

## Supplemental Figure 1

	3		98	
<i>Synechocystis</i> PCC 6803	...	KIRTVSDAKRKFTHYSRPISSIIYRRFVEELLVEMHLLSVNIDFTYDPIFALGIVTSFNSFMQGYQPAEQLPFAIFNALCHGVDQNPQVQRQDAKNV...		
<i>Prochlorococcus marinus</i> CCMP1375	...	EQTTSIDSKGLFHFKEFPYVIPVYRKRVLDEYLVELNLLSNQSNFKIDTIFSYGLIISFERFTVGYEPDSHISKILESLCNSCNIDKAIKEYSNNI...		
<i>Prochlorococcus marinus</i> CCMP1986	...	EKLTVSDSKLFLHEQFPYVIPGLYKRIVDEMLVELNLLNHQNEFIQDDLFCVGLTETTFKELTKGYKPEEHLRVLFESLCSNSNFEPPKKIKEASKKT...		
<i>Prochlorococcus marinus</i> MIT9313	...	DRKTIADSKRAFNHDFPHVIPSLYRRRTDELLELVELHLLSHQKHFHPDALFAIGLSQVFDVFTSGYRPEAHVKTLPFDALCRSCGFDPNALRQQAQQT...		
<i>Synechococcus</i> WH8102	...	ERHTIADSKRAFHQAFPHVIAPLYRRIADELLVELHLLSHQATFQANSLSFAVGLKTVFERFTQGYRPEMHPAALLSALCSSNGFDDEQLKQAAQHC...		
<i>Anabaena</i> PCC 7102	...	LLRTVSDTKRFFYALHTRPINTIYRRVVEELMVEMHLLSVNVDVSYNPIYALGVVTFDFRMEGYQPERDKESIFSAICQAVEQEPQRYRQDAERL...		
<i>Anabaena variabilis</i> ATCC29413	...	LLRTVSDTKRFFYALHTRPINTIYRRVVEELMVEMHLLSVNVDVSYNPIYALGVVTFDFRMEGYQPERDKESIFSAICQAVEQEPQRYRQDAERL...		
<i>Nostoc punctiforme</i> PCC 73102	...	NVRTVSDTKRFFYNLHTRPINTIYRRVVEELMVEMHLLSVNIDVSYNPIYALGVVTFDFRMEGYQPERDQESIFNALCRAIEQDPQHYRQDAERL...		
<i>Crocospaera watsonii</i> WH8501	...	NIRTVSDTKRFFYGYHTQPINSIYRRFVEELLVEMHLLSVNIDVSYNPIYALGVVTFDFRMEGYQPERDQESIFNALCQAVDGSSEKHYQEAEEI...		
<i>Tricodesmium ertythraeum</i> IMS101	...	NTRTVSDTKRFFYHFTRPINSIYRNVEIELLVEMHLLSVNVDVSYNPFYALGVVTFDFRMEGYQPERDQESIFNALIQQEEEDPNKYRSDAKGL...		
<i>Synechococcus elongatus</i> PCC 7942	...	SVPTVSDSKRAFYAAYPRPINFLYRRVVEELLVEMHLLSVNIDVSYNPFYALGVVTFDFRMEGYQPERDQESIFNALCQAVDGSSEKHYQEAEEI...		
<i>Thermosynechococcus elongatus</i> BP1	...	NPRTVSDTKRAFYAAHTRPIHSIYRRFIEELVEIHLRLVNVDFRYSPLFALGVVTFDFRMEGYQPEGDRDRIFHALCVAEEMNPQQLKEDAAQSW...		
<i>Gloeobacter violaceus</i> PCC 7421	...	SKRTVSDSKRAFFAAYPRPVNSIYRRVIDELLVVEHLLITNQDFRHDPLFATGLLTAYQALMEGYTPVEQRDAILRALCTALELSYQLHTDAAQW...		
<i>Solanum tuberosum</i>	...	DLPTVADTKLFLKFLTAYKRPIPTVYNTVLQELLVQQLHLMRYKSTYQYDPVDFALGFVTVYDQLMMEGYPSEEDRNAIFKAYIEALKEDPEQYRADAQKL...		
<i>Triticum aestivum</i>	...	IPPTVADTKMNFKLSYKRPISYISTVLQELLVQQLHLMRYKSTYQYDPVDFALGFVTVYDQLMMEGYPSTEDRDAIFKSYVTALNEDPEQYRADAQRM...		
<i>Oryza sativa</i> (japonica)	...	VPPTVAETKMNFLKSYKRPILSIYSTVLQELLVQQLHLMRYKSTYQYDAVDFALGFVTVYDQLMMEGYPSNEDRDAIFKAYITALNEDPEQYRADAQKM...		
<i>Chlamydomonas reinhardtii</i>	...	KPPTVAETKAKFLSGYNKPIASIYSTVLQELLVQQLHLMRYKSTYQYDAVDFALGFVTVYDQLMMEGYPSNEDRDAIFKAYITALNEDPEQYRADAQKM...		
<i>Arabidopsis thaliana</i>	...	DVPPVSETKSKFLKAYKRPIPSIYNTVLQELLVQQLHLMRYKSTYQYDAVDFALGFVTVYDQLMMEGYPSDQDRDAIFKAYIEALNEDPKYRIDAQKM...		
<i>Chlorella virus</i> PBCV-1	...	SPPTVSDTKRIFYANYKLLFLYNTPIQNMLVKQHIHRYNKNYTSVDSALGIVTTLDSVLNTPDDE-KTSIKNAFIIISLNEDEPEMYYSNIESL...		
	135	153	170	
	219			
<i>Synechocystis</i> PCC 6803	...	RHKFKYSRLFAIGLYTLA	...	LTRLSELDSLKDVVKDLDLYRSNLEKVDQLKLVLEDAEAERKKKEKQ...
<i>Prochlorococcus marinus</i> CCMP1375	...	GDKKYSSLRHLAIGIYELIS	...	ISECVEALGFSKDRVEKIDINQYKNSMEKIKEMMELIKLTVETTKRKAJLN
<i>Prochlorococcus marinus</i> CCMP1986	...	DSNLYSSRLNLNGIYILIIA	...	ISDIINKLNLNFKNAEKDIDIGIYKSSILKMEQAKELLQEAIKDKKEKKK-
<i>Prochlorococcus marinus</i> MIT9313	...	STTFHYSRLMAVGLLSLLA	...	AHELSESVGFSKARVEKDLNLYKSNLEKMAQAVELTEQILESERKKREQN...
<i>Synechococcus</i> WH8102	...	SDGAHYSRLMAVGLLALLE	...	AVKLSVDLGLPAERVEKDLTVFSSNSERMEQAVELMQETLAADRKKKEKR...
<i>Anabaena</i> PCC 7102	...	NSNFKYSRLFAIGLFTLLE	...	LKSI AAGLHLSDDKFSKDLLEYRSNLDKMTQALAVMADMLTADRKKREQR...
<i>Anabaena variabilis</i> ATCC29413	...	NPNFKYSRLFAIGLFTLLE	...	LKTI AAGLHLSDDKLSKDLLEYRSNLDKMTQALAVMADMLTADRKKREQR...
<i>Nostoc punctiforme</i> PCC 73102	...	NPNFKNRFLAIGVFSLLE	...	LKAI AAGLHVSDDKLNKDLLEYRSNLDKMAQALVVMADMLSADRKKREQR...
<i>Crocospaera watsonii</i> WH8501	...	NPKFKYSRLLAIGLYTLLM	...	IKEVSEALKFSPEKLRKDLLEYRSNLDKMQQLLTVIEDSLEADRKKRAST...
<i>Tricodesmium ertythraeum</i> IMS101	...	NSKFRYSRLFAIGLFTLLE	...	LKKICQSLNLVEKLLKIDIDLYLSNLERVAQARSAMEDTLAAMRKKREKR...
<i>Synechococcus elongatus</i> PCC 7942	...	RSEFKYSRLFAIGLFSLLE	...	VTAIAERFHLPSDKLQKDLLEYRSNLEKMEQARITMEEAIQADRKKREQR...
<i>Thermosynechococcus elongatus</i> BP1	...	NHTGKYSRLHAVGLYAFLO	...	LDQLAPVILPLPIEKVKRDLLEYRSNLDKINQARSLMKELVEQERKKRAQO...
<i>Gloeobacter violaceus</i> PCC 7421	...	AERFKYSRLFSLGLANILE	...	LQQICTYKLDYNRVVRDLDFHFSVLERIKRSKEVVDELSQTEERRKREER...
<i>Solanum tuberosum</i>	...	KDGFYSRLFAVGLFRLLE	...	LEKLCAALNVNKKSVDRDLVYRNLLSKLVQAKELLKEYVEREKKKRGER...
<i>Triticum aestivum</i>	...	KGNFSYSRFFAVGLFRLLE	...	LDKLCAALNINKKSVDRDLVYRNLLSKLVQAKELLKEYVEREKKKRGER...
<i>Oryza sativa</i> (japonica)	...	KGSFSYSRFFAVGLFRLLE	...	LDKLCAALNINKKSVDRDLVYRNLLSKLVQAKELLKEYVEREKKKRGER...
<i>Chlamydomonas reinhardtii</i>	...	AGAFSYNKFVAIGLFRLLE	...	LEKLVKAVGVKPEAVNRDLMYKGVLSKLAALKELMREFFVEREKKKQAE...
<i>Arabidopsis thaliana</i>	...	KEGFSYSRFFAVGLFRLLE	...	LDKLCASLNINKKSVDRDLVYRNLLSKLVQAKELLKEYVEREKKKRGER...
<i>Chlorella virus</i> PBCV-1	...	NDKYVYSSFAAVGIFKLLQ	...	VKHLSESIGFKGBLVHKDIATFFSLLKYIESSQKLADDIREESLKRKSKS...

WU-BLAST was used to identify proteins homologous to Psb29. The proteins identified in the analysis all had E-values < 10<sup>-14</sup>. Protein sequences were aligned using CLUSTAL-W, followed by manual alignment. Three well conserved blocks were identified and used for tree building, consisting of a total of 165 columns. Columns are numbered according to the *Synechocystis* 6803 protein sequence.