MBIC11017	103	EY.....
Acaryochloris sp. CCMEE 5410	103	EY.....
Anabaena v. ATCC 29413	103	EAQSSL.....
Arthrospira m. CS-328	103	NF.....
Arthrospira m. FACHB-438	103	NF.....
Arthrospira p. str. Paraca	103	NF.....
Crocospaera w. WH 0003	110	DP.....
Crocospaera w. WH 8501	103	DP.....
Cyanobium sp. PCC 7001	103	DALPLASPASEPDASLPVPSEGARSVDGLNPPA
Cyanothece sp. ATCC 51142	105	DP.....
Cyanothece sp. CCY0110	103	DP.....
Cyanothece sp. PCC 7424	103	ELDF.....
Cyanothece sp. PCC 7425	103	EAE.....
Cyanothece sp. PCC 7822	103	ELDF.....
Cyanothece sp. PCC 8801	103	EL.....
Cyanothece sp. PCC 8802	103	EL.....
Cylindrospermopsis r. CS-505	103	DNIDDDHKIEEIN.....
Fischerella sp. JSC-11	104	QTE.....
Leptolyngbya b. IAM M-101	103	DSQ.....
Lyngbya sp. PCC 8106	103	NL.....
Microcoleus c. PCC 7420	107	DF.....
Microcoleus v. FGP-2	103	NS.....
Microcystis a. NIES-843	103	DS.....
Microcystis a. PCC 7806	103	DS.....
Microcystis a. PCC 7820	103	DS.....
Microcystis a. PCC 7941	103	DS.....
Microcystis a. PCC 9443	103	DS.....
Microcystis a. PCC 9701	103	DG.....
Microcystis a. PCC 9717	103	DS.....
Microcystis a. PCC 9806	103	DS.....
Microcystis a. PCC 9807	103	DS.....
Microcystis sp. T1-4	103	DS.....
Moorea p. 3L	103	DF.....
Nodularia s. CCY9414	103	EE.....
Nostoc a. 0708	103	E.....
Nostoc c.	103	EESHNL.....
Nostoc p. PCC 73102	103	EE.....
Nostoc sp. PCC 7120	103	EAQSNL.....
Nostoc sp. PCC 9709	103	E.....
Oscillatoria sp. PCC 6506	103	NLHDRI.....
Raphidiopsis b. D9	103	DNIDDDHKIEEIN.....
Synechococcus e. PCC 6301	103	
Synechococcus sp. PCC 7335	103	N.....
Synechococcus sp. BL107	103	SSFLDDSDDEESAPATSDS.....
Synechococcus sp. CB0101	103	EPYSEEDSGGLL.....
Synechococcus sp. CB0205	103	ESLLDQDAVGDLV.....
Synechococcus sp. CC9311	103	SSLMDALGVDPDIEEADS.....
Synechococcus sp. CC9605	103	SSLIDAVDEETDTTISSDP.....
Synechococcus sp. CC9902	103	SSFLDDSDDEESAPATSDS.....
Synechococcus sp. JA-2-3B	101	
Synechococcus sp. JA-3-3Ab	101	
Synechococcus sp. PCC 7002	105	EL.....
Synechococcus sp. RCC307	103	QVLDDFSE.....
Synechococcus sp. RS9916	103	SSFRDLSLSDMDVSTDS.....
Synechococcus sp. RS9917	103	SSLMDALEADTDSTDS.....
Synechococcus sp. WH 5701	83	DEMMD..ESDEI.....
Synechococcus sp. WH 7803	103	STLMDALEADTTDSDS.....
Synechococcus sp. WH 7805	103	STLMDALEADTTDSDS.....
Synechococcus sp. WH 8016	83	SSLMDALEDPDIAEADS.....
Synechococcus sp. WH 8102	103	SG.....
Synechococcus sp. WH 8109	103	SSLIDAVDEETDTTISSDP.....
Synechocystis sp. PCC 6803	103	EDQ.....
Thermosynechococcus e. BP-1	104	DLGLE.....
Trichodesmium e. IMS101	103	NL.....
consensus		

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(C)

			10	20	30	$\beta 1$	$\alpha 1$
Synechococcus e. PCC 7942	1		MTSAEMTSPNNSEHQ	AK	RTMIEGFD	ISHG	
Acaryochloris m. MBIC11017	1		MKKSNSLSQYKKNDKNKVEVT	K	LTRIEGFD	ISHG	
Acaryochloris sp. CCME 5410	1		MKKSNSLSQYKKNDKNKVEVT	K	LTRIEGFD	ISHG	
Anabaena v. ATCC 29413	1		MSEKEQEQQNTSGNGVEK	RTMIEGFD	ISHG		
Arthrospira m. CS-328	1		MVNSNPQPEPNKRNLISVQK	RTMIEGFD	ISHG		
Arthrospira m. FACHB-438	1		MVNSNLGSEPNKRKLISVQK	RTMIEGFD	ISHG		
Arthrospira p. str. Paraca	1		MVNSNLGSEPNKRKLISVQK	RTMIEGFD	ISHG		
Crocospaera w. WH 0003	1		MNQPLKSENQPPQLTPKGVK	RTMIEGLD	ISHG		
Crocospaera w. WH 8501	1		MNQPLKSENQPPQLTPKGVK	RTMIEGLD	ISHG		
Cyanobium sp. PCC 7001	1		MQQPSSPDTPLMQVQK	PTGIEGFD	ISHG		
Cyanotheca sp. ATCC 51142	1		MNQPLPRENQPPQLAPKGVK	RTMIEGLD	ISHG		
Cyanotheca sp. CCY0110	1			MIEGLD	ISHG		
Cyanotheca sp. PCC 7424	1		MNEPIPNQGEQYKVTTKGVK	RTMIEGFD	ISHG		
Cyanotheca sp. PCC 7425	1		MNTVNSKRENNPGEGRDGDGFVRKEVQK	RTMIEGFD	ISHG		
Cyanotheca sp. PCC 7822	1		MNEPIPSRQYKYKVTSGVK	RTMIEGFD	ISHG		
Cyanotheca sp. PCC 8801	1		MNEPISNSSKKTENTTQGVK	RTMIEGFD	ISHG		
Cyanotheca sp. PCC 8802	1		MNEPISNSSKKTENTTQGVK	RTMIEGFD	ISHG		
Cylindrospermopsis r. CS-505	1		MSDQEKQEKEQPTPTSAVEK	RTMIEGFD	ISHG		
Fischerella sp. JSC-11	1		MREKDDQPEKNAPLFGGVEK	RTMIEGFD	ISHG		
Leptolyngbya b. IAM M-101	1		MTSINQTEPRNPADLSSVK	RTMIEGFD	ISHG		
Lyngbya sp. PCC 8106	1		MSESSQDATKKNGRKLIGVK	RTMIEGFD	ISHG		
Microcoleus c. PCC 7420	1		MSPFNLDQRPDEFTTPGVK	RTMIEGFD	ISHG		
Microcoleus v. FGP-2	1		MNEISQTVQRQLGTAGVEK	RTMIEGFD	ISHG		
Microcystis a. NIES-843	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 7806	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 7820	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 7941	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 9443	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 9701	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 9717	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 9806	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis a. PCC 9807	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Microcystis sp. Ti-4	1		MTQSNNSQLNPPIKPKGVK	RTMIEGFD	ISHG		
Moorea p. 3L	1		MNQFNQNELYQNSVTTVGVK	RTMIEGFD	ISHG		
Nodularia s. CCY9414	1		MSEKDQAAPNNTPIAGV	RTMIEGFD	ISHG		
Nostoc a. 0708	1		MSDKAQEQNPPIGGVEK	RTMIEGFD	ISHG		
Nostoc c.	1		MSENEQEKNQPPIGGVEK	RTMIEGFD	ISHG		
Nostoc p. PCC 73102	1		MSQNEQVEPNKAPKNGGVEK	RTMIEGFD	ISHG		
Nostoc sp. PCC 7120	1		MSEKEQEQQNTSGNGVEK	RTMIEGFD	ISHG		
Nostoc sp. PCC 9709	1		MIENEQVEPKQPTIIRGVEK	RTMIEGFD	ISHG		
Oscillatoria sp. PCC 6506	1		MDNNTQILHHEVLVTPGVK	RTMIEGFD	ISHG		
Raphidiopsis b. D9	1		MSDQEKQEKEQQTPTSAVEK	RTMIEGFD	ISHG		
Synechococcus e. PCC 6301	1		MTSAEMTSPNNSEHQ	AK	RTMIEGFD	ISHG	
Synechococcus sp. PCC 7335	1		MNSSYSSSSSSDQKAQTRLKGVK	RTMIEGFD	ISHG		
Synechococcus sp. BL107	1			MQVQK	ATGIEGFD	ISHG	
Synechococcus sp. CB0101	1		MQDPSATNHALSSVQK	PTGIEGFD	ISHG		
Synechococcus sp. CB0205	1		MQDPSQSNHLASVQK	PTGIEGFD	ISHG		
Synechococcus sp. CC9311	1		MQISSSSGSPQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. CC9605	1		MQFPPTSGQPQMQVQK	PTGIEGFD	ISHG		
Synechococcus sp. CC9902	1		MQFPPASGQSQMVGK	ATGIEGFD	ISHG		
Synechococcus sp. JA-2-3B	1	MNQSLGPSEPEKPDNTAEDSTEPTPDNHRADLSELRG	PKKQTCIEGFD	ISHG			
Synechococcus sp. JA-3-3Ab	1	MNQPLEPSESEKLQGNAAEG	IPSTYFLGGPEPRG	PKKQTCIEGFD	ISHG		
Synechococcus sp. PCC 7002	1		MNQPTSSNNGAIGVQK	RTMIEGLD	ISHG		
Synechococcus sp. RCC307	1		MTSPSPDPSPSEMRLSQVQK	PTGIEGFD	ISHG		
Synechococcus sp. RS9916	1		MKISGSTGSPQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. RS9917	1		MQQNVGVMQISSSSSPQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. WH 5701	1		MQDFNAVSPQSMK	QKPTGIEGLD	ISHG		
Synechococcus sp. WH 7803	1		MQISSSSGSPQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. WH 7805	1		MQISSSSGSPQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. WH 8016	1			MQVQK	PTGIEGFD	ISHG	
Synechococcus sp. WH 8102	1		MQFPPASGSTQMVGK	PTGIEGFD	ISHG		
Synechococcus sp. WH 8109	1			MQVQK	PTGIEGFD	ISHG	
Synechocystis sp. PCC 6803	1		MNLPVNERNRDVPKGVK	RTMIEGFD	ISHG		
Thermosynechococcus e. BP-1	1		MTNLPEHQSSPTEQSSAEVKK	PTMIEGFD	ISHG		
Trichodesmium e. IMS101	1		MNQNSESVKKDRIATVGVEK	RTMIEGFD	ISHG		
consensus							

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					$\beta 2$	$\beta 3$	$\alpha 2$	$\beta 4$	$\alpha 3$			
					40	50	60	70	80			
Synechococcus e. PCC 7942	35	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGIIEFDDEPG	FVTFEE	PQDI	KNARS	
Acaryochloris m. MBIC11017	36	G	P	GR	TL	SGTSGTGKT	FA	QFL	HGIVHFDDEPG	FVTFEESPKDI	QNALS	
Acaryochloris sp. CCME 5410	36	GLP	GR	TL	SGTSGTGKT	FA	QFL	HGIVHFDDEPG	FVTFEESPKDI	QNALS		
Anabaena v. ATCC 29413	34	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGISFDEDPG	FVTFEESPD	KN	NAHI	
Arthrospira m. CS-328	35	G	P	AGR	TL	SGTSGTGKT	FA	QFL	NGITYFDEDPG	LVTTEE	PTDI	KNAS
Arthrospira m. FACHB-438	35	G	P	AGR	TL	SGTSGTGKT	FA	QFL	NGITYFDEDPG	LVTTEE	PTDI	KNAS
Arthrospira p. str. Paraca	35	G	P	AGR	TL	SGTSGTGKT	FA	QFL	NGITYFDEDPG	LVTTEE	PTDI	KNAS
Crocospaera w. WH 0003	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	CHGIKYFDYPC	FVTFEESPRDI	ENAYS		
Crocospaera w. WH 8501	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	CHGIKYFDYPC	FVTFEESPRDI	ENAYS		
Cyanobium sp. PCC 7001	31	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGIROKDEPG	FVTFEESPLDI	KNAS		
Cyanothece sp. ATCC 51142	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPHDI	ENAYS		
Cyanothece sp. CCY0110	12	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPHDI	ENAYS		
Cyanothece sp. PCC 7424	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Cyanothece sp. PCC 7425	45	GLP	RGR	TL	SGTSGTGKT	LA	QFL	NGITLFDDEPG	FVTFEESPAD	IARNALS		
Cyanothece sp. PCC 7822	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Cyanothece sp. PCC 8801	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Cyanothece sp. PCC 8802	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Cylindrospermopsis r. CS-505	37	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGITHFDEPG	FVTFEESPD	KN	NAHI	
Fischerella sp. JSC-11	35	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGISYFDEPG	FVTFEESPHDI	KN	NAHI	
Leptolyngbya b. IAM M-101	35	GLP	GR	TL	SGTSGTGKT	LA	QFL	NGITQFDEAG	FVTFEESPD	KN	NAHS	
Lyngbya sp. PCC 8106	36	G	P	GR	TL	SGTSGTGKT	FA	QFL	NGIINFDEPG	FVTFEESPNDI	KNASS	
Microcoleus c. PCC 7420	36	GLP	R	TL	SGTSGTGKT	FA	QFL	NGITOFDDPG	FVTFEESPNDI	KN	NAHS	
Microcoleus v. FGP-2	35	G	P	GR	TL	SGTSGTGKT	FA	QFL	NGITHFDEAG	FVTFEESPAD	KNACS	
Microcystis a. NIES-843	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 7806	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 7820	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 7941	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 9443	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 9701	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 9717	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 9806	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis a. PCC 9807	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Microcystis sp. T1-4	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGIKYFDYPC	FVTFEESPD	KN	NAHS	
Moorea p. 3L	36	GLP	SGR	TL	SGTSGTGKT	LA	QFL	NGISNFDEAG	FVTFEESPD	KN	NAHS	
Nodularia s. CCY9414	34	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGITRFDEPG	FVTFEESPD	KN	NAHI	
Nostoc a. 0708	34	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGITYFDEPG	FVTFEESPD	KN	NAHI	
Nostoc c.	34	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGITYFDEPG	FVTFEESPD	KN	NAHI	
Nostoc p. PCC 73102	35	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGITYFDEAG	FVTFEESPD	KN	NAHI	
Nostoc sp. PCC 7120	34	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGISFDEPG	FVTFEESPD	KN	NAHI	
Nostoc sp. PCC 9709	35	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGITYFDEAG	FVTFEESPD	KN	NAHI	
Oscillatoria sp. PCC 6506	35	G	P	GR	TL	SGTSGTGKT	FA	QFL	NGIMNFDEAG	FVTFEESPD	KN	NAHS
Raphidiopsis b. D9	37	GLP	GR	TL	SGTSGTGKT	LS	QFL	NGITHFDEPG	FVTFEESPD	KN	NAHI	
Synechococcus e. PCC 6301	35	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGIIEFDDEPG	FVTFEE	PQDI	KNARS	
Synechococcus sp. PCC 7335	39	GLP	GR	TL	SGTSGTGKT	LA	QFL	HGITYFDEPG	FVTFEESPD	KN	NAHS	
Synechococcus sp. BL107	20	GLP	GR	TL	SGTSGTGKT	LS	HFL	HNGIKHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. CB0101	31	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGIROFDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. CB0205	31	GLP	GR	TL	SGTSGTGKT	FS	QFL	NGIROFDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. CC9311	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. CC9605	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIKHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. CC9902	31	GLP	GR	TL	SGTSGTGKT	LS	HFL	HNGIKHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. JA-2-3B	56	GLP	GR	TL	SGTSGTGKT	FA	QFL	NGIVKQDEPG	FVTFEE	PADI	KNASS	
Synechococcus sp. JA-3-3Ab	52	GLP	GR	TL	SGTSGTGKT	FA	QFL	NGIVKQDEPG	FVTFEE	PADI	KNASS	
Synechococcus sp. PCC 7002	32	GLP	SGR	TL	SGTSGTGKT	LA	QFL	HGIKHFDYPC	FVTFEESPRDI	KN	NAHS	
Synechococcus sp. RCC307	35	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIQHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. RS9916	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. RS9917	38	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 5701	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIRHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 7803	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 7805	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 8016	20	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIAHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 8102	31	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIKHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechococcus sp. WH 8109	20	GLP	GR	TL	SGTSGTGKT	FS	HFL	HNGIKHFDDEPG	FVTFEESPLDI	KN	NAHS	
Synechocystis sp. PCC 6803	36	GLP	GR	TL	SGTSGTGKT	LA	QFL	QGIHFDYPC	FVTFEESPD	KN	NAHS	
Thermosynechococcus e. BP-1	36	GLP	QGR	TL	SGTSGTGKT	FA	QFL	NGITIFDEPG	FVTFEESPD	KN	NAHS	
Trichodesmium e. IMS101	35	G	P	GR	TL	SGTSGTGKT	FA	QFL	NGITYFDEPG	FVTFEESPD	KN	NAHS
consensus												

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			$\alpha 4$	$\beta 5$	$\alpha 5$	$\beta 6$		
		90	90	100	110	120	130	140
Synechococcus e. PCC 7942	90	FGWDLAKLVDEGKLFILDA	SPDP	GGQEVVGGFDLSALIERI	QYATQKYAARRVSI			
Acaryochloris m. MBIC11017	91	FGWDLQQLHDDGKLFILDA	SPDP	GGQDIAGDFDLSALIERI	QYATSKYQARRVGI			
Acaryochloris sp. CCME 5410	91	FGWDLQQLHDDGKLFILDA	SPDP	GGQDIAGDFDLSALIERI	QYATSKYQARRVGI			
Anabaena v. ATCC 29413	89	FGWNLQRLINEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Arthrospira m. CS-328	90	FGWDLQQLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Arthrospira m. FACHB-438	90	FGWDLQQLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Arthrospira p. str. Paraca	90	FGWDLQQLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Crocospaera w. WH 0003	91	FGWDLQQLIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATNKYNAARLVSI			
Crocospaera w. WH 8501	91	FGWDLQQLIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATNKYNAARLVSI			
Cyanobium sp. PCC 7001	86	FGWNLQENLEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Cyanotheca sp. ATCC 51142	91	FGWDLQKLIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATHKYKARRVSI			
Cyanotheca sp. CCY0110	67	FGWDLQALIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATNKYKARRVSI			
Cyanotheca sp. PCC 7424	91	FGWDLNELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Cyanotheca sp. PCC 7425	100	FGWDLQKLIDEGKLFILDA	SPDP	GGQEVAGDFDLSALIERI	QYATIRKYKARRVSI			
Cyanotheca sp. PCC 7822	91	FGWDLNQLIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Cyanotheca sp. PCC 8801	91	FGWDLQELIVDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATNKYKARRVSI			
Cyanotheca sp. PCC 8802	91	FGWDLQELIVDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATNKYKARRVSI			
Cylindrospermopsis r. CS-505	92	FGWNLQKLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Fischerella sp. JSC-11	90	FGWDLQGLINDGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Leptolyngbya b. IAM M-101	90	FGWDLQRFVDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Lyngbya sp. PCC 8106	91	FGWDLQELIVDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Microcoleus c. PCC 7420	91	FGWNLQKLIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Microcoleus v. FGP-2	90	FGWDLAQLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. NIES-843	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 7806	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 7820	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 7941	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 9443	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 9701	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 9717	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 9806	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis a. PCC 9807	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Microcystis sp. T1-4	91	FGWDLQELIDEGKLFILDA	SPDP	GGQEVVGSFDLSALIERI	QYATIRKYKARRVSI			
Moorea p. 3L	91	FGWELNKLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATHKYKARRVSI			
Nodularia s. CCY9414	89	FGWNLQHLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Nostoc a. 0708	89	FGWNLQKLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Nostoc c.	89	FGWNLQRLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Nostoc p. PCC 73102	90	FGWNLQRLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Nostoc sp. PCC 7120	89	FGWNLQRLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Nostoc sp. PCC 9709	90	FGWNLQRLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Oscillatoria sp. PCC 6506	90	FGWDLQRLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Raphidiopsis b. D9	92	FGWNLQKLIDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus e. PCC 6301	90	FGWDLAKLVDEGKLFILDA	SPDP	GGQEVVGGFDLSALIERI	QYATQKYAARRVSI			
Synechococcus sp. PCC 7335	94	FGWDLKSLKSGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. BL107	75	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. CB0101	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. CB0205	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. CC9311	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. CC9605	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. CC9902	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. JA-2-3B	111	FGWDLQALIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. JA-3-3Ab	107	FGWDLQALIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. PCC 7002	87	FGWDLQSLVDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. RCC307	90	FGWDLQSMYNDKLFILDA	SPDP	GGQDVAGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. RS9916	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. RS9917	93	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 5701	86	FGWNLQELVAQKLFILDA	SPDP	GGQDVAGNFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 7803	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 7805	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 8016	75	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 8102	86	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechococcus sp. WH 8109	75	FGWNLQENVEQDKLFILDA	SPDP	GGQDVAGSFDLSALIERI	QYATIRKYKARRVSI			
Synechocystis sp. PCC 6803	91	FGWDLQQLIDEGKLFILDA	SPDP	GGQEVVGNFDLSALIERI	QYATIRKYKARRVSI			
Thermosynechococcus e. BP-1	91	FGWNLQSLIDEGKLFILDA	SPDP	GGQEVAGDFDLSALIERI	QYATIRKYKARRVSI			
Trichodesmium e. IMS101	90	FGWNLQQLVDEGKLFILDA	SPDP	GGQDIVGNFDLSALIERI	QYATIRKYKARRVSI			
consensus								

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			$\beta 6$	$\alpha 6$	$\alpha 7$	$\beta 7$	$\alpha 8$																																									
			150	160	170	180	190																																									
Synechococcus e. PCC 7942	145	DS	T	V	F	Q	Q	A	S	V	V	R	R	E	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F				
Acaryochloris m. MBIC11017	146	DS	T	A	F	Q	Q	Y	A	A	T	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Acaryochloris sp. CCME 5410	146	DS	T	A	F	Q	Q	Y	A	A	T	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Anabaena v. ATCC 29413	144	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Arthrospira m. CS-328	145	DS	T	A	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	S	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Arthrospira m. FACHB-438	145	DS	T	A	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Arthrospira p. str. Paraca	145	DS	T	A	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Crocospaera w. WH 0003	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Crocospaera w. WH 8501	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanobium sp. PCC 7001	141	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	E	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. ATCC 51142	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. CCY0110	122	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. PCC 7424	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. PCC 7425	155	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. PCC 7822	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. PCC 8801	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cyanotheca sp. PCC 8802	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Cylindrospermopsis r. CS-505	147	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Fischerella sp. JSC-11	145	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Leptolyngbya b. IAM M-101	145	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Lyngbya sp. PCC 8106	146	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcoleus c. PCC 7420	146	DS	T	A	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Microcoleus v. FGP-2	145	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. NIES-843	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 7806	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 7820	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 7941	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 9443	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 9701	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 9717	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 9806	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis a. PCC 9807	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Microcystis sp. TI-4	146	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	S	I	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Moorea p. 3L	146	DS	T	A	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	S	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Nodularia s. CCY9414	144	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Nostoc a. 0708	144	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Nostoc c.	144	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Nostoc p. PCC 73102	145	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	I	S	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Nostoc sp. PCC 7120	144	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Nostoc sp. PCC 9709	145	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	L	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Oscillatoria sp. PCC 6506	145	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Raphidiopsis b. D9	147	DS	T	A	V	F	Q	Q	Y	A	A	G	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Synechococcus e. PCC 6301	145	DS	T	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	A	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F	
Synechococcus sp. PCC 7335	149	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	Q	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Synechococcus sp. BL107	130	DS	T	A	V	F	Q	Q	Y	A	A	S	V	V	R	R	E	I	F	R	L	A	R	K	E	I	G	V	T	T	M	T	T	E	R	E	E	Y	G	P	A	R	F	G	V	E	E	F
Synechococcus sp. CB0101	141	DS	T	A	V	F	Q	Q	Y	A	A	S																																				

			$\alpha 8$	$\beta 8$	$\beta 9$	$\beta 10$	$\beta 11$	
			200	210	220	230	240	250
Synechococcus e. PCC 7942	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDHGIN	IFPLGAMRL
Acaryochloris m. MBIC11017	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDDGIN	IFPLGAMRL
Acaryochloris sp. CCMEE 5410	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDDGIN	IFPLGAMRL
Anabaena v. ATCC 29413	199	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Arthrospira m. CS-328	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TQNGIS	IFPLGAMRL
Arthrospira m. FACHB-438	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TQNGIS	IFPLGAMRL
Arthrospira p. str. Paraca	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TQNGIS	IFPLGAMRL
Crocospaera w. WH 0003	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Crocospaera w. WH 8501	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanobium sp. PCC 7001	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GSHGIS	IFPLGAMRL
Cyanothece sp. ATCC 51142	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanothece sp. CCY0110	177	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanothece sp. PCC 7424	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanothece sp. PCC 7425	210	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNQGIS	IFPLGAMRL
Cyanothece sp. PCC 7822	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanothece sp. PCC 8801	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cyanothece sp. PCC 8802	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Cylindrospermopsis r. CS-505	202	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Fischerella sp. JSC-11	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNEGIS	IFPLGAMRL
Leptolyngbya b. IAM M-101	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNQGIS	IFPLGAMRL
Lyngbya sp. PCC 8106	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNNGIS	IFPLGAMRL
Microcoleus c. PCC 7420	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	SNQGIS	IFPLGAMRL
Microcoleus v. FGP-2	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. NIES-843	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 7806	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 7820	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 7941	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 9443	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 9701	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 9717	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 9806	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis a. PCC 9807	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Microcystis sp. T1-4	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Moorea p. 3L	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDQGIS	IFPLGAMRL
Nodularia s. CCY9414	199	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Nostoc a. 0708	199	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Nostoc c.	199	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Nostoc p. PCC 73102	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNEGIS	IFPLGAMRL
Nostoc sp. PCC 7120	199	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Nostoc sp. PCC 9709	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNEGIS	IFPLGAMRL
Oscillatoria sp. PCC 6506	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNNGIS	IFPLGAMRL
Raphidiopsis b. D9	202	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Synechococcus e. PCC 6301	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDHGIN	IFPLGAMRL
Synechococcus sp. PCC 7335	204	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TDGGIS	IFPLGAMRL
Synechococcus sp. BL107	185	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. CB0101	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GSHGIS	IFPLGAMRL
Synechococcus sp. CB0205	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GSHGIS	IFPLGAMRL
Synechococcus sp. CC9311	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. CC9605	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GTHGIS	IFPLGAMRL
Synechococcus sp. CC9902	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. JA-2-3B	221	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Synechococcus sp. JA-3-3Ab	217	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNEGIS	IFPLGAMRL
Synechococcus sp. PCC 7002	197	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL
Synechococcus sp. RCC307	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GLHGIS	IFPLGAMRL
Synechococcus sp. RS9916	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. RS9917	203	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. WH 5701	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. WH 7803	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. WH 7805	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. WH 8016	185	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GAHGIS	IFPLGAMRL
Synechococcus sp. WH 8102	196	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GTHGIS	IFPLGAMRL
Synechococcus sp. WH 8109	185	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	GTHGIS	IFPLGAMRL
Synechocystis sp. PCC 6803	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	THDGIN	IFPLGAMRL
Thermosynechococcus e. BP-1	201	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNAGIS	IFPLGAMRL
Trichodesmium e. IMS101	200	VSDNVVILRNVL	EGERRRRT	LEILKLRGT	THMKGE	PFT	TNDGIN	IFPLGAMRL

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			$\beta 12$	$\alpha 9$	$\beta 13$	$\beta 14$	$\alpha 10$
			260	270	280	290	300
Synechococcus e. PCC 7942	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Acaryochloris m. MBIC11017	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Acaryochloris sp. CCME 5410	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Anabaena v. ATCC 29413	254	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Arthrospira m. CS-328	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Arthrospira m. FACHB-438	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Arthrospira p. str. Paraca	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Crocospaera w. WH 0003	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Crocospaera w. WH 8501	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanobium sp. PCC 7001	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. ATCC 51142	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. CCY0110	232	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. PCC 7424	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. PCC 7425	265	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. PCC 7822	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. PCC 8801	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cyanotheca sp. PCC 8802	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Cylindrospermopsis r. CS-505	257	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Fischerella sp. JSC-11	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Leptolyngbya b. IAM M-101	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Lyngbya sp. PCC 8106	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcoleus c. PCC 7420	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcoleus v. FGP-2	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. NIES-843	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 7806	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 7820	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 7941	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 9443	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 9701	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 9717	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 9806	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis a. PCC 9807	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Microcystis sp. T1-4	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Moorea p. 3L	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nodularia s. CCY9414	254	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nostoc a. 0708	254	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nostoc c.	254	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nostoc p. PCC 73102	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nostoc sp. PCC 7120	254	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Nostoc sp. PCC 9709	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Oscillatoria sp. PCC 6506	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Raphidiopsis b. D9	257	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus e. PCC 6301	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. PCC 7335	259	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. BL107	240	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. CB0101	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. CB0205	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. CC9311	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. CC9605	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. CC9902	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. JA-2-3B	276	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. JA-3-3Ab	272	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. PCC 7002	252	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. RCC307	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. RS9916	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. RS9917	258	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 5701	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 7803	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 7805	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 8016	240	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 8102	251	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechococcus sp. WH 8109	240	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Synechocystis sp. PCC 6803	256	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Thermosynechococcus e. BP-1	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
Trichodesmium e. IMS101	255	TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN
consensus		TQRSSNV	SSGVVRLD	MCGGGFFKDS	IILATGATGTGKTL	LVSKF	EDACN

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			$\beta 15$	$\alpha 11$	$\alpha 12$	$\beta 16$	$\alpha 13$	
			310	320	330	340	350	360
Synechococcus e. PCC 7942	310	ERAILFAYEESRAQLRNAY	SWG	DFBE	ERQNL	LLK	IVCA	PESA
Acarochloris m. MBIC11017	311	RAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Acarochloris sp. CCME 5410	311	RAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Anabaena v. ATCC 29413	309	ERAILFAYEESRAQLRNAY	SWGIDFBE	ESQ	GLLK	ICTY	PESA	AGLEDHLQ
Arthrospira m. CS-328	310	ERAILFAYEESRAQLRNAY	SWG	DFBE	ERK	GLLK	ICTY	PESA
Arthrospira m. FACHB-438	310	ERAILFAYEESRAQLRNAY	SWG	DFBE	ERK	GLLK	ICTY	PESA
Arthrospira p. str. Paraca	310	ERAILFAYEESRAQLRNAY	SWG	DFBE	ERK	GLLK	ICTY	PESA
Crocospaera w. WH 0003	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Crocospaera w. WH 8501	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanobium sp. PCC 7001	306	RAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. ATCC 51142	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. CCY0110	287	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. PCC 7424	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. PCC 7425	320	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. PCC 7822	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. PCC 8801	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cyanothece sp. PCC 8802	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Cylindrospermopsis r. CS-505	312	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Fischerella sp. JSC-11	310	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Leptolyngbya b. IAM M-101	310	RAILFAYEESRAQLRNAY	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Lyngbya sp. PCC 8106	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Microcoleus c. PCC 7420	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	EEK	GLLK	ICTY	PESA	AGLEDHLQ
Microcoleus v. FGP-2	310	NRALFAYEESRAQLRNAY	SWGIDFBE	EEK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. NIES-843	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 7806	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 7820	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 7941	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 9443	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 9701	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 9717	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 9806	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis a. PCC 9807	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Microcystis sp. T1-4	311	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Mocrea p. 3L	311	RAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Nodularia s. CCY9414	309	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Nostoc a. 0708	309	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQH	GLLK	ICTY	PESA	AGLEDHLQ
Nostoc c.	309	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQH	GLLK	ICTY	PESA	AGLEDHLQ
Nostoc p. PCC 73102	310	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Nostoc sp. PCC 7120	309	ERAILFAYEESRAQLRNAY	SWGIDFBE	ESQ	GLLK	ICTY	PESA	AGLEDHLQ
Nostoc sp. PCC 9709	310	ERAILFAYEESRAQLRNAY	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Oscillatoria sp. PCC 6506	310	RAILFAYEESRAQLRNAY	SWGIDFBE	EQK	GLLK	ICTY	PESA	AGLEDHLQ
Raphidiopsis b. D9	312	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus e. PCC 6301	310	ERAILFAYEESRAQLRNAY	SWG	DFBE	ERQNL	LLK	IVCA	PESA
Synechococcus sp. PCC 7335	314	ERAILFAYEESRAQLRNAY	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. BL107	295	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. CB0101	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. CB0205	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. CC9311	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. CC9605	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. CC9902	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. JA-2-3B	331	ERCILFAYEESRAQLRNAT	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. JA-3-3Ab	327	ERCILFAYEESRAQLRNAT	SWGIDFBE	ERQ	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. PCC 7002	307	ERAILFAYEESRAQLRNAT	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. RCC307	310	ERAILFAYEESRAQLRNAT	SWGIDFBE	ESD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. RS9916	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. RS9917	313	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 5701	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 7803	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 7805	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 8016	295	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 8102	306	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechococcus sp. WH 8109	295	ERAILFAYEESRAQLRNAT	SWGIDFBE	EQD	GLLK	ICTY	PESA	AGLEDHLQ
Synechocystis sp. PCC 6803	311	ERAILFAYEESRAQLRNAT	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Thermosynechococcus e. BP-1	310	ERAILFAYEESRAQLRNAT	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
Trichodesmium e. IMS101	310	ERAILFAYEESRAQLRNAT	SWGIDFBE	ERK	GLLK	ICTY	PESA	AGLEDHLQ
consensus		*****	*****	*****	*****	*****	*****	*****

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			$\alpha 13$	$\beta 17$	$\alpha 14$	$\beta 18$																																																
			370	380	390	400	410																																															
Synechococcus e. PCC 7942	365	E	I	N	D	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	L	F	T	N	T	S	I	Q	F
Acaryochloris m. MBIC11017	366	E	I	A	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Acaryochloris sp. CCMEE 5410	366	E	I	A	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Anabaena v. ATCC 29413	364	E	I	A	Y	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Arthrospira m. CS-328	365	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Arthrospira m. FACHB-438	365	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Arthrospira p. str. Paraca	365	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Crocospaera w. WH 0003	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Crocospaera w. WH 8501	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanobium sp. PCC 7001	361	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. ATCC 51142	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. CCY0110	342	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. PCC 7424	366	E	I	A	L	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. PCC 7425	375	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. PCC 7822	366	E	I	A	L	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. PCC 8801	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cyanotheca sp. PCC 8802	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Cylindrospermopsis r. CS-505	367	E	I	A	D	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Fischerella sp. JSC-11	365	E	I	A	D	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Leptolyngbya b. IAM M-101	365	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Lyngbya sp. PCC 8106	366	E	I	T	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcoleus c. PCC 7420	366	E	I	Q	I	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F
Microcoleus v. FGP-2	365	E	I	A	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. NIES-843	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 7806	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 7820	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 7941	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 9443	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 9701	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 9717	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 9806	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis a. PCC 9807	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Microcystis sp. T1-4	366	E	I	S	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Moorea p. 3L	366	E	I	A	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nodularia s. CCY9414	364	E	I	A	V	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nostoc a. 0708	364	E	I	A	D	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nostoc c.	364	E	I	A	N	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nostoc p. PCC 73102	365	E	I	A	V	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nostoc sp. PCC 7120	364	E	I	A	Y	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Nostoc sp. PCC 9709	365	E	I	A	I	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Oscillatoria sp. PCC 6506	365	E	I	A	E	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V	I	G	V	T	G	Y	A	K	Q	E	E	I	T	G	F	F	N	T	T	I	H	F	
Raphidiopsis b. D9	367	E	I	A	D	F	K	P	S	R	A	I	D	S	L	S	A	L	A	R	G	V	S	N	N	A	F	R	Q	F	V																							

				$\beta 19$	$\beta 20$	$\beta 21$			
		420	430	440	450	460	470		
Synechococcus e. PCC 7942	420	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREF	MTSD
Acaryochloris m. MBIC11017	421	LG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Acaryochloris sp. CCME 5410	421	LG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Anabaena v. ATCC 29413	419	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Arthrospira m. CS-328	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Arthrospira m. FACHB-438	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Arthrospira p. str. Paraca	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Crocospaera w. WH 0003	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	MTNQ
Crocospaera w. WH 8501	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	MTNQ
Cyanobium sp. PCC 7001	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Cyanothece sp. ATCC 51142	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	MTNQ
Cyanothece sp. CCY0110	397	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	MTNQ
Cyanothece sp. PCC 7424	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Cyanothece sp. PCC 7425	430	LG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREY	TDIA
Cyanothece sp. PCC 7822	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Cyanothece sp. PCC 8801	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STNE
Cyanothece sp. PCC 8802	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STNE
Cylindrospermopsis r. CS-505	422	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Fischerella sp. JSC-11	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Leptolyngbya b. IAM M-101	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Lyngbya sp. PCC 8106	421	MG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTID
Microcoleus c. PCC 7420	421	MG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREY	MTSK
Microcoleus v. FGP-2	420	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Microcystis a. NIES-843	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 7806	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 7820	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 7941	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 9443	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 9701	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 9717	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 9806	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis a. PCC 9807	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Microcystis sp. Ti-4	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSQ
Moorea p. 3L	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREY	MTSK
Nodularia s. CCY9414	419	LG	HSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Nostoc a. 0708	419	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Nostoc c.	419	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Nostoc p. PCC 73102	420	LG	HSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Nostoc sp. PCC 7120	419	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Nostoc sp. PCC 9709	420	LG	HSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Oscillatoria sp. PCC 6506	420	MG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Raphidiopsis b. D9	422	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	NTIA
Synechococcus e. PCC 6301	420	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREF	MTSD
Synechococcus sp. PCC 7335	424	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREF	MTSD
Synechococcus sp. BL107	405	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. CB0101	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	MTSD
Synechococcus sp. CB0205	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIS
Synechococcus sp. CC9311	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	FLTG
Synechococcus sp. CC9605	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. CC9902	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. JA-2-3B	441	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	LTSD
Synechococcus sp. JA-3-3Ab	437	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	LTSD
Synechococcus sp. PCC 7002	417	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Synechococcus sp. RCC307	420	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	IREF	MTSD
Synechococcus sp. RS9916	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. RS9917	423	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. WH 5701	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. WH 7803	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. WH 7805	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. WH 8016	405	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	FLTG
Synechococcus sp. WH 8102	416	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechococcus sp. WH 8109	405	MG	SHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIG
Synechocystis sp. PCC 6803	421	MG	AHSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	STSH
Thermosynechococcus e. BP-1	420	MG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIE
Trichodesmium e. IMS101	420	MG	NSIT	SHISTITDTIL	LQYVEIRGEM	SRA	NVFKMRGSWHDK	GIREY	VTIS
consensus			*	*****	*****	*	*****	*****	*

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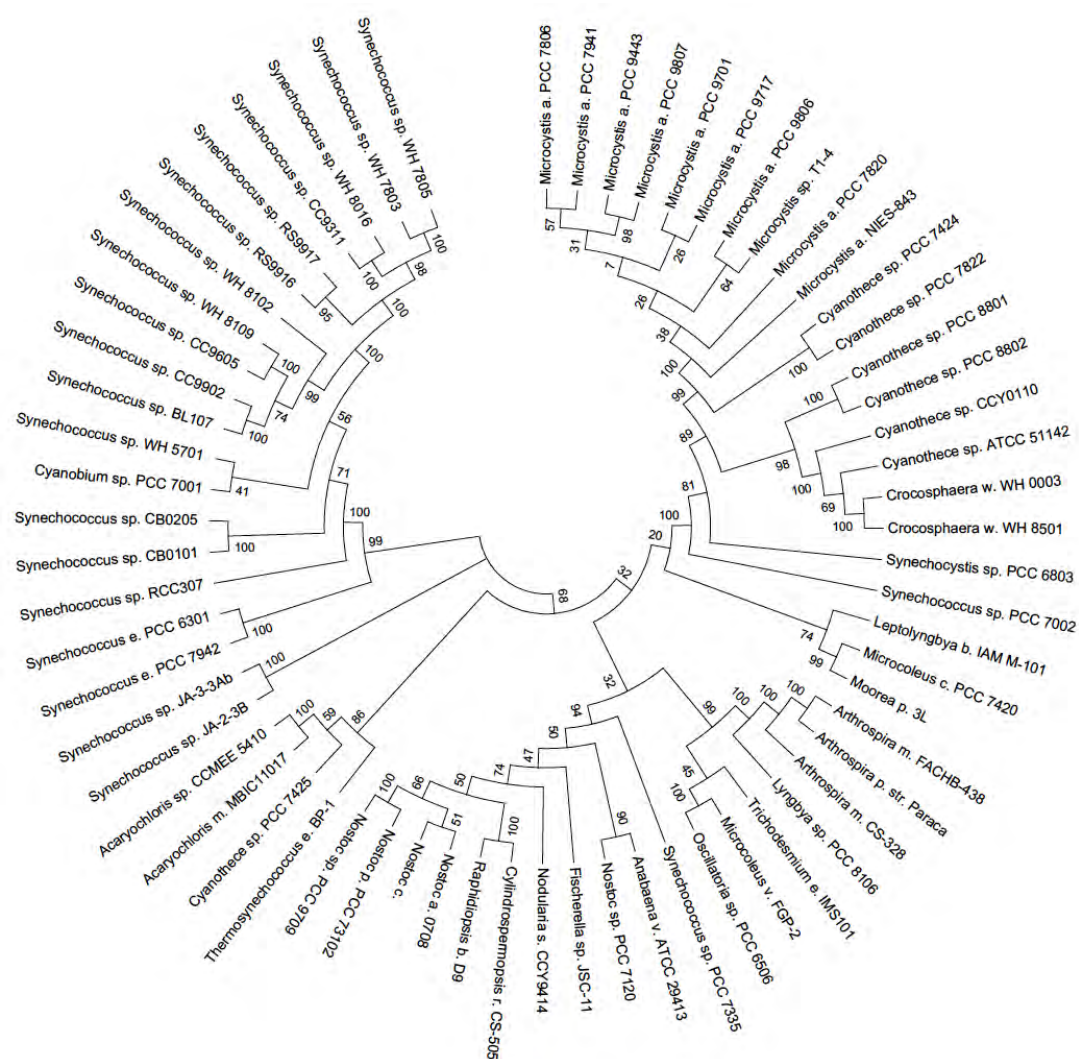


			$\beta 22$	$\beta 23$	$\beta 24$			
			480	490	500	510		
Synechococcus e. PCC 7942	475	KCPD	IQDSFRN	YERII	SGSPTRITVDEKSELSRI	VGVQEKGPES...		
Acaryochloris m. MBIC11017	476	HCP	IKMFAHNF	EGII	SGTPTRVSD	DEKRLSLRI	VQDVKGLSDDDLL.	
Acaryochloris sp. CCME 5410	476	HCP	IKMFAHNF	EGII	SGTPTRVSD	DEKRLSLRI	VQDVKGLSDDDLL.	
Anabaena v. ATCC 29413	474	DGPEI	IQDSFRN	YERII	SGSPTRVSD	DEKAELSRI	VRFEDKQGS...	
Arthrospira m. CS-328	475	HCPHIT	DSFRN	YERII	SGSPTRISIE	EKSELSRI	IEGVQDDQD....	
Arthrospira m. FACHB-438	475	HCPHIT	DSFRN	YERII	SGSPTRISIE	EKSELSRI	IEGVQDDQD....	
Arthrospira p. str. Paraca	475	HCPHIT	DSFRN	YERII	SGSPTRISIE	EKSELSRI	IEGVQDDQD....	
Crocospaera w. WH 0003	476	DGPII	IQDSFRN	YERII	SGSPSRISV	DEKTELSRI	VGVKDKTEE....	
Crocospaera w. WH 8501	476	DGPII	IQDSFRN	YERII	SGSPSRISV	DEKTELSRI	VGVKDKTEE....	
Cyanobium sp. PCC 7001	471	NCPEI	IQDSFRN	YERII	SGSPSRITV	DEKSELSRI	VGVSDDR.....	
Cyanothece sp. ATCC 51142	476	DGPII	IQDSFRN	YERII	SGSPSRITV	DEKSELSRI	VGVKDKTEE....	
Cyanothece sp. CCY0110	452	DGPII	IQDSFRN	YERII	SGSPSRITV	DEKSELSRI	VGVKDKTEE....	
Cyanothece sp. PCC 7424	476	DGAEI	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VGVKDKTLDE..	
Cyanothece sp. PCC 7425	485	QCPEI	IQDSFRN	YERII	SGTPTRVIT	DEKTELSRI	VADVQSLEGL....	
Cyanothece sp. PCC 7822	476	DGAEI	IQDSFRN	YERII	SGSPTRITV	DEKTELSRI	VGVKDKTLDE..	
Cyanothece sp. PCC 8801	476	SCPTI	IQDSFRN	YERII	SGSPTRITV	DEKNEL	SLRI	VGVKDKTLDEE..
Cyanothece sp. PCC 8802	476	SCPTI	IQDSFRN	YERII	SGSPTRITV	DEKNEL	SLRI	VGVKDKTLDEE..
Cylindrospermopsis r. CS-505	477	DGPNI	IQDSFRN	YERII	SGAPTRVSD	DEKAELSRI	VRFEDK.....	
Fischerella sp. JSC-11	475	DGPEI	IQDSFRN	YERII	SGAPTRVSD	DEKAELSRI	VSFQDKDSSDASS	
Leptolyngbya b. IAM M-101	475	QCPEI	IQDSFRN	YERII	SGSPTRIA	DEKVL	SLRI	TGERSDK.....
Lyngbya sp. PCC 8106	476	KCPRI	ITDSFRN	YERII	SGSPTRISV	EKSELSRI	TEGVQGGGEIDD..	
Microcoleus c. PCC 7420	476	DGADI	IQDSFRN	YERII	SGSPSRITV	DEKSELSRI	VGVVREKLQDE...	
Microcoleus v. FGP-2	475	RCPQI	IQDSFRN	YERII	SGSPSRITV	EKSELSRI	LQGVQGGDEEDI..	
Microcystis a. NIES-843	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Microcystis a. PCC 7806	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Microcystis a. PCC 7820	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Microcystis a. PCC 7941	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VNVREKLQDE...	
Microcystis a. PCC 9443	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VNVREKLQDE...	
Microcystis a. PCC 9701	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Microcystis a. PCC 9717	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VNVREKLQDE...	
Microcystis a. PCC 9806	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Microcystis a. PCC 9807	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VNVREKLQDE...	
Microcystis sp. T1-4	476	DGADI	IQDSFRN	YERII	SGSPTRISV	DEKTELSRI	VNVREKLQDE...	
Moorea p. 3L	476	DGPEI	IQDSFRN	YERII	SGSPTRIP	DEKSELSRI	VGVKDKSGE....	
Nodularia s. CCY9414	474	DGPEI	IQDSFRN	YERII	SGAPTRVSD	DEKAELSRI	VQSFENKPSDSSI	
Nostoc a. 0708	474	DGPNI	IQDSFRN	YERII	SGAPTRVIT	DEKAELSRI	VRFEEK.....	
Nostoc c.	474	DGPEI	IQDSFRN	YERII	NGAPTRVSD	DEKAELSRI	VSFYRSDS...	
Nostoc p. PCC 73102	475	DGPD	IQDSFRN	YERII	SGAPTRVSD	DEKAELSRI	VRFEDKQSSSEP..	
Nostoc sp. PCC 7120	474	DGPEI	IQDSFRN	YERII	SGSPTRVSD	DEKAELSRI	VRFEDKQGSDS..	
Nostoc sp. PCC 9709	475	DGPD	IQDSFRN	YERII	SGAPTRVSD	DEKAELSRI	VRFEDKQSSSEP..	
Oscillatoria sp. PCC 6506	475	KGPEI	IQDSFRN	YERII	SGSPTRISV	EKSELSRI	VGVQGGGDGEEI	
Raphidiopsis b. D9	477	DGPNI	IQDSFRN	YERII	SGAPSRVSD	DEKAELSRI	VRFEDK.....	
Synechococcus e. PCC 6301	475	KGPD	IQDSFRN	YERII	SGSPTRITV	DEKSELSRI	VGVQEKGPES...	
Synechococcus sp. PCC 7335	479	DGPD	IQDSFRN	YERII	SGSARRIPS	REKNEI	RVSAVRSDDD....	
Synechococcus sp. BL107	460	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AXSVSSDD.....	
Synechococcus sp. CB0101	471	NGPEI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	VQGVTSDDRL....	
Synechococcus sp. CB0205	471	NGPEI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	VGVGEGQF.....	
Synechococcus sp. CC9311	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVSP.....	
Synechococcus sp. CC9605	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVSTED.....	
Synechococcus sp. CC9902	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVSVSSDD.....	
Synechococcus sp. JA-2-3B	496	ACIQI	IQDSFRN	YERII	SGSPTRIN	DEKNEL	SLRI	VQNVQALEEEGL..
Synechococcus sp. JA-3-3Ab	492	SCIQI	IQDSFRN	YERII	SGSPTRIN	DEKNEL	SLRI	VQNVQSLDE.....
Synechococcus sp. PCC 7002	472	GCAAI	IQDSFRN	YERII	SGSPTRIA	DEKSELSRI	VGVQDKTLPE...	
Synechococcus sp. RCC307	475	NGPEI	IQDSFRN	YERII	SGVPHRVSD	DEKNEI	RVGVDSVDPG....	
Synechococcus sp. RS9916	471	NGPEI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	VGVNPD.....	
Synechococcus sp. RS9917	478	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	VGVVTD.....	
Synechococcus sp. WH 5701	471	NGPEI	IQDSFRN	YERII	SGVPHRVSD	DEKTELSRI	AVSFEGEESF....	
Synechococcus sp. WH 7803	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVDPDA.....	
Synechococcus sp. WH 7805	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVDPDA.....	
Synechococcus sp. WH 8016	460	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVDPDA.....	
Synechococcus sp. WH 8102	471	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVSSD.....	
Synechococcus sp. WH 8109	460	NGPQI	IQDSFRN	YERII	SGVPHRVSD	DEKSELSRI	AVGVSTED.....	
Synechocystis sp. PCC 6803	476	DGPD	IQDSFRN	YERII	SGSPTRISV	DEKSELSRI	VGVKDKTAE....	
Thermosynechococcus e. BP-1	475	KGAEI	IQDSFRN	YERII	SGTPTRISV	DEKTEL	RIAGVQDLESE...	
Trichodesmium e. IMS101	475	QCPD	IQDSFRN	YERII	SGSPTRITV	EKSELSRI	VGVQEKTTDD...	
consensus		!	!!!!	*****	!*****	*****	*****	*****

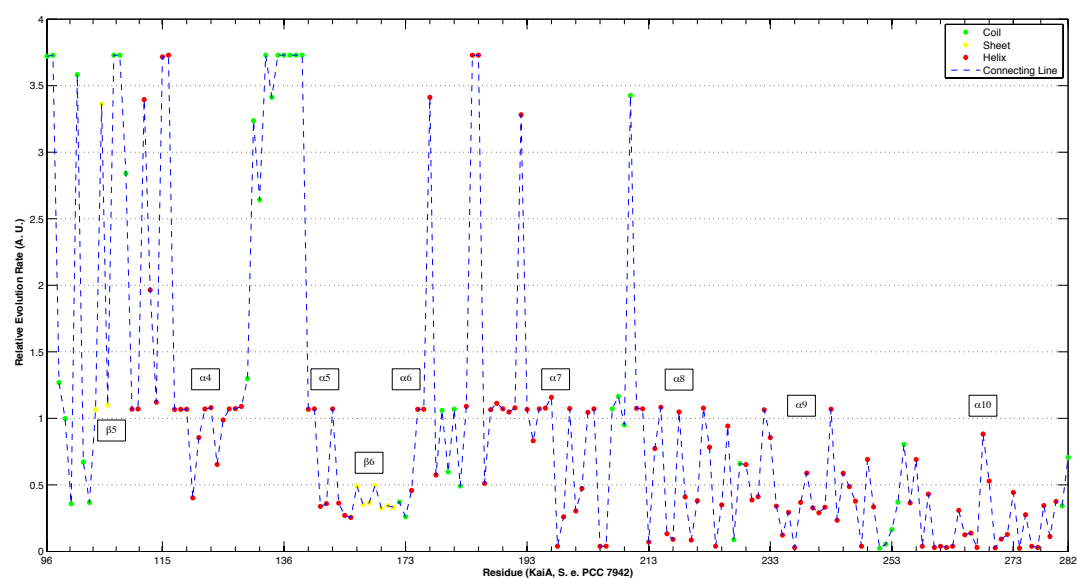
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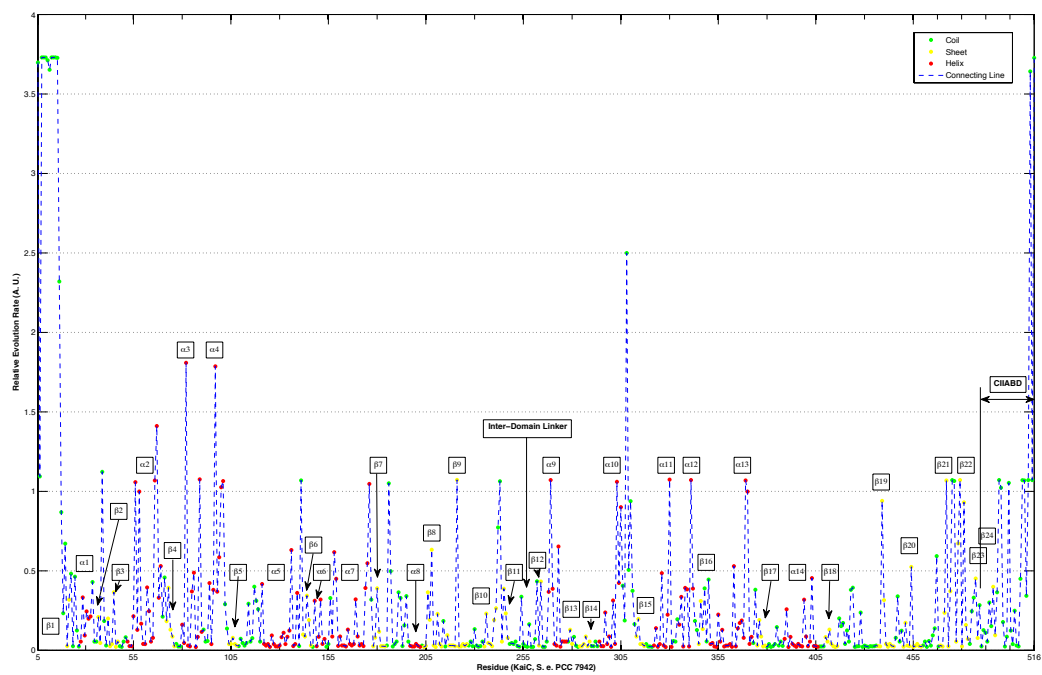
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(E)

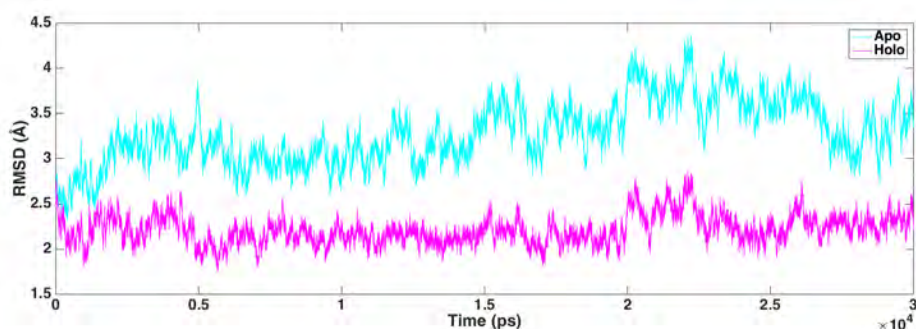


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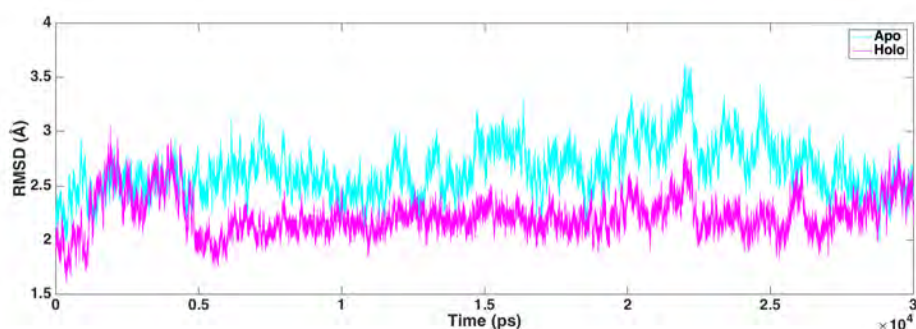


Supplementary Figure S2. The molecular dynamics simulation of the apo structure of KaiA C-terminal domains (1Q6B). (A) The backbone RMSD variation of the whole C-terminal domains compared to the apo structure (1Q6B) and the holo structure (1SV1). (B) The backbone RMSD variation of the helices 9 and 10 compared to the apo structure (1Q6B) and the holo structure (1SV1) respectively. Compared to the simulation of the holo structure described in the main text, the conformational change of the apo structure had a lower oscillation frequency between these two conformations.

(A)



(B)



Supplementary Video S1. The snapshots of the molecular dynamics simulation of the apo structure of KaiA C-terminal domains. The apo structure (1R6B, in cyan) and the holo structure (1SV1, in magenta) are shown for comparison. Only 20 ns simulations were shown.

Supplementary Video S2. The snapshots of the molecular dynamics simulation of the holo structure of KaiA C-terminal domains. The apo structure (1R6B, in cyan) and the holo structure (1SV1, in magenta) are shown for comparison. Only 20 ns simulations were shown.