

Phenomics reveals a novel chloroplast fatty acid transporter in the marine diatom *Skeletonema marinoi* involved in temperature acclimation

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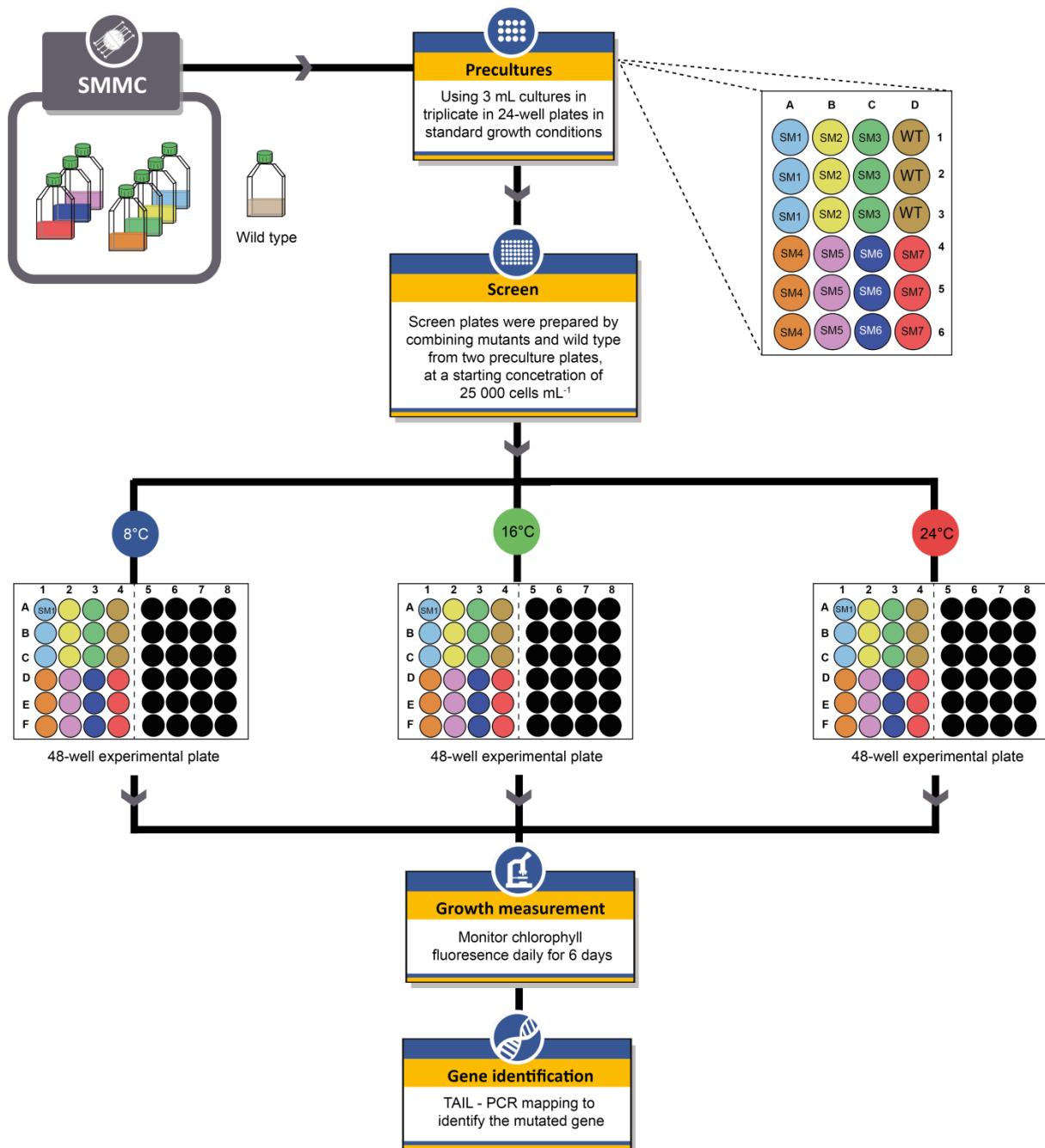
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Supplementary Figure 1. Overview of the temperature screening for *S. marinoi* mutant collection.



Supplementary Figure 2. List of primers

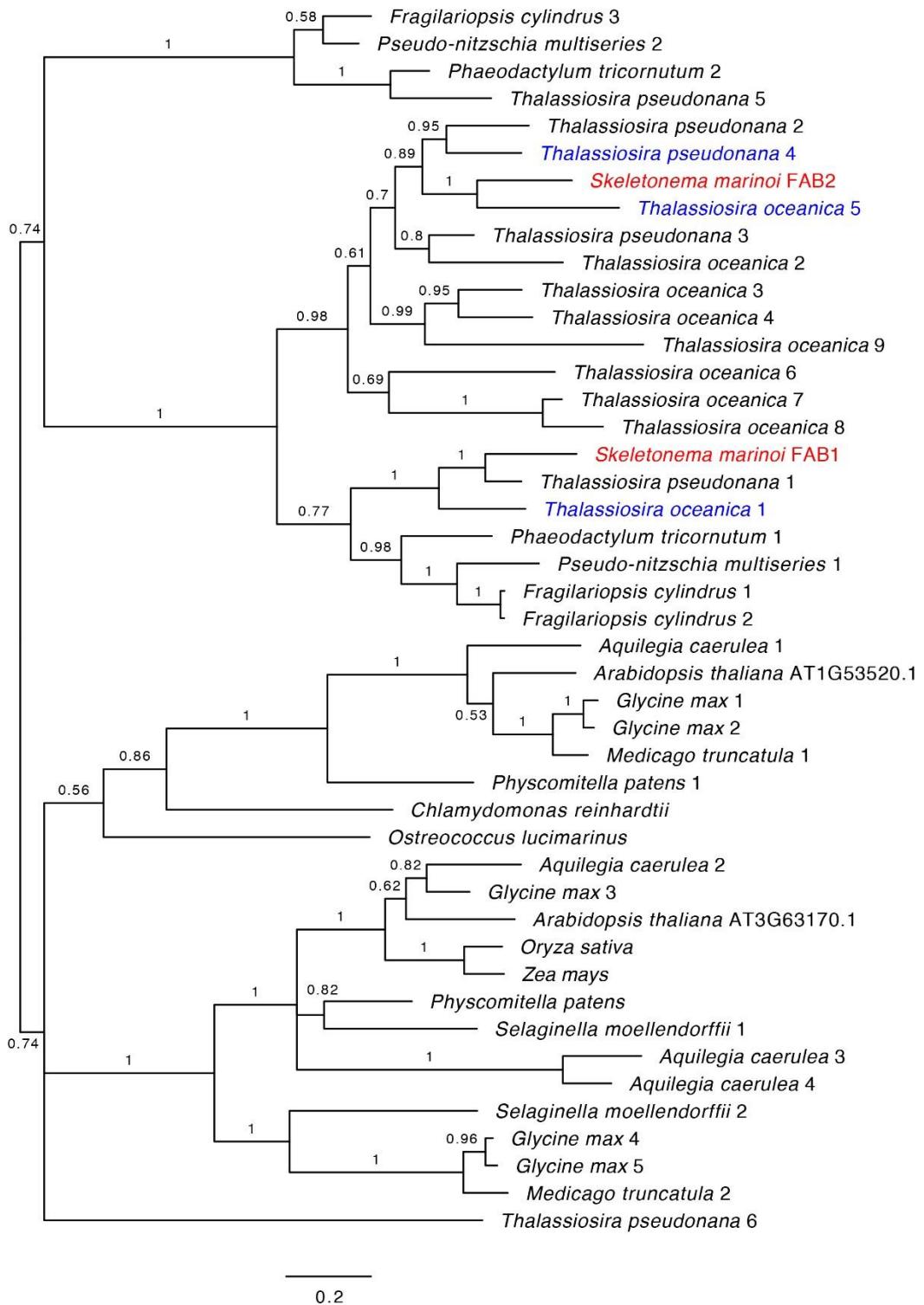
Primer	Sequence (5'->3')
TAIL-PCR	
<i>Insert specific primers</i>	
Promoter	
pISP[1]	CCTGCGAAATCATCCTCCACAAAATCTG
pISP[2]	GAGAATCCGAGACGATCTGTCCAGAACTC
pISP[3]	AGCACTGGAACAGCACTTGTCAACTTTGC
Terminator	
tISP[1]	CGAGAATTGTGGAGGATGATTCGCAGG
tISP[2]	GAATGGTCTGAAGTAGTGTCTACAAATTCCGTG
tISP[3]	GCGTGCATTCGTGGCAGAAGAGCAAG
<i>Degenerate primers</i>	
Fusion-1	GGGCACGTCTACTATAGGGCACGCGTGGTNTCGASTWTSGWGTT
Fusion-2	GGGCACGTCTACTATAGGGCACGCGTGGTNGTCGASWGANAWGAA
Fusion-3	GGGCACGTCTACTATAGGGCACGCGTGGTWGTGNAGWANCANAGA
Fusion-4	GGGCACGTCTACTATAGGGCACGCGTGGTAGWGNAGWANCAWAGG
Fusion-5	GGGCACGTCTACTATAGGGCACGCGTGGTNGTAWAASGTNTSCAA
Fusion-6	GGGCACGTCTACTATAGGGCACGCGTGGTNGACGASWGANAWGAA
Fusion-7	GGGCACGTCTACTATAGGGCACGCGTGGTGTNCGASWCANAWGTT
Fusion-8	GGGCACGTCTACTATAGGGCACGCGTGGTNCAGCTWSCTNTSCTT
<i>Amplicon specific primer</i>	
ASP1	GGGCACGTCTACTATAGGGC
ASP2	ACTATAGGGCACGCGTGGT
Genotyping	
SM127-fwd	ACTGGACAAAAATTACTGCAAAGCAAGAGACGGC
SM127-rev	ATCCTATCTCGTCCATCAACAGATACTTGACTCCC
qRT-PCR	
q_Lsu4e_Fwd	CATGGGAACTGGACGTGCCG
q_Lsu4e_Rev	GGTGTACTTACGGCCACGGGC
q_fat1_Fwd	AACAACTTATTCTTGATGGCGTAGCGG
q_fat1_Rev	CGCTCCTGCGATAGTTTGATCC

Supplementary Figure 3. Thermal conditions for the modified TAIL-PCR approach used to map the genomic insertion sites in *S. marinoi* transformants.

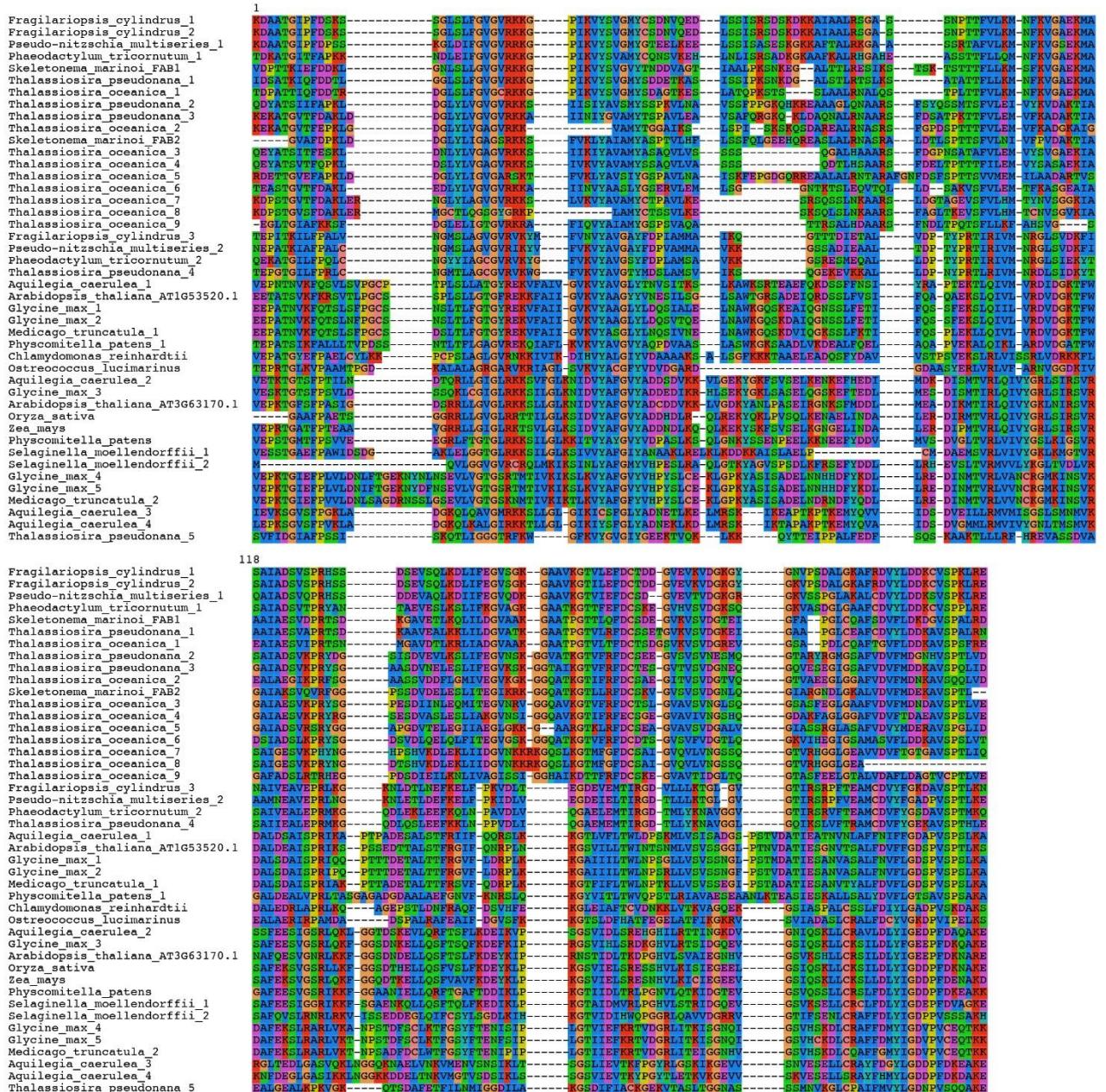
Primary reaction (1°)			Secondary reaction (2°)			Tertiary reaction (3°)		
Step	Temp.	Time	Step	Temp.	Time	Step	Temp.	Time
1	96°C	5 min	1	98°C	20 s	1	98°C	30 s
2	98°C	10 s	2	65°C	1 min	2	98°C	10 s
3	70°C	15 s	3	72°C	2 min	3	57°C	15 s
4	72°C	2 min	4	Go to 1	1 x	4	72°C	2 min
5	Go to 2	30 x	5	94°C	20 s	5	Go to 2	14 x
6	98°C	10 s	6	68°C	1 min	6	72°C	5 min
7	25°C	1 min	7	72°C	2 min	7	4°C	∞
8	72°C	2 min	8	94°C	20 s			
	ramp 0.5 °C / sec							
9	Go to 6	1 x	9	68°C	1 min			
10	98°C	10 s	10	72°C	2 min			
11	70°C	15 s	11	94°C	20 s			
12	72°C	2 min	12	55°C	1 min			
13	98°C	10 s	13	72°C	2 min			
14	70°C	15 s	15	72°C	5 min			
15	72°C	2 min	16	4°C	∞			
16	98°C	10 s						
17	40°C	15 s						
18	72°C	2 min						
19	Go to 10	10 x						
21	72°C	5 min						
22	4°C	∞						



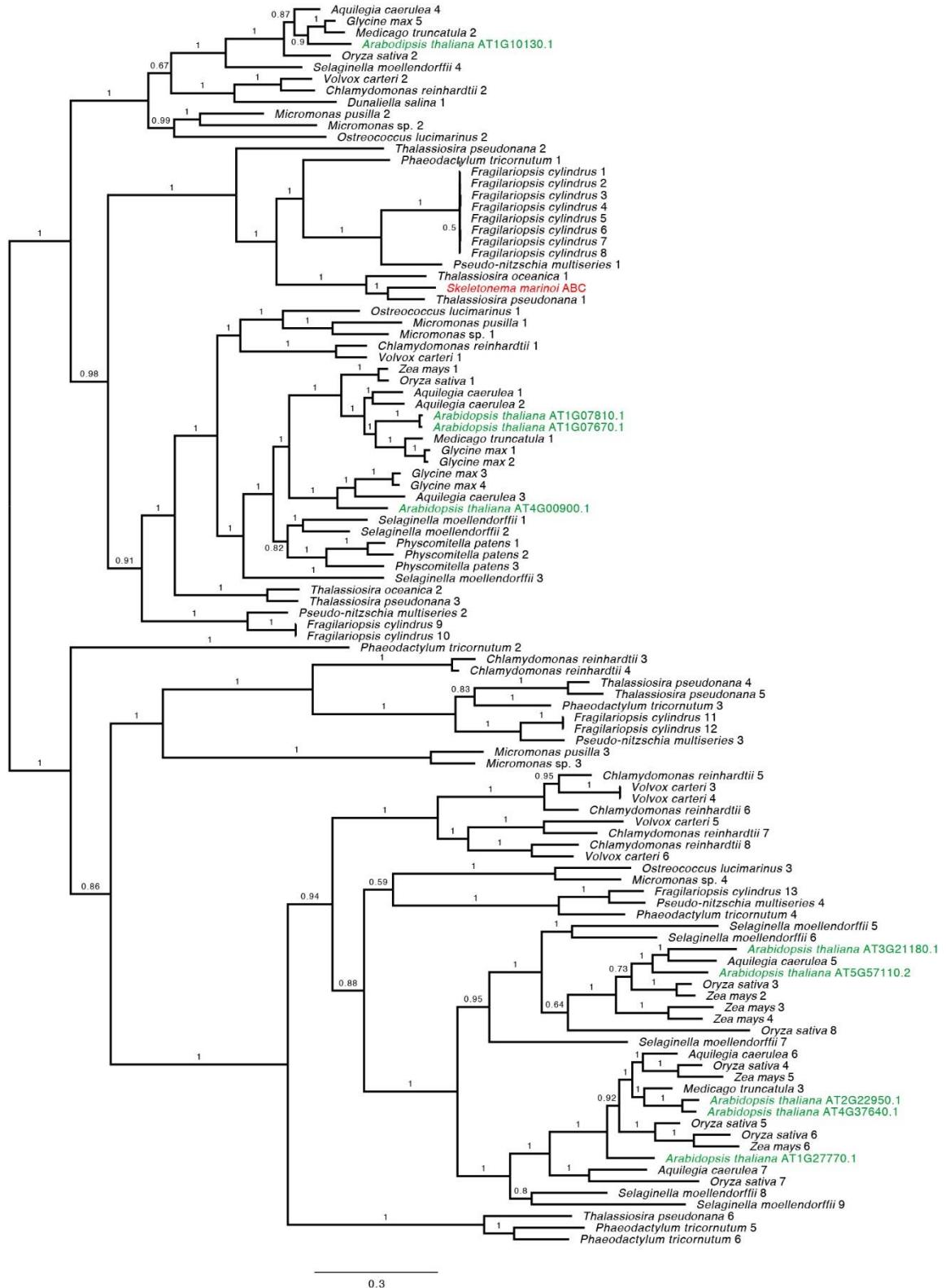
Supplementary Figure 4. Amino acid sequence alignment of the *S. marinoi* FAT1 protein with the orthologs in *T. oceanica* and *T. pseudonana*. The sequence from *T. pseudonana* consists of several ORF predictions (generated using the program getorf in the EMBOSS v6.3.1 package) grafted together after putative intron sequences were removed. Conserved sites are indicated by a black background, and similar amino acids by a grey background.



Supplementary Figure 5. Unrooted phylogenetic tree of the fatty acid-binding (FAB) domains presented in supplementary table SX. Highlighted in red are the two FAB regions found in the *S. marinoi* FAT1 protein. Highlighted in blue are the three domains from *T. oceanica* and *T. pseudonana* that are part of the proteins shown in Supplementary Figure 4. Posterior probability values appear above the branches, and the expected number of changes per site along the branches is indicated by the scale bar.



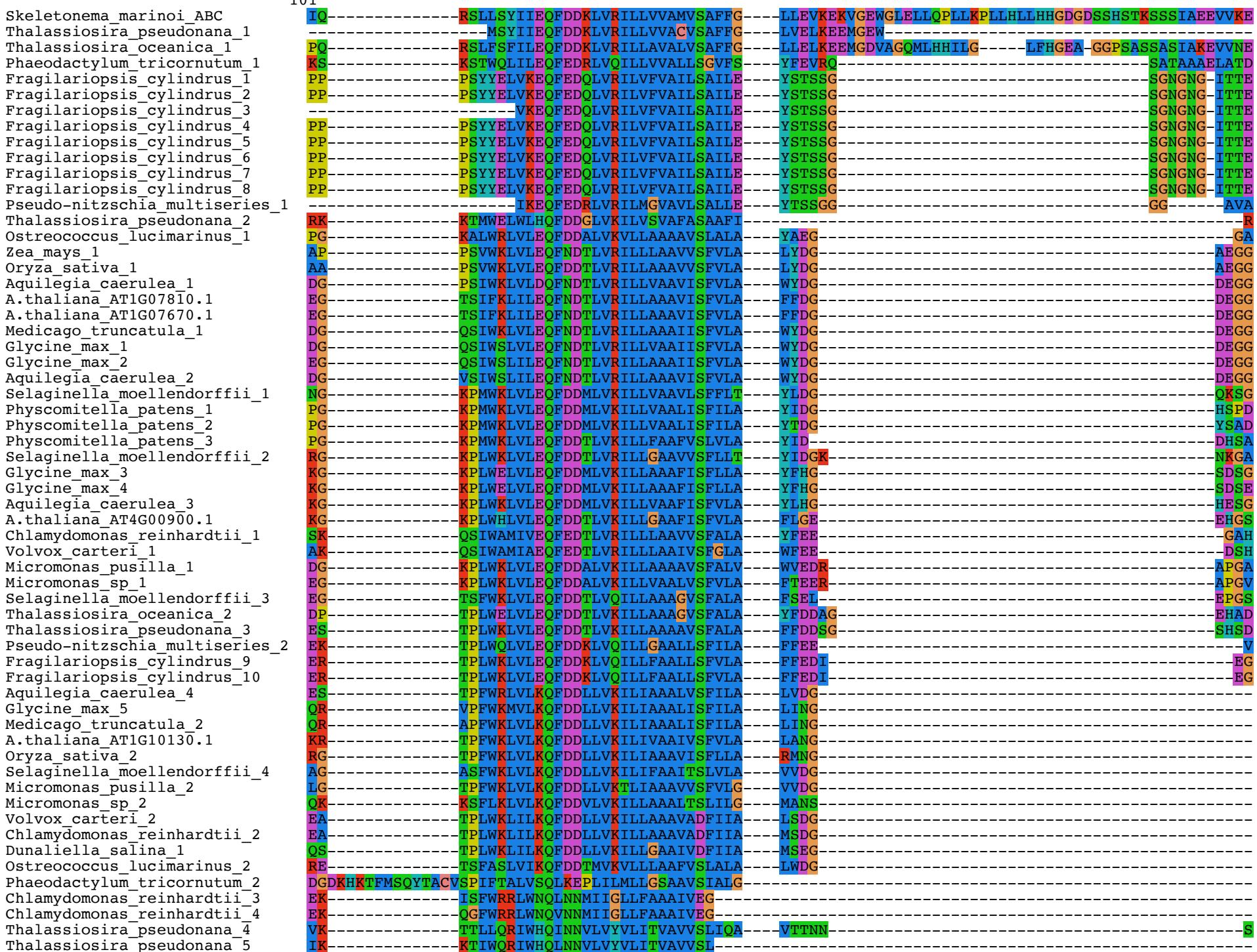
Supplementary Figure 6. Alignment of the FAB domains identified in green algae, land plants and diatoms and used for the phylogenetic analysis presented in Supplementary Figure 5. The length of the sequences has been adjusted to the length of the two domains found in *S. marinoi*. Numbers above the sequences indicate the positions in the alignment and the background color of the letters indicates size, shape, solubility, and ionization properties of the different amino acids.



Supplementary Figure 7. Unrooted phylogenetic tree of the ABC transporter domains presented in Supplementary Table SX. The position of the *S. marinii* sequence is highlighted in red and eight homologous domains found in *Arabidopsis thaliana* are highlighted in green. Posterior probability values appear above the branches, and the expected number of changes per site along the branches is indicated by the scale bar.

Skeletonema_marinoi_ABC	1	YLSVPTS-DDGO	V-IGLTS	QDRHNRLKYIFGPN	ELEQP
Thalassiosira_pseudonana_1		YLNVPAL-DKG	KTLLGLTE	QDRQRKV-AVYGKN	ELDQP
Thalassiosira_oceanica_1			EGLSE	SOASARI-VQFGKN	SLEQS
Phaeodactylum_tricornutum_1		TT-TT-DDNY	LF-NGLSS	TKAKOLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_1			LF-NGLSS	TKAKQLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_2					
Fragilariopsis_cylindrus_3					
Fragilariopsis_cylindrus_4		TT-TT-DDNY	LF-NGLSS	TKAKOLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_5		TT-TT-DDNY	LF-NGLSS	TKAKOLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_6			GLSS	TKAKOLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_7		TT-TT-DDNY	LF-NGLSS	TKAKOLL-EIYGKN	ELQDL
Fragilariopsis_cylindrus_8				RIYGKN	ELQDL
Pseudo-nitzschia_multiseries_1					
Thalassiosira_pseudonana_2			MH-TGLSN	OOASALL-SQIGPN	SLOPP
Ostreococcus_lucimarinus_1		D	ET-VGIAS	ASVDDRRARAGGHN	ELERE
Zea_mays_1		GV	AD-RGLSS	EEAAARL-ORHGPN	ELERH
Oryza_sativa_1		GV	A	EEAAARL-RRYGPN	ELERH
Aquilegia_caerulea_1		KV	AD-RGLSS	QEVEKRO-QIYGLN	ELEKH
A.thaliana_AT1G07810.1		VV	KH-QGLTT	DEVILKRH-QIYGLN	ELEKP
A.thaliana_AT1G07670.1		GV	RE-KGLSS	DEVLKRH-QIYGLN	ELEKP
Medicago_truncatula_1		KV	RE-KGLST	DEVENRNR-KIYGFN	ELEKH
Glycine_max_1		KV	VK-TGLSH	DEVENRNR-KIHGLN	ELEKH
Glycine_max_2		KV	VK-VGLNH	DEVEERR-KIYGLN	ELEKH
Aquilegia_caerulea_2		KV	N	DEVEERR-KIHGLN	ELEKH
Selaginella_moellendorffii_1		IT	KE-LGLRV	SSVEKRR-ELYAWN	ELEKE
Physcomitella_patens_1		EV	LA-HGLSK	SSIAKRR-KOYAWN	ELEKE
Physcomitella_patens_2		EV	VR-SGLSQ	SSIENRR-KOYAWN	ELEKE
Physcomitella_patens_3		DV	VK-SGLSC	SVVQSSR-AKYAWN	ELOKE
Selaginella_moellendorffii_2		GV	PK-KGLSK	EEVIARR-ERYAWN	ELKRE
Glycine_max_3		GV	PS-KGLSG	YEVQKRL-EKYGMN	ELAKE
Glycine_max_4		GV	K	YEVQKRL-EKYGMN	ELAKE
Aquilegia_caerulea_3		NV	LD-KGLST	YDVENRK-EKCGWN	ELDKQ
A.thaliana_AT4G00900.1		KT	LE-KGLSS	EDVQIRR-QKYGMN	ELAKE
Chlamydomonas_reinhardtii_1		DV	LD-KGLTS	QKVEEKR-ATYGYN	ELEKE
Volvox_carteri_1		DV	PK-TGLTD	EKVAARR-AQYGMN	ELEKE
Micromonas_pusilla_1		GV	VK-EGLTD	AAARDAL-ARHGPN	ELDKE
Micromonas_sp_1		GV	D	TKIAKLR-SEYGFN	ELDKE
Selaginella_moellendorffii_3		GV	PA-NGLTS	DAIDKKR-TTYAWN	ELQKP
Thalassiosira_oceanica_2		GV	PE-TGLDE	AQVEARR-AEYAWN	ELDKE
Thalassiosira_pseudonana_3		DV	IS-KGLNF	AEVEAKR-EEFAWN	ELDKE
Pseudo-nitzschia_multiseries_2		KV	T	ARVEELR-KIHGTN	ELEHE
Fragilariopsis_cylindrus_9		HT	LE-QGLSE	DRVHALR-ATYGSN	ELEKE
Fragilariopsis_cylindrus_10		HT	D	DRVHALR-ATYGSN	ELEKE
Aquilegia_caerulea_4		GV	LK-SGLST	SOVANNA-RIYGRN	VLPQE
Glycine_max_5		GV	QT-KGLND	AEVVQHA-RLYGKN	VLAED
Medicago_truncatula_2		GV	PT-KGLSD	TOVAQHG-RLYGTN	VLHED
A.thaliana_AT1G10130.1		GV	PT-KGLSD	SQVHHHS-RLYGRN	VLPEE
Oryza_sativa_2		GV	PT-KGLSD	FOVASGS-KLLVKSCCWWSMIGFTAKTSCPKKNITNTPSVYMSVLILTDQA	
Selaginella_moellendorffii_4		GV	PS-HGLAD	SOVFEEAR-SLYGCN	GMLQI
Micromonas_pusilla_2		GV	PA-VGLSD	REAAALR-ARHGPN	EMPPE
Micromonas_sp_2		AV	PS-EGLSA	DDVWRLR-RTWGRN	ELSKG
Volvox_carteri_2		NV	LD-AGLSD	TDVFKAR-SRYGRN	ELAPE
Chlamydomonas_reinhardtii_2		NV	LD-KGLSD	TDVFKAR-TRYGRN	ELAPE
Dunaliella_salina_1		HV	LD-KGLSD	RDVQQAR-IKYGRN	OMEAE
Ostreococcus_lucimarinus_2		GV	LA-HGLDR	ADVORRR-DACGAN	ALPAQ
Phaeodactylum_tricornutum_2		SV	SSSTOLLS-HGWST	ROIISTLR-LEFGAN	RMPGD
Chlamydomonas_reinhardtii_3		GS	S-NGLSD	SEAQRRL-QLFGPN	KLTEV

Chlamydomonas_reinhardtii_4	--GT-S	--AE-NGLIST	--AEEERRI	--QDFGPN	--KLTEV
Thalassiosira_pseudonana_4	--GLKE-D	--LPK-AGLST	--DEAASRL	--TKYGPNA	--QMTEK
Thalassiosira_pseudonana_5	--SLKD-D	--LPK-MGLST	--EEACDRL	--AKYGPNA	--OMAEK
Fragilariaopsis_cylindrus_11	--GLPS-D	--IRK-KGLTA	--AAQARL	--EKYGEN	--KLTTET
Fragilariaopsis_cylindrus_12	--GLPS-D	--IRK-KGLTA	--AAQARL	--EKYGEN	--KLTTET
Pseudo-nitzschia_multiseries_3	--DLPA-D	--IRK-IGLTT	--AQAKERL	--EKYGEN	--KLTEK
Phaeodactylum_tricornutum_3	--GCSP-N	--SRR-EGLTS	--QEAKARL	--ERYGPNA	--OLTEK
Micromonas_pusilla_3	--QT-D	--KV-KGLTA	--AAAKRL	--LEDGPNA	--ELEKP
Micromonas_sp_3	--QT-N	--AE-TGLSA	--AAAKRL	--EEDGPNA	--ELEKP
Chlamydomonas_reinhardtii_5	--VT-D	--LH-AGLNE	--SPAAAAAADGLATAAAASELGSVEAHR	--KAYGEN	--KFPEK
Chlamydomonas_reinhardtii_6	--GT-D	--LK-EGLSD	--AGDSSK	--QAFGVN	--SFPEK
Volvox_carteri_3	--CT-D	--LH-HGLTE	--ODDKTGVNAHR	--AAYGPN	--TFPEK
Volvox_carteri_4	--CT-D	--LH-HGLTE	--ODDKTGVNAHR	--AAYGPN	--TFPEK
Volvox_carteri_5	--GS-S	--PD-RGLNP	--TGHDVSSEHR	--RIFGAN	--KHAEV
Chlamydomonas_reinhardtii_7	--SS-S	--VE-SGLNA	--AGDDSVLEHR	--RVFGEN	--KHAET
Chlamydomonas_reinhardtii_8	--AT-S	--LH-EGLDP	--STVDAHA	--EAYGN	--KFKET
Volvox_carteri_6	--LS-D	--LH-KGLDP	--QOGLASIEAHV	--DAYGEN	--KFPEV
Ostreococcus_lucimarinus_3	--GC-D	--LK-RGLCD	--WASEERK	--ESYGVN	--EFEYP
Micromonas_sp_4	--KT-D	--PK-VGLCGT	--ESLARRK	--EAFGVN	--EFEYP
Selaginella_moellendorffii_5	--ST-S	--TK-NGIED	--PKIERR	--LLYGSN	--TYPQQ
A.thaliana_AT3G21180.1	--KS-N	--ME-QGINE	--KEVIDRK	--NAFGSN	--TYPKK
Aquilegia_caerulea_5	--KT-N	--PE-KGING	--ADILNRK	--NTFGSN	--TYPKK
A.thaliana_AT5G57110.2	--KT-N	--PE-KGISG	--DLLLKRK	--TIYGSN	--TYPRK
Oryza_sativa_3	--KT-D	--TE-KGISG	--SDLTARR	--NAFGSN	--TYPRK
Zea_mays_2	--KT-D	--TO-KGISG	--SDLLARK	--NAFGSN	--TYPRK
Zea_mays_3	--KS-D	--LD-RGLSP	--SELMRRR	--DIFGAN	--TYPRK
Zea_mays_4	--KS-N	--LE-KGVSP	--DELLQRK	--NVYGSN	--TYPRK
Selaginella_moellendorffii_6	--HV-N	--LE-KGIEP	--ESVQHRR	--EAFGAN	--SYPTK
Selaginella_moellendorffii_7	--KI-D	--PO-KGIDA	--VDIKARR	--DAFGPN	--TYPLK
Aquilegia_caerulea_6	--CT-S	--TA-NGLTT	--DILSRQ	--EIYGIN	--KFTES
Medicago_truncatula_3	--ST-S	--AT-EGISN	--DILDKRQ	--QIYGIN	--KFTES
A.thaliana_AT2G22950.1	--KA-C	--PN-AGLSTG	--EOLSKRQ	--ELFGIN	--KFAES
A.thaliana_AT4G37640.1	--KA-S	--PT-DGLST	--AQLSQRQ	--ELFGIN	--KFAES
Oryza_sativa_4	--CT-S	--PE-DGLPK	--RROAVRE	--ELFGIN	--RFAET
Zea_mays_5	--ST-S	--AS-DGLDDS	--PMTAARQ	--ELFGVN	--RFAEA
Oryza_sativa_5	--AT-S	--PA-DGLST	--ESIKRRQ	--DVYGLN	--KFTES
A.thaliana_AT1G27770.1	--ST-S	--IA-SGIST	--DILSVRK	--EIYGIN	--QFTES
Oryza_sativa_6	--GT-S	--LT-NGIVT	--DILNQRQ	--DIYGVN	--KFAET
Zea_mays_6	--AT-S	--LA-DGITT	--LSLNQRQ	--GMYGVN	--KFTES
Aquilegia_caerulea_7	--SV-S	--LD-DGVSE	--KDLPRRQ	--EIYGFN	--RYVEK
Oryza_sativa_7	--KA-S	--LE-DGAKE	--TDIATRQ	--MLYGAN	--RHAEK
Selaginella_moellendorffii_8	--LV-S	--LD-DGVSK	--DEIDKRK	--EAFGSN	--MYEEK
Selaginella_moellendorffii_9	--HVH-G	--IE-HGIDP	--SELLARR	--RAFGSN	--TYKES
Oryza_sativa_8	--AS-G	--AE-RGIRG	--ADVARRK	--KAFGSN	--TYPKP
Fragilariaopsis_cylindrus_13	--AT-H	--PD-HGAAE	--TSINKRQ	--ELFGSN	--LLPSS
Pseudo-nitzschia_multiseries_4	--AT-N	--RD-HGAAE	--STIETRR	--ETFGSN	--VLPST
Phaeodactylum_tricornutum_4	--GT-D	--PK-AGLDR	--ETIETRR	--ACFGAN	--RLPSA
Thalassiosira_pseudonana_6	--RS-K	--PE-DGITN	--SDIEFRR	--EAFGTN	--AIADK
Phaeodactylum_tricornutum_5	--RS-S	--PE-VGIFF	--SEVEKRR	--EAFGSN	--RIAPK
Phaeodactylum_tricornutum_6	--RS-S	--PE-SGIDP	--REVEHRO	--SVFGSN	--AIAAK



Fragilariopsis_cylindrus_11	EK	EGLLKKIW NQVNNILVLILVIVAVI SVVTAA	FVIPS	AV
Fragilariopsis_cylindrus_12	EK	EGLLKKIW NQVNNILVLILVIVAVI SVVTAA	FVIPS	AV
Pseudo-nitzschia_multiseries_3	EK	ETLLQK IWNQVNNILVLILVIVAVI SLISA	FVIPP	DI
Phaeodactylum_tricornutum_3	EK	VTLQ RQIWKQVNVLGV GILVFVAVVSLAKG	IST	SG
Micromonas_pusilla_3	PR	VSSLVLFLIQ LNSVIMYLLM GAVVASAAIKATG		DD
Micromonas_sp_3	PR	VSSLVLFLIQ LNSVIMYLLM AAAVASAAIKATG		DD
Chlamydomonas_reinhardtii_5	PP	PNFFMMLLE AQDPMIIILLI VAIVTIVLGAA	VP	E
Chlamydomonas_reinhardtii_6	PP	PSFL SMILLEAKDPMIVILLI VAVTIVLGAA	VP	E
Volvox_carteri_3	PP	PSFL SMILLEASKDPMIILLI VVALVTIVLGAA	VP	E
Volvox_carteri_4	PP	PSFL SMILLEASKDPMIILLI VVALVTIVLGAA	VP	E
Volvox_carteri_5	PP	KNFFVLVWE VVQDPILILLIAAAAV STILGSA	IP	E
Chlamydomonas_reinhardtii_7	PP	KNFFFLVWE VVQDPILILLIAAAAT TVSTVLGAA	IP	E
Chlamydomonas_reinhardtii_8	PP	KSSFL SVLWENLQDPVIII CVAAAVSTALGAA	IP	E
Volvox_carteri_6	PP	KSF FLALVWGNLQDPVIII IAALVSTILGAA	IA	E
Ostreococcus_lucimarinus_3	PP	KSF WEI CKDALGDLTVRILIAASVVS ¹ SLAVGAG	MK	S
Micromonas_sp_4	PP	KSF QLCRDALDDITVQII CVAIIISLGIGAG	LP	K
Selaginella_moellendorffii_5	SP	KGFLAFLWEACQDLTLV ILGVC AVVSLALA	LA	T
A.thaliana_AT3G21180.1	KG	KNFFMFLWEAWQD DLTLI IAAVTSLALG	IK	T
Aquilegia_caerulea_5	KG	RSFL FLWEAQD DLTLIILMVAAAASLALG	IK	T
A.thaliana_AT5G57110.2	KG	KGFLRFLWDACHDLTLIILMVAAAASLALG	IK	T
Oryza_sativa_3	KG	RSFLAFLWDA CKDLTLI ILMVAAAASLALG	IT	T
Zea_mays_2	KG	RSFLAFVWDACKDLTLIILMVAAAASLALG	IT	T
Zea_mays_3	ER	RSIWHFVFE ACQDLTLV ILMVAAAASFSLG	MA	T
Zea_mays_4	KR	KNILRFVFE ACQDLTLV ILMIAAAASLTLG	MT	T
Selaginella_moellendorffii_6	AG	KSF W		
Selaginella_moellendorffii_7	KR	TPF YMYVWEALQDE TLMLIILCAIVSLAVG	LT	T
Aquilegia_caerulea_6	EV	RSFWFVFW WEALQDM TLMLGVC ¹ SLIVG	LI	M
Medicago_truncatula_3	QA	KSF WWVWEALQDM TLMLGVC ¹ SLIVG	IA	T
A.thaliana_AT2G22950.1	EL	RSFWFVFW WEALQDM TLMLGVC ¹ SLIVG	IA	T
A.thaliana_AT4G37640.1	EM	RGFWFVFW EAQDM TLMLGVC ¹ SLIVG	IA	T
Oryza_sativa_4	ES	RSFWFVFW EAQDM TLMLAACAFFS ¹ LVVG	IA	T
Zea_mays_5	EP	RSFWFVFW EAQDM TLMLAACALVS ¹ LVVG	IA	T
Oryza_sativa_5	EV	RSFWFVFW EAQDT TLMLAACACAFV ¹ SLIVG	IA	M
A.thaliana_AT1G27770.1	PS	RGFWLFVWE EAQDT TLMLAACACAFV ¹ SLIVG	IL	M
Oryza_sativa_6	FI	RSFWEFVWE EALED TTLIL SACAI FS ¹ LVVG	IT	T
Zea_mays_6	EA	RSLWEFW WEALQDT TLVILIACALVS ¹ LVVG	IA	T
Aquilegia_caerulea_7	PS	KGFFMFVWE ALQDL TLMLVMCAVVS ¹ IVVG	LA	T
Oryza_sativa_7	PP	RSFWMFVWDALHDLTLIILVV ¹ CALVS ¹ IVVG	LA	T
Selaginella_moellendorffii_8	PP	KGF WFVWEAMHD TLAILGFC ¹ LSL ¹ VG	VL	T
Selaginella_moellendorffii_9	PQ	RSVFSYILDASQDLTLLILVV ¹ CALVS ¹ IVVG	IA	T
Oryza_sativa_8	KP	KGFFR HVWDALAD VFLIVLLVCAAVSLA ¹ FG	IK	E
Fragilariopsis_cylindrus_13	PR	KSFYQLFIDTFDDATLQILIVAAIVS ¹ LCVG	MY	
Pseudo-nitzschia_multiseries_4	PR	OSFWOLFVD IDTFDDATLQILIVAAIVS LA ¹ FG	IY	
Phaeodactylum_tricornutum_4	PR	KTFGQLFLD IDTFDDATLQILIVAAIVS LA ¹ VG	LY	
Thalassiosira_pseudonana_6	KL	DSFLKLCWN ¹ AVODFVLIMLIVLGVI GIVVETT IGLD		PG
Phaeodactylum_tricornutum_5	KI	ESFC ¹ KL ¹ CWN ¹ AVODFVLIMLIVLGVI GIVVETT IGLD		ND
Phaeodactylum_tricornutum_6	SL	DSFFRLCWEAVQDFVLIMLIVLGIVSIVIEVG ¹ GT	LE	DG

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Skeletonema_marinoi_ABC	AKNILSGKT-E SKELND FG IK D LLE AL VE PI V ITT IL VINA LV GGY QSL N ASKG I SALK EM QASK A	VVRVC-RGG	NDVDEVELDASSLVPGDVVR
Thalassiosira_pseudonana_1	-AL VE PI V ITT IL I NAL VGGY QSL N ASKG I SALK OM QA QKA	-	SAIDEVEVDASSLVPGDVVI
Thalassiosira_oceanica_1	ATTIVTGHSDDDTKLHKIGIKH VIE AL VE PI V ITT IL VINA LV GGY QSL D ASKG I SALK SM QADKA	VIRVS-SGDR	STFDEVEVDSSSLVPGDITVV
Phaeodactylum_tricornutum_1	EA-LWKS FVE PLV ILLA I L VVNA AVVGVWQSKASGSIEALKAMQPNIC	TVIR	DGVWKSSLEASDLVPGDIIE
Fragilaropsis_cylindrus_1	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_2	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_3	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_4	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_5	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_6	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_7	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Fragilaropsis_cylindrus_8	SS-IWHRFAEPVVIISSLI I NAVVGVWQSKASGSIEALKAMQPNIC	TVIRN	NGTEYNDYPSCDLVVGDIII
Pseudo-nitzschia_multiseries_1	ES-LWHRFAEPPIVIISSLI VINASAVGVWQSKASGSIEALKAMQPSVC	-	TAAEIIDYPSADLVPGD LIV
Thalassiosira_pseudonana_2	ST-ILQS FVE PFI VAI VAILLNACVGWVQDL SARSS LEALKM KQPRKA	TVLRYDED TNNN YSDWI TDY DATQLVPGDIIR	-
Ostreococcus_lucimarinus_1	EE-GLAA YAE PAVIAL I L VNA I VGWVQ E SNA E KALE AL KE I QSE HA	KCLR	DGRWNGS LEAREL VP GD VVE
Zea_mays_1	EV-GLTA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	TVRR	DGRWSHGLPARDLVPGDIVE
Oryza_sativa_1	EM-GATA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	TVKR	DGRWSHGLPARDLVPGDIVE
Aquilegia_caerulea_1	EM-GITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	NVIR	NGKRI PSLPAKELVPGDIVE
A.thaliana_AT1G07810.1	EM-GITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSQQA	TVMR	DGTKVSSLPAKELVPGDIVE
A.thaliana_AT1G07670.1	EM-GITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSQQA	TVMR	DGTKVSSLPAKELVPGDIVE
Medicago_truncatula_1	EM-EITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE QA	SVIR	NNEKIPSLPAKDLVPGDIVE
Glycine_max_1	EM-EITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	VVIR	EGAKIPNLPAKELVPGDIVE
Glycine_max_2	EM-EITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	VVIR	EGAKISNLPAKELVPGDIVE
Aquilegia_caerulea_2	EM-EITA FVE PLV IFL I L I VNA I VGWVQ E SNA E KALE AL KE I QSE HA	TVIR	EGKKIHNLPAKELVPGDIVE
Selaginella_moellendorffii_1	ES-DLTAYVEPLV IFL I L I I NAVVGVWQ E TNA E SA E LEALKM QPEHA	KVLR	DGHWIPDLAARELVPGDVVE
Physcomitella_patens_1	TS-GIGAYVEPLV IFL I L I I NAVVGVWQ E SNA E NA E AL KEM QSENA	KVFR	DGKYISDL PARELVPGDIVE
Physcomitella_patens_2	ES-GIGAYVEPLV IFL I L I I NAVVGVWQ E SNA E NA E AL KEM QSENA	KVFR	DGEFISNL PARELVPGDIVE
Physcomitella_patens_3	EE-GATAYVEPMV IFL I L I I NAVVGVWQ E SNA E NA E AL KEM QSAQA	EVIR	DGVAISDL PARELVPGDIVE
Selaginella_moellendorffii_2	ES-ELTAYVEPLV IFL I L I I NAVVGVWQ E SNA E AL KEM QPENA	KVLR	DGOLLGDL PARELVPGDVVE
Glycine_max_3	ES-GFEAYVEPLV IFL I L I I L VNA I VGWVQ E NNA E KALE AL KE LQESGG	KVLR	DGYFVPDLPAKELVPGDIVE
Glycine_max_4	ES-GFEAYVEPLV IFL I L I I L VNA I VGWVQ E NNA E KALE AL KE LQESGG	KVLR	DGYFVPDLPARELVPGDIVE
Aquilegia_caerulea_3	EA-GFEAYVEPFV IFL I L I I L VNA I VGWVQ E SNA E KALE AL KE LQESA	KVLR	DGYYVPDL PARELVPGDIVE
A.thaliana_AT4G00900.1	GS-GFEAFVEPFV IFL I L I I L VNA I VGWVQ E SNA E KALE AL KEM QCESA	KVLR	DGNVLPNLPARELVPGDIVE
Chlamydomonas_reinhardtii_1	EE-GLRAFIEPLV IFL I L I I L NAGVGWVQ E SNA E SA E LEALKL QETETA	HVTR	NGKMVSDLPSRELLPGDIVH
Volvox_carteri_1	EE-GIRAFIEPLV IFL I L I I L NLA I VGWVQ E SNA E SA E LEALKL QETETA	HVTR	NGKLLSDL PARELVPGDVVE
Micromonas_pusilla_1	PI-DLVDFVEPGV IFL I L I I L NLA I VGWVQ E SNA E NA E AL KEM QSDTA	RVLR	DGKWDHAFOQARDLVPGDVVE
Micromonas_sp_1	EL-SLVD FVE PGV IFL I L I I L NLA I VGWVQ E SNA E SA E AL KEM QSETA	RCLR	AGEWISDL PARELVPGDVVE
Selaginella_moellendorffii_3	KL-GPGAFTEPLV IFL I L I I L N A E K A S T L Q A L K E M Q S E E A	RVLR	DGKEIVDL PARELVPGDIVE
Thalassiosira_oceanica_2	EE-GILAYIEPLV IFL I L I I L N A E K A S T L Q A L K E M Q S E H A	RVLR	DGK-MGTVCSRELVPGDVVE
Thalassiosira_pseudonana_3	EE-GILAYVEPLV IFL I L I I L N A M V G V W Q E S N A E A A L E A L K E L Q S E T A	RVLR	DGK-MATINSREIVPGDIIE
Pseudo-nitzschia_multiseries_2	DH-WVEAFVEPAV IFL I L I I L N A I VGWVQ E SNA E N A E A L E A L K K M Q S L H A	PCLR	DGVWHDALPTEELVPGDIVK
Fragilaropsis_cylindrus_9	GH-WIEAFVEPAV IFL I L I I L N A I VGWVQ E SNA E N A E A L E A L K K M Q S L H T	PCLR	GGIWYDQLPTEELVPGDIVK
Fragilaropsis_cylindrus_10	GH-WIEAFVEPAV IFL I L I I L N A I VGWVQ E SNA E N A E A L E A L K K M Q S L H T	PCLR	GGIWYDQLPTEELVPGDIVK
Aquilegia_caerulea_4	ET-GLTA FLEP S V I L M I L A A N A A V G V I T E T N A E K A E E L R A Y Q A D V A	TVLR	NGC-FSILPATDLVPGDIVE
Glycine_max_5	ET-GLMAFLEP S V I L M I L A A N A A V G V I T E T N A E K A E E L R A Y Q A D V A	TVLR	NGC-FSILPATELVPGDIVE
Medicago_truncatula_2	ET-GLTA FLEP S V I L M I L A A N A A V G V I T E T N A E K A E E L R A Y Q A N I A	TVLR	NGC-FSILPATELVPGDIVE
A.thaliana_AT1G10130.1	ET-GLAA FLEP S V I L I L A A N A A V G V I T E T N A E K A E E L R A Y Q A D V A	TVLR	NGC-FSILPATELVPGDIVE
Oryza_sativa_2	ET-GFTA FVE PFV IFL I L I A A N A T V G V V T E T N A E K A L K E L Q A D V A	TVLR	NGL-LSIVPASNLVPGDIVE
Selaginella_moellendorffii_4	ET-GFTA FVE PFV IFL I L I A A N A T V G V V T E T N A E K A L K E L Q A D V A	TVLR	SGR-LKVLPAAEELVPGDVVE
Micromonas_pusilla_2	D-GSGAFVEPGV IFL I L I L V N A T V G V L T E T N A E R A I E E L K A Y Q A N L A	TVLR	GGSLTVC PAAELVPGDIVE
Micromonas_sp_2	E-GIYSLIEPSVIACILIANAIVGVMTETNAAKA E E L G A Y Q A E V A	TVCR	GGSLTVC PAAELVPGDIVE
Volvox_carteri_2	EG-VLGALVEPV IFL I L I L I L A N A T V G V V T E R N A E Q A I E E L K A Y E A E S A	TVLR	SGV-LOLVP S GDLVPGDVVE
Chlamydomonas_reinhardtii_2	EG-VLGSLVEPV IFL I L I L I L A N A T V G V V T E R N A E Q A I E E L K A Y E A E S A	TVLR	NGV-LOLVPAGDLVPGDVVE
Dunaliella_salina_1	ES-IQSGLIEPMV ILL I L V N A T V G V V T E R N A E K A I E E L K S Y E	-	-
Ostreococcus_lucimarinus_2	EG-GSEAFLEP G V I A V I L I A N A A V G V A T E K N A E R A I E E L K K Y E A D V A	TCTR	DGE-KRKVN A E A L V P G D I V E
Phaeodactylum_tricornutum_2	-NOSEA I S I A F A L L I V S M V A A V O E Y R S E A A L E K L A T L V P H T C	TVLR	DGOVIDGFFAKELVVGDVL
Chlamydomonas_reinhardtii_3	-ALESWAEFG L I L G V I V I N T A L G Y E G R A E K A A D A I K A L L S P N A	TVLR	DGO-AAVLP A E S L V P G D V V L
Chlamydomonas_reinhardtii_4	-ALESWAEFG L I L G V I V I N T A L G L I Q E G R A E K A A D A I K A L L S P N A	TVLR	DGO-AAVLP A E S L V P G D V V L
Thalassiosira_pseudonana_4	NT-RFTAWFOQ A L I V S V I T L N T W I G I Y Q E G N A E K A A D A L K N M L S T D A	RVIR	GGK-EVMIPAGD I V P G D V C L
Thalassiosira_pseudonana_5	NT-FTAWFOQ A L I V S V I T L N T W I G I Y Q E G N A E K A A D A L K N M L S T D A	RVIR	SGK-EIMISAGE I V P G D V C L

Fragilariopsis_cylindrus_11	NP	TYTNWIQVAAIVGVIVANTVIGIVQEGSAENAAEALKNMLSSDA	VLVR	DGV	ETKVP	SQFIVPGD	VVV
Fragilariopsis_cylindrus_12	NP	TYTNWIQVAAIVGVIVANTVIGIVQEGSAENAAEALKNMLSSDA	VLVR	DGV	ETKVP	SQFIVPGD	VVV
Pseudo-nitzschia_multiseries_3	NP	RYTNFIQIGIILGVIVINTTIGLIOEGSAEEAAEALKNMLSSDA	IVIR	DGV	TTKVP	AQLLVPGDAVI	
Phaeodactylum_tricornutum_3	ED	RVTNFIEVGLTFVITLNNTIGIYQEGSAEKAEEALKNMLSSDA	ILIR	DGK	EVKIP	PASDVVPGD	VVV
Micromonas_pusilla_3	KD	QFLSYIDSIAITIIIVFINASIAAKAENNANDALEALSSLQAPI	TLIR	DGE	EIRVE	ESSDIVRGDLVK	
Micromonas_sp_3	KD	KFLSYVDSIAISIIILINATIAAVTEN SANDALEALSSLQSPMC	TVIR	GGE	EVS	IIESKNMVRGDIVK	
Chlamydomonas_reinhardtii_5	QR	AHQGWSEGLAVLIGTALIVIFLGAGQDFSKERQFQKLNAALKDVIDV	KVTR	GGK	OVLVP	NPTEVVVGDIMF	
Chlamydomonas_reinhardtii_6	QR	AHQGWSEGLAVLIGTALIVVFQAGQDYSKERQFQKLNAALKDNEIV	KVTR	GGK	OVLVP	NPTEIVVVGDVMF	
Volvox_carteri_3	QR	AHDGWSEGLAVLIGTALIVIFGAGQDYSKELQFQKLNLKDNDIV	KVTR	SGR	OVLIP	NPTDVVVGDILF	
Volvox_carteri_4	OR	AHDGWSEGLAVLIGTALIVIFGAGQDYSKELQFQKLNLKDNDIV	KVTR	SGR	OVLIP	NPTDVVVGDILF	
Volvox_carteri_5	ER	KKNHWIEGVAIWVAVIVVTLVGAGNDYSKDLQFRKLNAQKDRIQI	KVIR	GGE	OILVENT	DLVVGDIVI	
Chlamydomonas_reinhardtii_7	ER	AKSAWVEGVAIWVAVIVVTLVGAGNDYSKDLQFRKLNAQKDRIEI	KVVR	GGG	OILVP	NPTDLVVGDVML	
Chlamydomonas_reinhardtii_8	QR	KHGEWIEGVAIWVAAIIILVVSVGAGNDYQDKQFRKLNAQKDICKM	KVVR	GHO	TLLVEN	VELVVGDVYL	
Volvox_carteri_6	OR	KHGEWIEGVAIWVAAIIILVVSVGAGNDYQDKQFRKLNAQKDICKM	KVVR	GGH	TELIENT	OLVVGDVYL	
Ostreococcus_lucimarinus_3	HR	EEYGYLEGIAIVLVVFFVFLQAFIDYAKEMKFRQLNSVKDNQYQV	KVHR	DGK	SVAVPAGE	IMVGDVLVE	
Micromonas_sp_4	HR	EEYGYLEGIAIVLVVFFVFLQAFIDYAKEMKFRQLNSVKDNQYQV	KVVR	NGE	VHAVTA	GEVLVGDVVE	
Selaginella_moellendorffii_5	KV	KFASWYDGASIAFVILVVCVTACSDYKQSLQFQRLNAEKRKIHV	EVLR	GGR	RIGVS	IIFELVVGDVVP	
A.thaliana_AT3G21180.1	EG	LKEGWLDDGSIAFAVLLVIVVTAVDYRQSLQFQNLNDEKRNIQL	EIVR	GGR	TVKISIY	DIVVVGDVIP	
Aquilegia_caerulea_5	EG	IKEGWYDGGSIAFAVLLVIVVTAISDYRQSLQFQNLNDEKRNIHL	EIVR	GGK	RVE	ISIYDIVVVGDVIP	
A.thaliana_AT5G57110.2	EG	IKEGWYDGGSIAFAVILVIVVTAVDYRQSLQFQNLNDEKRNIHL	EIVR	GGR	RVE	ISIYDIVVVGDVIP	
Oryza_sativa_3	EG	IKEGWYDGASIAFAVLLVVVTTATSDYKQSLQFQNLNEEKQNIKL	EVVR	GGR	RISVIY	DLVAGDVVP	
Zea_mays_2	EG	IKEGWYDGASIAFAVLLVVVFTATSDYKQSLQFQNLNEEKQNIRL	EVVR	GGR	RTVSIY	DLVVGDVVP	
Zea_mays_3	EG	VKGDWYDGGSIFFAVFLVIFTATSDYRQSLQFQHLNEEKRNIQV	EVIR	GGK	RLVASI	FDLVVGDVVP	
Zea_mays_4	EG	VDEGWYDGGSIFFFLAVFLVILVTTATSDYRQSLQFQHLNEEKRNQIV	EVVR	GGK	RFGAS	IFDLVVGDVVP	
Selaginella_moellendorffii_6		VKEGWYDGASICICFAVLVVIFVTAFSDYRQSLQFQRLSLSOEKRNIQI	OVVR	GGK	RFTTS	IFDLVVGDIVP	
Selaginella_moellendorffii_7	EA	RWYDGGGTCFAIVCVMVASLSDYNQANQFQQLSAEKRKIYI	NVTR	GGH	RTKVS	IIFELVVGDVMH	
Aquilegia_caerulea_6	EG	WPKGGAHDGLGIAASILLVVFTATSDYRQSLQFQKDLLKEKKKIITI	QVTR	NGY	ROKISIY	DLLPGD	IVH
Medicago_truncatula_3	EG	WPKGGAHDGLGIVASILLVVFTATSDYRQSLQFQKDLLKEKKKISI	QVTR	NGY	ROKMSIY	YELLPGD	IVH
A.thaliana_AT2G22950.1	EG	WPQGSHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIITV	QVTR	NGF	ROKMSIY	DLLPGD	VVH
A.thaliana_AT4G37640.1	EG	WPKGSHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIITV	QVTR	NGF	ROKLSIY	DLLPGD	IVH
Oryza_sativa_4	EG	WPKGGAHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIITV	QVTR	NGY	ROKLSIY	DLLAGD	IVH
Zea_mays_5	EG	WPKGGAHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIITV	QVTR	NGY	ROKLSIY	DLLAGD	IVH
Oryza_sativa_5	EG	WPKGGAHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIITV	QVTR	NGF	RORLSIY	DLLPGD	VVH
A.thaliana_AT1G27770.1	EG	WPKGGAHDGLGIVASILLVVFTATSDYRQSLQFQFRDLDKEKKKIIVV	QVTR	DKL	ROKISIY	DLLPGD	VVH
Oryza_sativa_6	EG	WPOGAHDGVGIVASILLVSVTGTSNYQQSLQFRDLDKEKRKILV	QVTR	NGL	RQRVL	IDDDLPGD	AVH
Zea_mays_6	EG	WPSGAHDGIGIFTSILLVVFTATSNYQQSLQFRDLDREKRKISV	QVTR	DGF	RORIL	IIDDDLPGD	VVH
Aquilegia_caerulea_7	EG	WPKGMYDGLGILLSILLVVMVTAVSDYKOSLQFQKDLLKEKKKIMV	QVTR	NGY	ROKVS	IFDLVVGDIVH	
Oryza_sativa_7	KG	WPMGIYDGFGLLISILLVVLVTATSDYQQARKFMDLDREKQKIYI	RVTR	DKK	TKEVLV	HDLVVGDILH	
Selaginella_moellendorffii_8	EG	WKEGWYDGTLGALSILLVVFTAA SDYQOSLQFRDLDKEKKKILV	QVTR	NHK	RQKVS	IFDLVVGDVH	
Selaginella_moellendorffii_9	KG	FRDGWCAGILVSVVLVITVSASSDYQQAVQFRALDKEKGKVYI	QVTR	SAK	RRRILASE	LVVGDIVH	
Oryza_sativa_8	HG	IKDGYWDGVSIIFLAVFLVAASAVSNHSQGKRFDKLARESENIMV	SVVR	AAR	ROEVSI	FDVVVGDVVV	
Fragilariopsis_cylindrus_13	DD	PAAGYVEGMAILSACLIVSVVTALNDYQKETQFREL	LVIR	SGK	FQKIP	PVGEVVGDLS	
Pseudo-nitzschia_multiseries_4	DD	PEVGYVEGMAILAACLVVSVVTALNDYQKETQFREL	LVIR	SGK	FQKIP	PVGDLVVGDLS	
Phaeodactylum_tricornutum_4	DD	PATGYVEGCAILAAVLVVSFTA VNVDFQKESQFREL	LVVR	NNV	HWQIP	PVDELVVGDVVC	
Thalassiosira_pseudonana_6	EK	CGMCWLEGAAAILASVCIVVLFVTAGIDYAKQFAFIRL	TRSLNDTNTKMVIR	NGH	OMSVT	DDEIVVGDI	LSS
Phaeodactylum_tricornutum_5	EE	CTTCWIEGAAILSVCIVVVFVSAGIDYAKQFAFLRLTRSLHET	TNTKQVIR	EGK	QVSII	DDDLVVGDILS	
Phaeodactylum_tricornutum_6	ED	CKTCWIEGAAILSVCIVVLFVTAISDYAKQFAFLRLTRSLHDTNTKQVIR	DAK	QVSIV	DDDDIVVGDI	LSS	

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Skeletonema_marinoi_ABC
 Thalassiosira_pseudonana_1
 Thalassiosira_oceanica_1
 Phaeodactylum_tricornutum_1
 Fragilariaopsis_cylindrus_1
 Fragilariaopsis_cylindrus_2
 Fragilariaopsis_cylindrus_3
 Fragilariaopsis_cylindrus_4
 Fragilariaopsis_cylindrus_5
 Fragilariaopsis_cylindrus_6
 Fragilariaopsis_cylindrus_7
 Fragilariaopsis_cylindrus_8
 Pseudo-nitzschia_multiseries_1
 Thalassiosira_pseudonana_2
 Ostreococcus_lucimarinus_1
 Zea_mays_1
 Oryza_sativa_1
 Aquilegia_caerulea_1
 A.thaliana_AT1G07810.1
 A.thaliana_AT1G07670.1
 Medicago_truncatula_1
 Glycine_max_1
 Glycine_max_2
 Aquilegia_caerulea_2
 Selaginella_moellendorffii_1
 Physcomitella_patens_1
 Physcomitella_patens_2
 Physcomitella_patens_3
 Selaginella_moellendorffii_2
 Glycine_max_3
 Glycine_max_4
 Aquilegia_caerulea_3
 A.thaliana_AT4G00900.1
 Chlamydomonas_reinhardtii_1
 Volvox_carteri_1
 Micromonas_pusilla_1
 Micromonas_sp_1
 Selaginella_moellendorffii_3
 Thalassiosira_oceanica_2
 Thalassiosira_pseudonana_3
 Pseudo-nitzschia_multiseries_2
 Fragilariaopsis_cylindrus_9
 Fragilariaopsis_cylindrus_10
 Aquilegia_caerulea_4
 Glycine_max_5
 Medicago_truncatula_2
 A.thaliana_AT1G10130.1
 Oryza_sativa_2
 Selaginella_moellendorffii_4
 Micromonas_pusilla_2
 Micromonas_sp_2
 Volvox_carteri_2
 Chlamydomonas_reinhardtii_2
 Dunaliella_salina_1
 Ostreococcus_lucimarinus_2
 Phaeodactylum_tricornutum_2
 Chlamydomonas_reinhardtii_3
 Chlamydomonas_reinhardtii_4
 Thalassiosira_pseudonana_4
 Thalassiosira_pseudonana_5

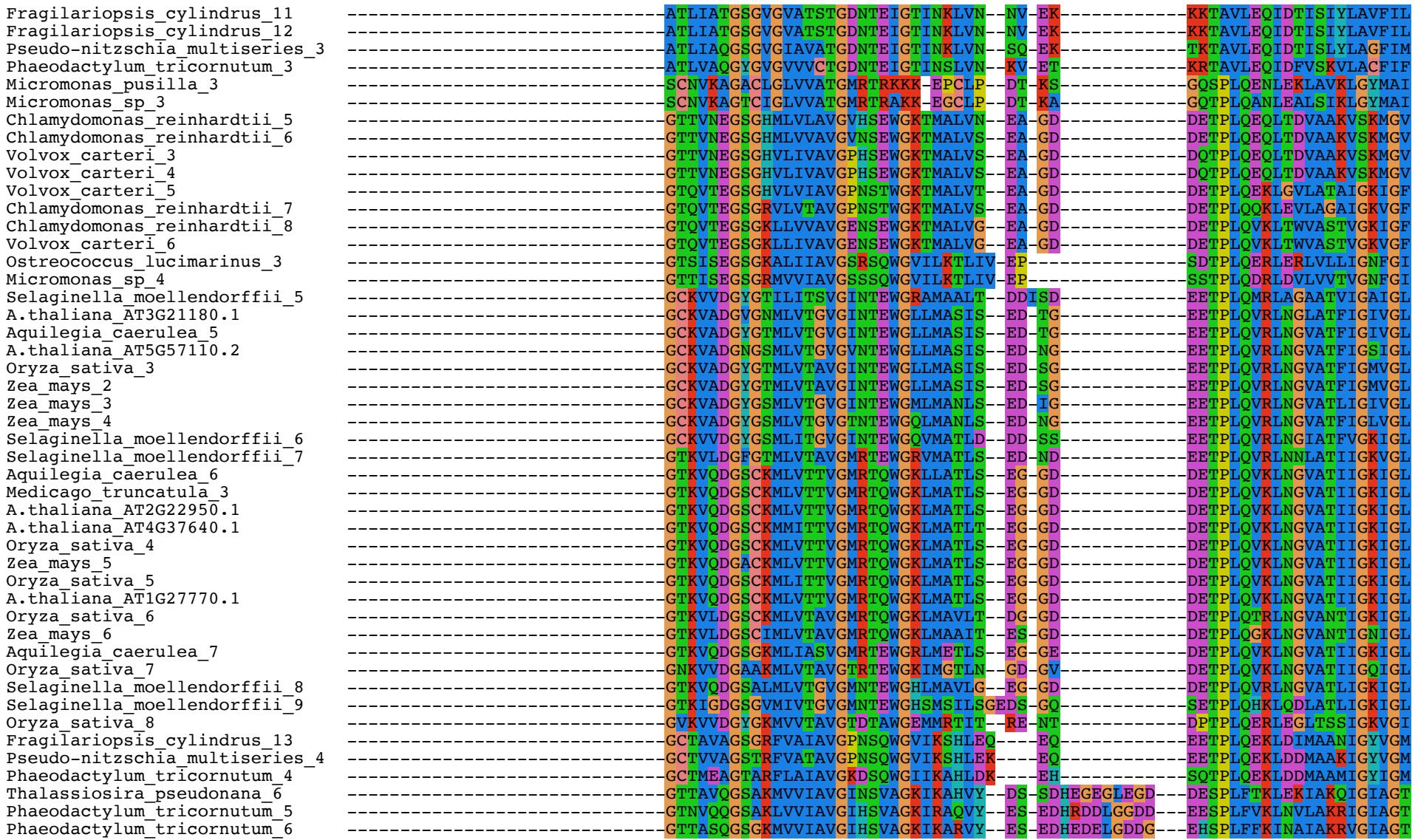
<img alt="Sequence alignment of protein fragments from various organisms. The alignment shows a series of colored boxes representing amino acid sequences. The first column contains the sequence 'LSVGO' for most organisms. The second column contains 'KIPADIRL' for most, except for some which have variations like 'YIKRRT' or 'YIKRR'. The third column contains 'SS' for most, except for some which have variations like 'QS' or 'TA'. The fourth column contains 'FTVDEACILTGESD' for most, except for some which have variations like 'FTVDEACILTGESD' or 'FTVDEACILTGESD'. The fifth column contains 'VNP' for most, except for some which have variations like 'VNP' or 'VNP'. The sixth column contains 'GDIKNDMQ' for most, except for some which have variations like 'GDIKNDMQ' or 'GDIKNDMQ'. The seventh column contains 'NG' for most, except for some which have variations like 'NG' or 'NG'. The eighth column contains 'GGTMGANSNGMLYG' for most, except for some which have variations like 'GGTMGANSNGMLYG' or 'GGTMGANSNGMLYG'. The ninth column contains 'NGHHANGMLYG' for most, except for some which have variations like 'NGHHANGMLYG' or 'NGHHANGMLYG'. The tenth column contains 'SGS' for most, except for some which have variations like 'SGS' or 'SGS'. The eleventh column contains 'GSPNRPVQDOKGMLYS' for most, except for some which have variations like 'GSPNRPVQDOKGMLYS' or 'GSPNRPVQDOKGMLYS'. The twelfth column contains 'NSISNHQFNDGMCYS' for most, except for some which have variations like 'NSISNHQFNDGMCYS' or 'NSISNHQFNDGMCYS'. The thirteenth column contains 'NSISNHQFNDGMCYS' for most, except for some which have variations like 'NSISNHQFNDGMCYS' or 'NSISNHQFNDGMCYS'. The fourteenth column contains 'NSISNHQFNDGMCYS' for most, except for some which have variations like 'NSISNHQFNDGMCYS' or 'NSISNHQFNDGMCYS'. The fifteenth column contains 'NSISNHQFNDGMCYS' for most, except for some which have variations like 'NSISNHQFNDGMCYS' or 'NSISNHQFNDGMCYS'. The sixteenth column contains 'NSISNHQFNDGMCYS' for most, except for some which have variations like 'NSISNHQFNDGMCYS' or 'NSISNHQFNDGMCYS'. The seventeenth column contains 'PSKAPVOSQSGMVFS' for most, except for some which have variations like 'PSKAPVOSQSGMVFS' or 'PSKAPVOSQSGMVFS'. The eighteenth column contains 'DDKKTIPIQDQSSMLFS' for most, except for some which have variations like 'DDKKTIPIQDQSSMLFS' or 'DDKKTIPIQDQSSMLFS'. The nineteenth column contains 'DEDIELOGKTCMLFA' for most, except for some which have variations like 'DEDIELOGKTCMLFA' or 'DEDIELOGKTCMLFA'. The twentieth column contains 'LEDTDILOGKECMVFA' for most, except for some which have variations like 'LEDTDILOGKECMVFA' or 'LEDTDILOGKECMVFA'. The twenty-first column contains 'V-DTDILOGKRCMLFA' for most, except for some which have variations like 'V-DTDILOGKRCMLFA' or 'V-DTDILOGKRCMLFA'. The twenty-second column contains 'E-NADIOGKKCMVFA' for most, except for some which have variations like 'E-NADIOGKKCMVFA' or 'E-NADIOGKKCMVFA'. The twenty-third column contains 'E-NADIOGKKCMVFA' for most, except for some which have variations like 'E-NADIOGKKCMVFA' or 'E-NADIOGKKCMVFA'. The twenty-fourth column contains 'E-DADIOQGKRCMVFA' for most, except for some which have variations like 'E-DADIOQGKRCMVFA' or 'E-DADIOQGKRCMVFA'. The twenty-fifth column contains 'E-DADIOQGKRCMVFA' for most, except for some which have variations like 'E-DADIOQGKRCMVFA' or 'E-DADIOQGKRCMVFA'. The twenty-sixth column contains 'V-ETDILOGKCMVFA' for most, except for some which have variations like 'V-ETDILOGKCMVFA' or 'V-ETDILOGKCMVFA'. 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The thirty-third column contains 'MDDCELOGKENMVFA' for most, except for some which have variations like 'MDDCELOGKENMVFA' or 'MDDCELOGKENMVFA'. The thirty-fourth column contains 'DPNCELOSKECMLFA' for most, except for some which have variations like 'DPNCELOSKECMLFA' or 'DPNCELOSKECMLFA'. The thirty-fifth column contains 'DPNCELOQAKECMVFA' for most, except for some which have variations like 'DPNCELOQAKECMVFA' or 'DPNCELOQAKECMVFA'. The thirty-sixth column contains 'DPDAAELQAKGCMLF' for most, except for some which have variations like 'DPDAAELQAKGCMLF' or 'DPDAAELQAKGCMLF'. The thirty-seventh column contains 'DAGCELOGKECVLFG' for most, except for some which have variations like 'DAGCELOGKECVLFG' or 'DAGCELOGKECVLFG'. The thirty-eighth column contains 'DEVEILOGKDNMVFA' for most, except for some which have variations like 'DEVEILOGKDNMVFA' or 'DEVEILOGKDNMVFA'. The thirty-ninth column contains 'DDELVVQAKTNIMFA' for most, except for some which have variations like 'DDELVVQAKTNIMFA' or 'DDELVVQAKTNIMFA'. The四十th column contains 'EDELVVQAKTNIMFA' for most, except for some which have variations like 'EDELVVQAKTNIMFA' or 'EDELVVQAKTNIMFA'. The四十-one column contains 'S-GEKISDQKNMVFA' for most, except for some which have variations like 'S-GEKISDQKNMVFA' or 'S-GEKISDQKNMVFA'. The四十-two column contains 'SSNKSSISDQKNMVFA' for most, except for some which have variations like 'SSNKSSISDQKNMVFA' or 'SSNKSSISDQKNMVFA'. The四十-three column contains 'ATNAVFDKTNILFS' for most, except for some which have variations like 'ATNAVFDKTNILFS' or 'ATNAVFDKTNILFS'. The四十-four column contains 'TTNAVYQDKTNILFS' for most, except for some which have variations like 'TTNAVYQDKTNILFS' or 'TTNAVYQDKTNILFS'. The四十-five column contains 'AANAVYQDKTNILFS' for most, except for some which have variations like 'AANAVYQDKTNILFS' or 'AANAVYQDKTNILFS'. The四十-six column contains 'TTNAVYQDKKNILFS' for most, except for some which have variations like 'TTNAVYQDKKNILFS' or 'TTNAVYQDKKNILFS'. The四十-seven column contains 'TMNAVYQDKTNILFSNITNRDN' for most, except for some which have variations like 'TMNAVYQDKTNILFSNITNRDN' or 'TMNAVYQDKTNILFSNITNRDN'. The四十-eight column contains 'VINPVPYQDKTSILFS' for most, except for some which have variations like 'VINPVPYQDKTSILFS' or 'VINPVPYQDKTSILFS'. The四十-nine column contains 'SAKAVVQDKTCMLYS' for most, except for some which have variations like 'SAKAVVQDKTCMLYS' or 'SAKAVVQDKTCMLYS'. The五十柱 contains 'ATKAVLODKTCIAFS' for most, except for some which have variations like 'ATKAVLODKTCIAFS' or 'ATKAVLODKTCIAFS'. The fifty-one column contains 'KDNPVYQDKTNMLFS' for most, except for some which have variations like 'KDNPVYQDKTNMLFS' or 'KDNPVYQDKTNMLFS'. The fifty-two column contains 'KDNPVYQDKTNMLFS' for most, except for some which have variations like 'KDNPVYQDKTNMLFS' or 'KDNPVYQDKTNMLFS'. The fifty-three column contains 'NKOAVYQDKLNMLFS' for most, except for some which have variations like 'NKOAVYQDKLNMLFS' or 'NKOAVYQDKLNMLFS'. The fifty-four column contains 'HLGECLVQDKTCMVYS' for most, except for some which have variations like 'HLGECLVQDKTCMVYS' or 'HLGECLVQDKTCMVYS'. The fifty-five column contains 'GASPLTQQKNNVFA' for most, except for some which have variations like 'GASPLTQQKNNVFA' or 'GASPLTQQKNNVFA'. The fifty-six column contains 'APAQAGLGDRCMCFS' for most, except for some which have variations like 'APAQAGLGDRCMCFS' or 'APAQAGLGDRCMCFS'. The fifty-seven column contains 'APAQAGLGDRCMCFS' for most, except for some which have variations like 'APAQAGLGDRCMCFS' or 'APAQAGLGDRCMCFS'. The fifty-eight column contains 'ANPDQVPLGDRKNMAYS' for most, except for some which have variations like 'ANPDQVPLGDRKNMAYS' or 'ANPDQVPLGDRKNMAYS'. The fifty-nine column contains 'MDPDQVPLGDRKNMYS' for most, except for some which have variations like 'MDPDQVPLGDRKNMYS' or 'MDPDQVPLGDRKNMYS'.
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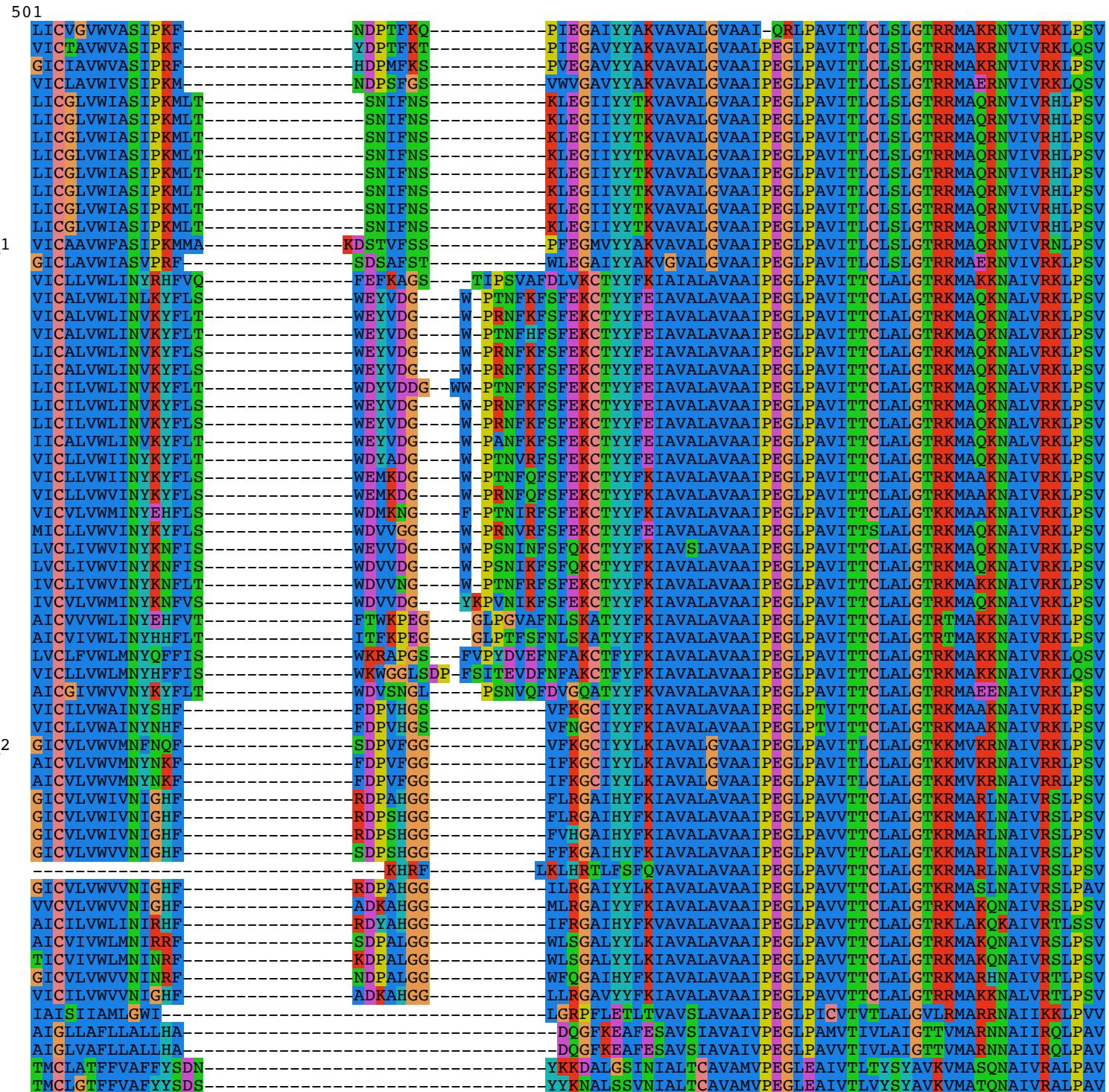
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Fragilariopsis_cylindrus_12	LALGD--KIPCDLDRV	-EV-SN-MASAEAALTGESVPIEK	TPLP-TIELNEG	QLPKQVPLGDRKNMCF
Pseudo-nitzschia_multiseries_3	LGLGD--RIPADLDRV	-EV-SN-MASAEAALTGESVPIEK	TPQA-TFLNEG	QLAKQVPLGDRKNMCF
Phaeodactylum_tricornutum_3	LGTGD--RVPGDIRML	-EV-NN-LACQEAALTGESVPIEK	VTDA-IDCGSG	KPEQVPLGDRKNMCF
Micromonas_pusilla_3	LGTGD--VVPADVRCI	-TA-ND-LRVNEMLLTGESEDVAK	STKVKP	RVPGHPEKLTADNMAFS
Micromonas_sp_3	LGTGD--VVPADCRCI	-KA-ND-FRVNEMLLTGESEDVAK	NTKIKA	RVPGQPEKLTADNMAFS
Chlamydomonas_reinhardtii_5	LDTGD--KVIADGIVI	-DS-QG-LVLDEASLTGESDPIKK	D	PVSDPWRS
Chlamydomonas_reinhardtii_6	LDTGD--KVIADGVVI	-DS-QG-IVLDEASLTGESDPIKK	D	AVSDPWIRS
Volvox_carteri_3	LDTGD--KVIADGIVI	-DS-QG-LVLDEASLTGESDPIKK	D	PLNDPWVRS
Volvox_carteri_4	LDTGD--KVIADGIVI	-DS-QG-LVLDEASLTGESDPIKK	D	PLNDPWVRS
Volvox_carteri_5	LDTGD--KVVADAIWI	-DS-QG-LTMDEASLTGESDPMKK	N	INEDPWVMS
Chlamydomonas_reinhardtii_7	LDTGD--KVVADAIWI	-DS-QG-LTMDEASLTGESDPMKK	N	TTADPWVMS
Chlamydomonas_reinhardtii_8	LDTGD--KVVADGVCF	-DC-QG-LVIDEASLTGESDPIKK	N	TDEDPWVRS
Volvox_carteri_6	LDTGD--KVVADGICF	-DS-QG-LVVDEASLTGESDPIKK	N	PEDDCWVRS
Ostreococcus_lucimarinus_3	LAAGD-KVPADALFV	-EG-SK-FKANEAAAMTGEPIDISK	S	REKDPWVLS
Micromonas_sp_4	L\$AGD-KVPADGVFL	-EG-SK-LRADESAMTGEPIGIASK	S	HDKDPFLMS
Selaginella_moellendorffii_5	LKTGD-QIPADGVLV	-DG-YS-LVVDESSILTGESDPVSM	PK	GLDHPFFMS
A.thaliana_AT3G21180.1	LRIGD-QVPADGVLI	-SG-HS-LAIDESSMTGESKIVHK	D	QKSPFLMS
Aquilegia_caerulea_5	LKIGD-QVPADGVLV	-VG-HS-LAIDESSMTGEAKIVHK	D	HRKDPFLMS
A.thaliana_AT5G57110.2	LNIGN-QVPADGVLI	-SG-HS-LALDESSMTGESKIVNK	D	ANKDPFLMS
Oryza_sativa_3	LKIGD-QVPADGILI	-SG-HS-LSVDESSMTGESKIVHK	D	OKSPFLMS
Zea_mays_2	LKIGD-QVPADGILI	-SG-HS-LSIDEDESSMTGESKIVHK	D	OKSPFLMS
Zea_mays_3	LKIGD-QVPADGILI	-YG-HS-LAIDESSSMTGESKIVNK	D	ORAPFLMS
Zea_mays_4	LKIGD-QVPADGVLI	-SG-HS-LAIDESSSMTGESKVVHK	D	OKAPFLMS
Selaginella_moellendorffii_6	LNIGD-QVPADGVLV	-SG-HS-LSIDEDESSMTGESEPVHV	D	GKSPFLHS
Selaginella_moellendorffii_7	LAIGD-QIPADGLVY	-VG-HS-LIVDESSMTGESDPLPK	D	EEEKPFLMS
Aquilegia_caerulea_6	LAIGD-QVPADGLFV	-SG-FS-LSINESSLTGESDPVMV	N	AQNPYMLS
Medicago_truncatula_3	LAIGD-QVPADGLFV	-SG-FS-LLIDEDESSLTGESEPVVV	N	TENPFLLS
A.thaliana_AT2G22950.1	LAIGD-QVPADGLFL	-SG-FS-VVIDEDESSLTGESEPVMV	T	AQNPFLS
A.thaliana_AT4G37640.1	LAIGD-QVPADGLFL	-SG-FS-VVIDEDESSLTGESEPVMV	N	AQNPFLMS
Oryza_sativa_4	LSIGD-QVPADGLFL	-SG-FS-LLINEDESSLTGESEPVAV	N	AENPFLLS
Zea_mays_5	LSIGD-QVPADGLFV	-SG-FS-MLIDEDESSLTGESEPVAV	S	AENPFLLS
Oryza_sativa_5	LAIGD-QVPADGLFI	-SG-FS-LLINEDESSLTGESEPVVV	N	EDNPFLLS
A.thaliana_AT1G27770.1	LGIGD-QIPADGLFI	-SG-FS-VLINEDESSLTGESEPVS	S	VEHPFLLS
Oryza_sativa_6	LAVGD-QVPADGLFI	-SG-FS-VLVDEDESSLTGESEPVFV	N	EDNPYLLS
Zea_mays_6	LGVD-QVPADGLFV	-SG-YS-VLVNESSSLTGESEPVV	S	EDNPFLLS
Aquilegia_caerulea_7	LSIGD-QVPADGFFI	-SG-YS-LVIDEDESSLSGESEPVN	N	QRNPFLLA
Oryza_sativa_7	LSIGD-QVPADGLFI	-SG-DC-LMIDEDESSLSGESEPVN	S	EERPFLHA
Selaginella_moellendorffii_8	LSIGD-QVPADGLFI	-SG-YS-LVIDEDESSMTGESEPOHV	G	KNKPFLLS
Selaginella_moellendorffii_9	LGIID-QIPADGLLL	-YG-QS-LLVDESCMTGESEMRAK	S	AEQPFLLS
Oryza_sativa_8	LKIGD-VVPADGVFL	-DG-HA-LQVDEDESSMTGEPHPVEV	D	AVKSPFLAS
Fragilariopsis_cylindrus_13	LEAGD-SIPCDGVLV	-NY-DG-LEVDESALITGEPEDIDK	D	FENDPFLLS
Pseudo-nitzschia_multiseries_4	LEAGD-SIPCDGVVV	-OY-DG-LTVDESALITGEPEDIEK	D	FENDPFLLS
Phaeodactylum_tricornutum_4	VEAGD-QIPCDGVLL	-VA-DD-VQVDESALITGEPTDV	S	LONGPFVLS
Thalassiosira_pseudonana_6	INAHLNLASIPADCVVL	-GPS-GG-LKMDDESSLTGESVLIAK		NPGDDVVL
Phaeodactylum_tricornutum_5	INAHLNLASIPADCVLL	-GPA-TD-LKMDDESTLTGESKAVSK		KPGDDVVL
Phaeodactylum_tricornutum_6	VNAHLNLASIPADCVLL	-GPA-GD-LKMDDESTLTGESKAVSK		KPGDIILS

401

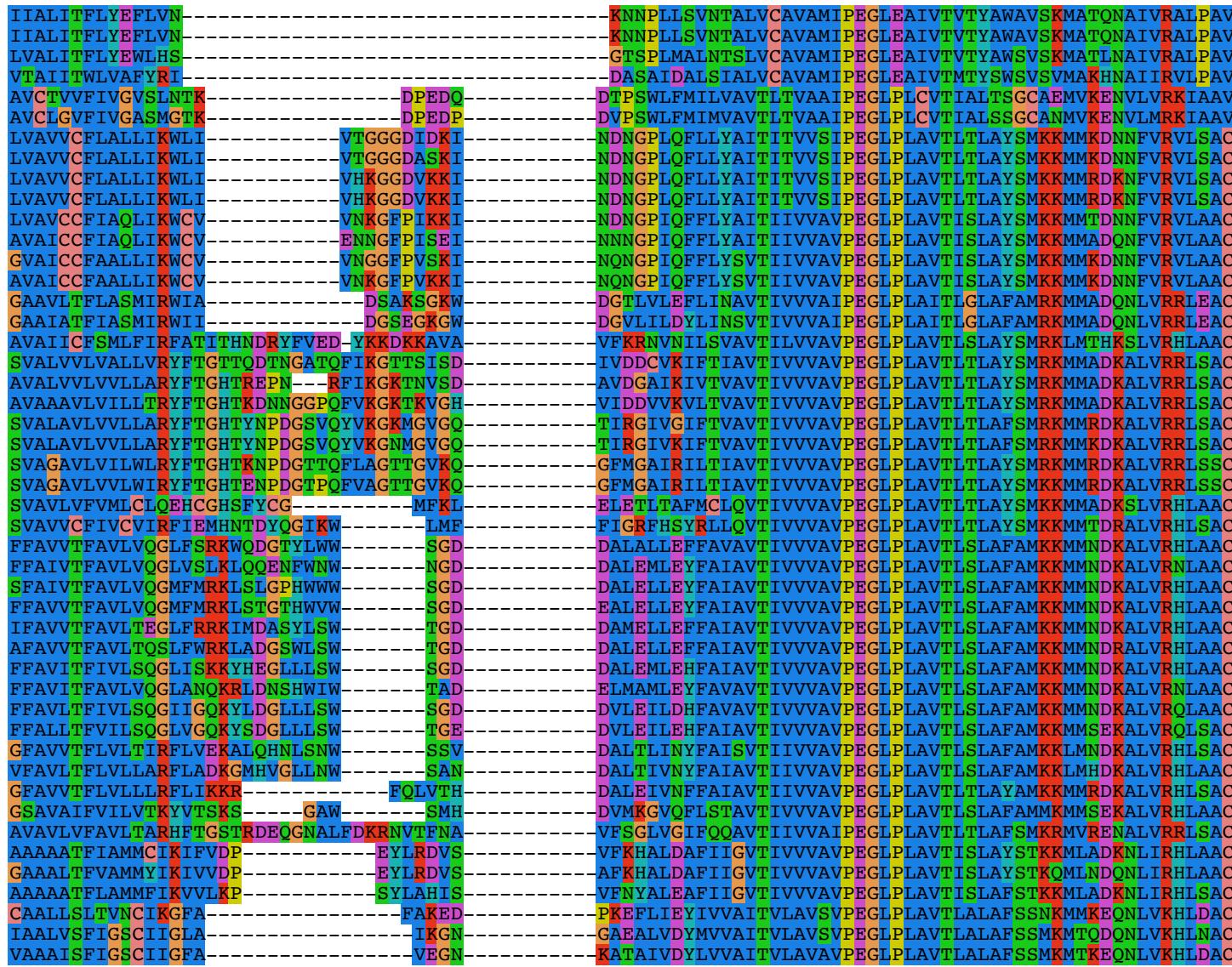
Skeletonema_marinoi_ABC
 Thalassiosira_pseudonana_1
 Thalassiosira_oceanica_1
 Phaeodactylum_tricornutum_1
 Fragilaropsis_cylindrus_1
 Fragilaropsis_cylindrus_2
 Fragilaropsis_cylindrus_3
 Fragilaropsis_cylindrus_4
 Fragilaropsis_cylindrus_5
 Fragilaropsis_cylindrus_6
 Fragilaropsis_cylindrus_7
 Fragilaropsis_cylindrus_8
 Pseudo-nitzschia_multiseries_1
 Thalassiosira_pseudonana_2
 Ostreococcus_lucimarinus_1
 Zea_mays_1
 Oryza_sativa_1
 Aquilegia_caerulea_1
 A.thaliana_AT1G07810.1
 A.thaliana_AT1G07670.1
 Medicago_truncatula_1
 Glycine_max_1
 Glycine_max_2
 Aquilegia_caerulea_2
 Selaginella_moellendorffii_1
 Physcomitella_patens_1
 Physcomitella_patens_2
 Physcomitella_patens_3
 Selaginella_moellendorffii_2
 Glycine_max_3
 Glycine_max_4
 Aquilegia_caerulea_3
 A.thaliana_AT4G00900.1
 Chlamydomonas_reinhardtii_1
 Volvox_carteri_1
 Micromonas_pusilla_1
 Micromonas_sp_1
 Selaginella_moellendorffii_3
 Thalassiosira_oceanica_2
 Thalassiosira_pseudonana_3
 Pseudo-nitzschia_multiseries_2
 Fragilaropsis_cylindrus_9
 Fragilaropsis_cylindrus_10
 Aquilegia_caerulea_4
 Glycine_max_5
 Medicago_truncatula_2
 A.thaliana_AT1G10130.1
 Oryza_sativa_2
 Selaginella_moellendorffii_4
 Micromonas_pusilla_2
 Micromonas_sp_2
 Volvox_carteri_2
 Chlamydomonas_reinhardtii_2
 Dunaliella_salina_1
 Ostreococcus_lucimarinus_2
 Phaeodactylum_tricornutum_2
 Chlamydomonas_reinhardtii_3
 Chlamydomonas_reinhardtii_4
 Thalassiosira_pseudonana_4
 Thalassiosira_pseudonana_5

The alignment shows a highly conserved sequence at the top followed by a more variable region. The first 200 positions are mostly 'G' (green) or 'A' (green). The second 200 positions show a mix of colors, with 'D' (orange), 'E' (red), 'F' (yellow), 'H' (purple), 'I' (pink), 'K' (dark blue), 'L' (light blue), 'M' (teal), 'N' (light purple), 'P' (brown), 'Q' (light orange), 'R' (dark red), 'S' (light pink), 'T' (light teal), 'V' (light yellow), 'W' (light brown), and 'Y' (light red) appearing frequently.





Fragilariopsis_cylindrus_11
 Fragilariopsis_cylindrus_12
 Pseudo-nitzschia_multiseries_3
 Phaeodactylum_tricornutum_3
 Micromonas_pusilla_3
 Micromonas_sp_3
 Chlamydomonas_reinhardtii_5
 Chlamydomonas_reinhardtii_6
 Volvox_carteri_3
 Volvox_carteri_4
 Volvox_carteri_5
 Chlamydomonas_reinhardtii_7
 Chlamydomonas_reinhardtii_8
 Volvox_carteri_6
 Ostreococcus_lucimarinus_3
 Micromonas_sp_4
 Selaginella_moellendorffii_5
 A.thaliana_AT3G21180.1
 Aquilegia_caerulea_5
 A.thaliana_AT5G57110.2
 Oryza_sativa_3
 Zea_mays_2
 Zea_mays_3
 Zea_mays_4
 Selaginella_moellendorffii_6
 Selaginella_moellendorffii_7
 Aquilegia_caerulea_6
 Medicago_truncatula_3
 A.thaliana_AT2G22950.1
 A.thaliana_AT4G37640.1
 Oryza_sativa_4
 Zea_mays_5
 Oryza_sativa_5
 A.thaliana_AT1G27770.1
 Oryza_sativa_6
 Zea_mays_6
 Aquilegia_caerulea_7
 Oryza_sativa_7
 Selaginella_moellendorffii_8
 Selaginella_moellendorffii_9
 Oryza_sativa_8
 Fragilariopsis_cylindrus_13
 Pseudo-nitzschia_multiseries_4
 Phaeodactylum_tricornutum_4
 Thalassiosira_pseudonana_6
 Phaeodactylum_tricornutum_5
 Phaeodactylum_tricornutum_6

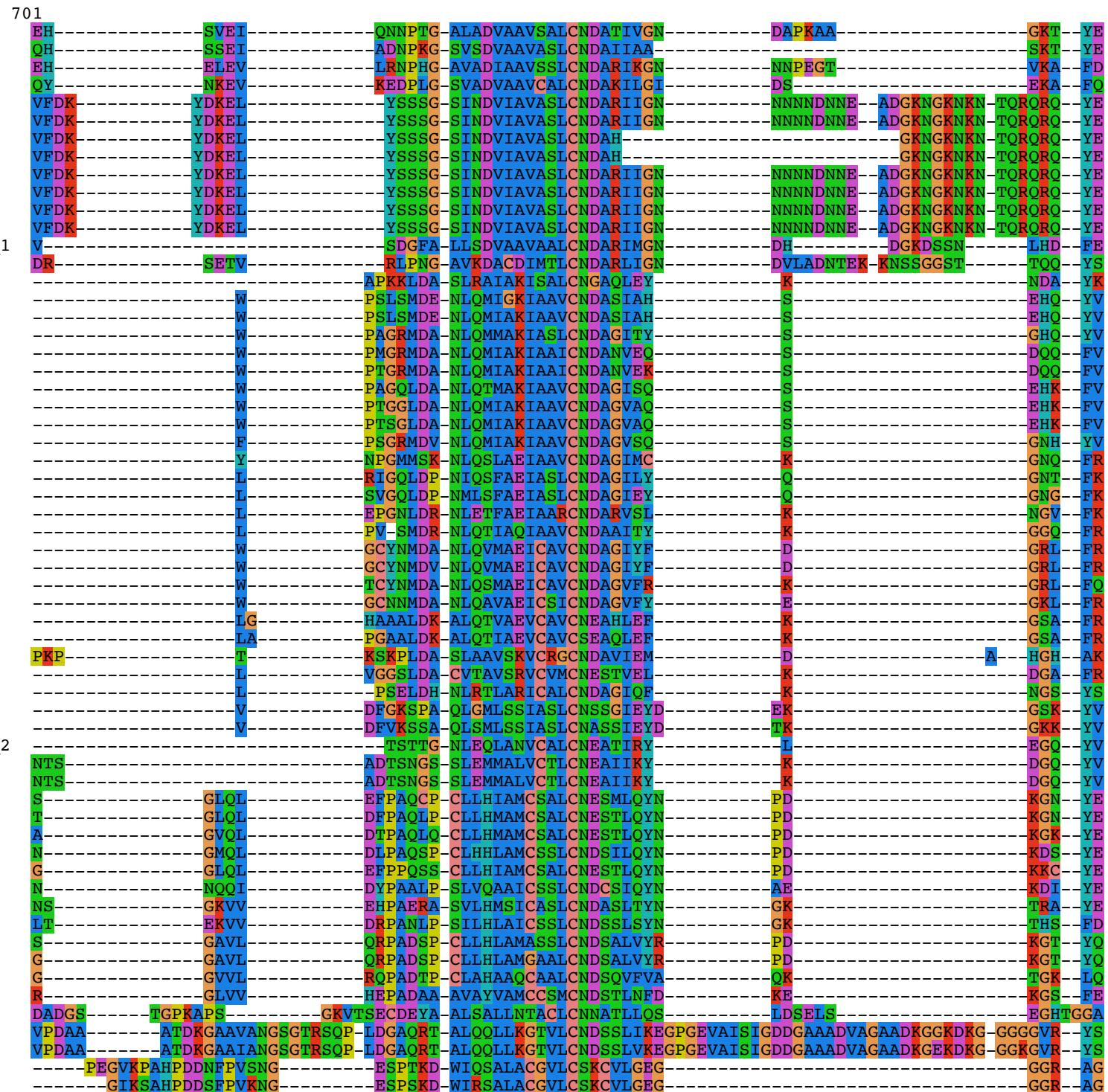


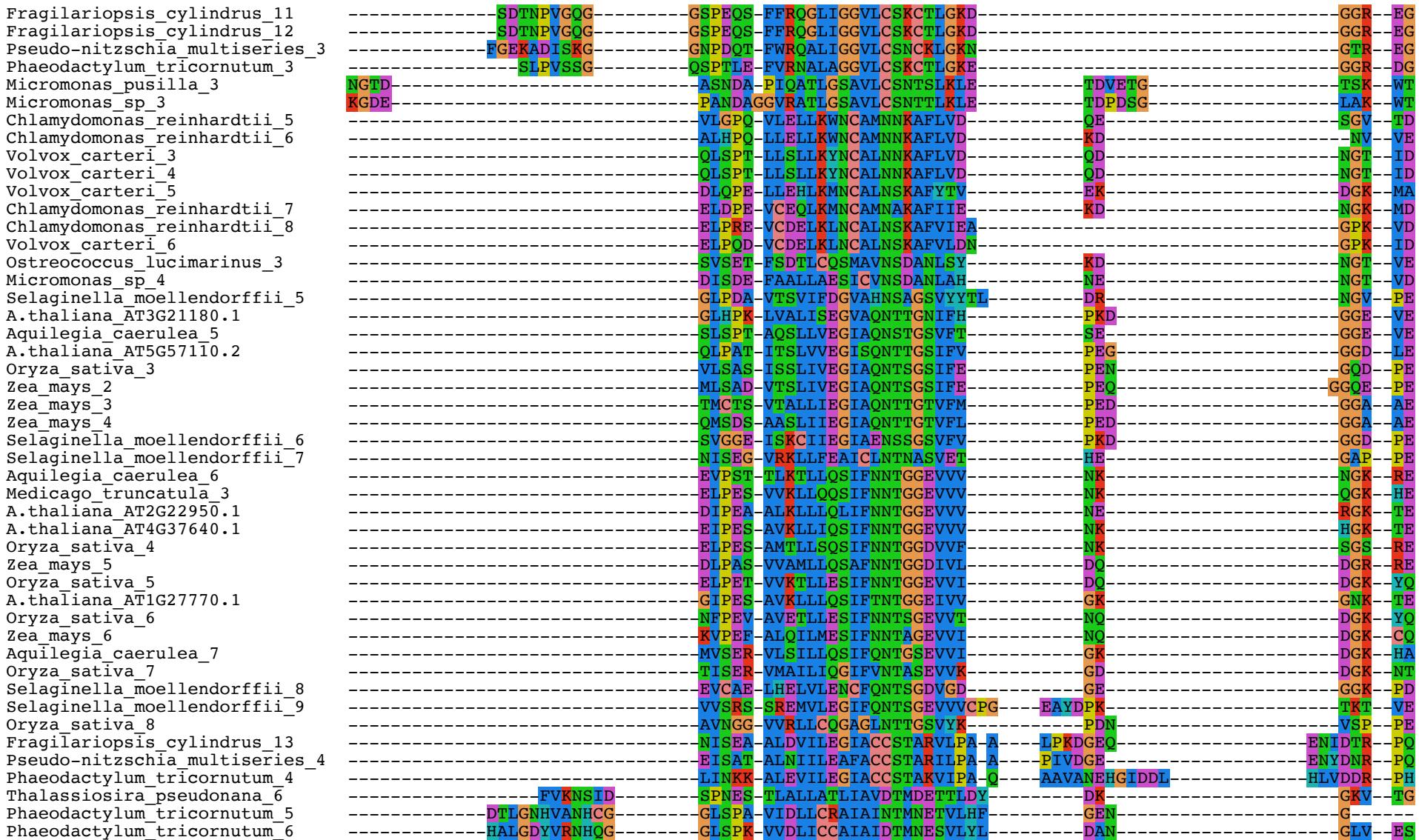
601

Skeletonema_marinoi_ABC
 Thalassiosira_pseudonana_1
 Thalassiosira_oceanica_1
 Phaeodactylum_tricornutum_1
 Fragilariaopsis_cylindrus_1
 Fragilariaopsis_cylindrus_2
 Fragilariaopsis_cylindrus_3
 Fragilariaopsis_cylindrus_4
 Fragilariaopsis_cylindrus_5
 Fragilariaopsis_cylindrus_6
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 Ostreococcus_lucimarinus_1
 Zea_mays_1
 Oryza_sativa_1
 Aquilegia_caerulea_1
 A.thaliana_AT1G07810.1
 A.thaliana_AT1G07670.1
 Medicago_truncatula_1
 Glycine_max_1
 Glycine_max_2
 Aquilegia_caerulea_2
 Selaginella_moellendorffii_1
 Physcomitella_patens_1
 Physcomitella_patens_2
 Physcomitella_patens_3
 Selaginella_moellendorffii_2
 Glycine_max_3
 Glycine_max_4
 Aquilegia_caerulea_3
 A.thaliana_AT4G00900.1
 Chlamydomonas_reinhardtii_1
 Volvox_carteri_1
 Micromonas_pusilla_1
 Micromonas_sp_1
 Selaginella_moellendorffii_3
 Thalassiosira_oceanica_2
 Thalassiosira_pseudonana_3
 Pseudo-nitzschia_multiseries_2
 Fragilariaopsis_cylindrus_9
 Fragilariaopsis_cylindrus_10
 Aquilegia_caerulea_4
 Glycine_max_5
 Medicago_truncatula_2
 A.thaliana_AT1G10130.1
 Oryza_sativa_2
 Selaginella_moellendorffii_4
 Micromonas_pusilla_2
 Micromonas_sp_2
 Volvox_carteri_2
 Chlamydomonas_reinhardtii_2
 Dunaliella_salina_1
 Ostreococcus_lucimarinus_2
 Phaeodactylum_tricornutum_2
 Chlamydomonas_reinhardtii_3
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 Thalassiosira_pseudonana_4
 Thalassiosira_pseudonana_5

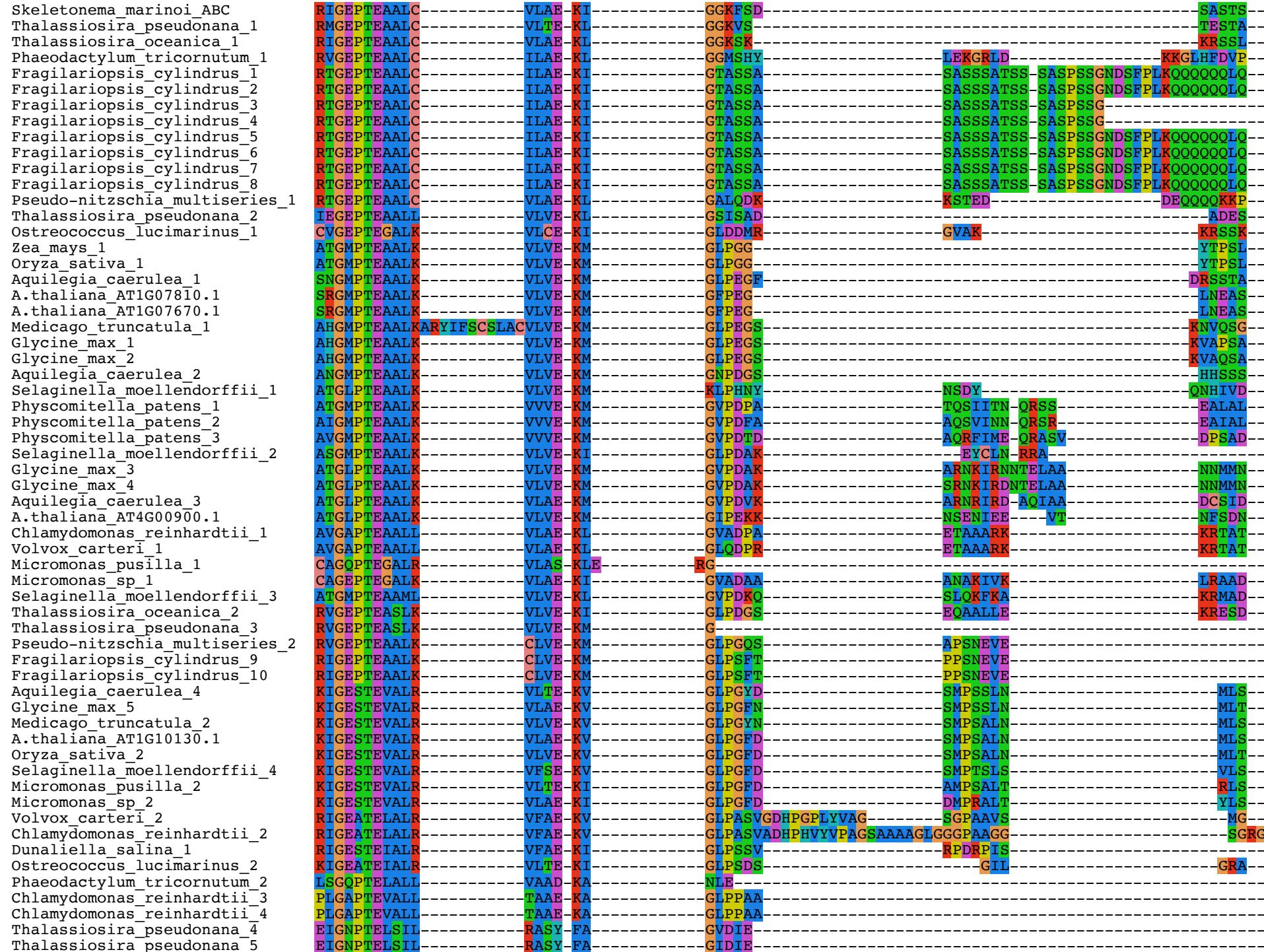
The alignment shows the conservation of a specific sequence motif across various eukaryotic species. The sequences are color-coded by residue identity, with each column having a unique color scheme. A vertical dashed line marks position 601. The alignment is color-coded by residue identity, with each column having a unique color scheme.

Fragilariopsis_cylindrus_11	ETLGSVTVICSDKTGTLTTNVMSLTAFVTS	NAHYKNNVVSS	VMAVDT	
Fragilariopsis_cylindrus_12	ETLGSVTVICSDKTGTLTTNVMSLTAFVTS	NAHYKNNVVSS	VMAVDT	
Pseudo-nitzschia_multiseries_3	ETLGSVTVICSDKTGTLTTNVMSMTAFVTS	NAHYKNNVHAS	DRTPA NLVRDDTF	
Phaeodactylum_tricornutum_3	ETLGSVTVICSDKTGTLTKNEMSLIGFVTS	NARYKIDVDISK	NRSAD NFVRDDT	
Micromonas_pusilla_3	ETLGSASIICTDKTGTLTEGKMLVAMHAG	GVDYTVTGKGFD	PT	VGKITTS
Micromonas_sp_3	ETLGSASIICTDKTGTLTEGKMLVAMYAG	KVDYTVTGKGFD	PT	VGKITTT
Chlamydomonas_reinhardtii_5	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GTAYP	QVPEGS	
Chlamydomonas_reinhardtii_6	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GTAYP	QVPEAS	
Volvox_carteri_3	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GTSFE	SVPPPE	
Volvox_carteri_4	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GTSFE	SVPPPE	
Volvox_carteri_5	ETMGGATAICSDKTGTLTENRMTVVEGWFG	GKKYD	QAPRSG	
Chlamydomonas_reinhardtii_7	ETMGGATAICSDKTGTLTENRMTVVEGWFG	GRHFS	TAPKAN	
Chlamydomonas_reinhardtii_8	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GQQFD	HLPDPS	
Volvox_carteri_6	ETMGGATAICSDKTGTLTENRMTVVEGWFA	GKSYD	HCPQPE	
Ostreococcus_lucimarinus_3	ETMGSATQLNADKTGTLTQNRMTVTACWLG	GKVC	EQVPPP	
Micromonas_sp_4	ETMGSATQLNADKTGTLTQNRMTVTDAYLG	GTQY	DSVPPD	
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Phaeodactylum_tricornutum_4	ETMGNATNICSDKTGTLTENRMTVVKGIFA	DTRC	DD TINRNPV	
Thalassiosira_pseudonana_6	ETMGCATTICSDKTGTLTANKMTARAIYTT	KTDFS		
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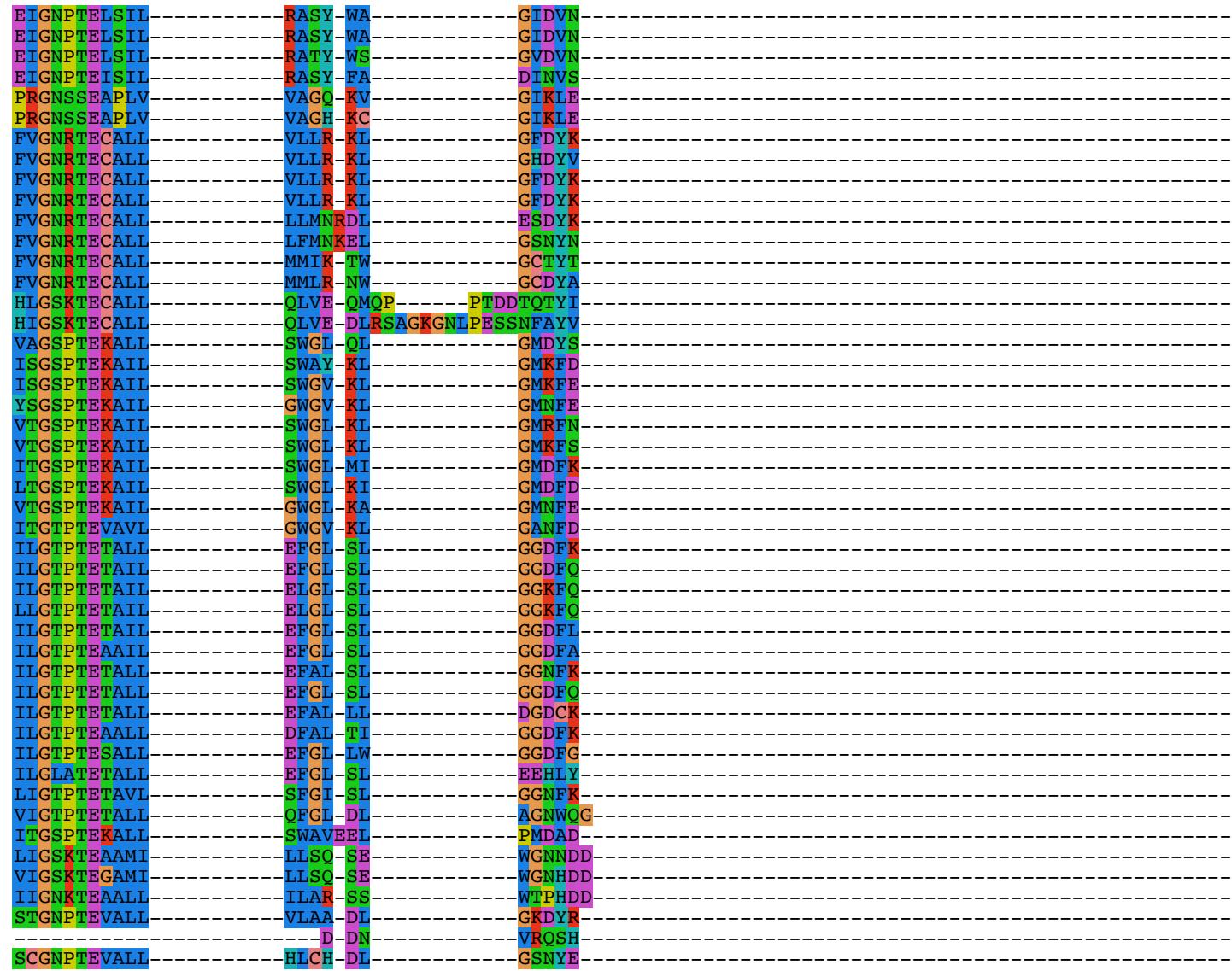




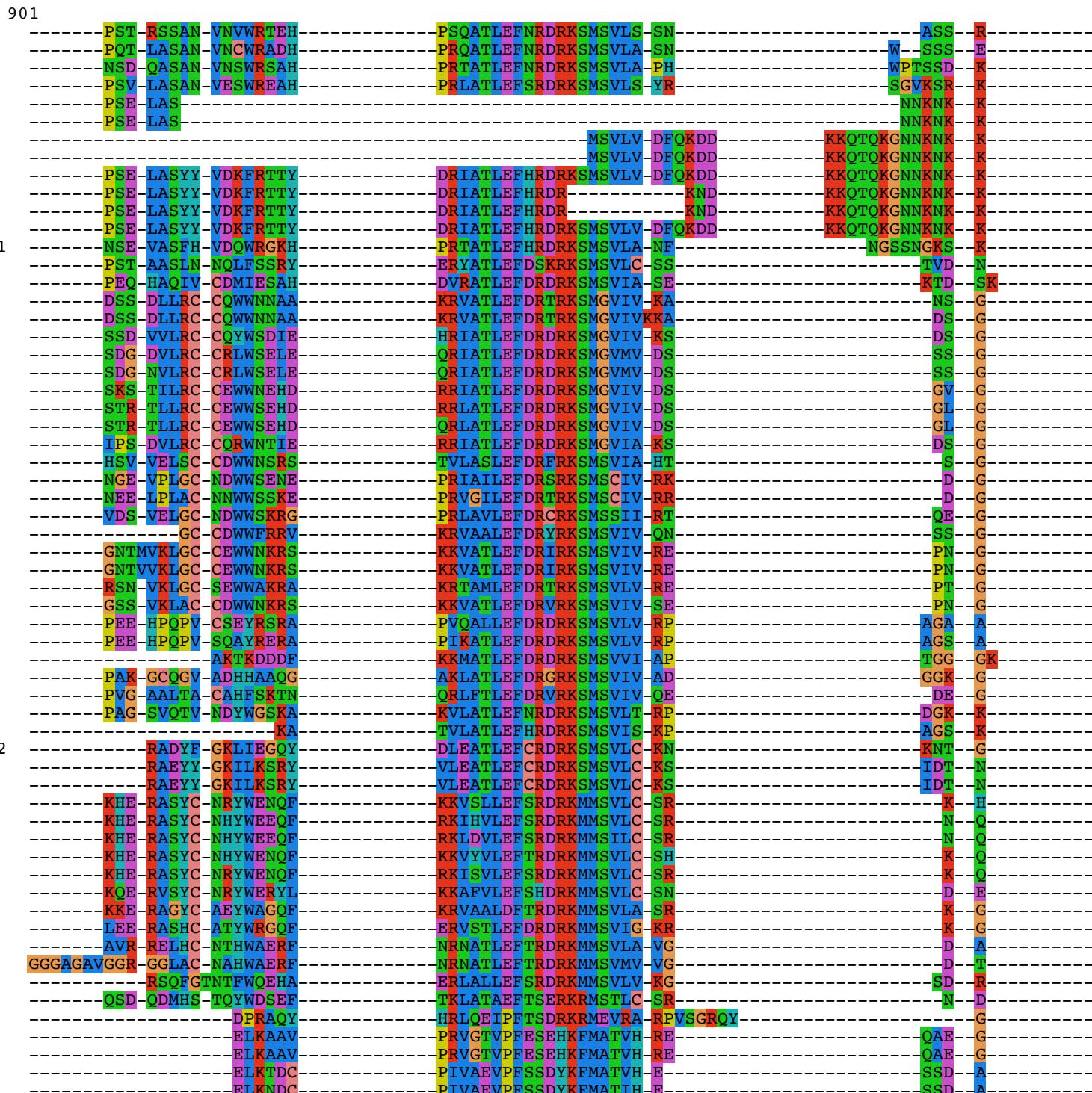
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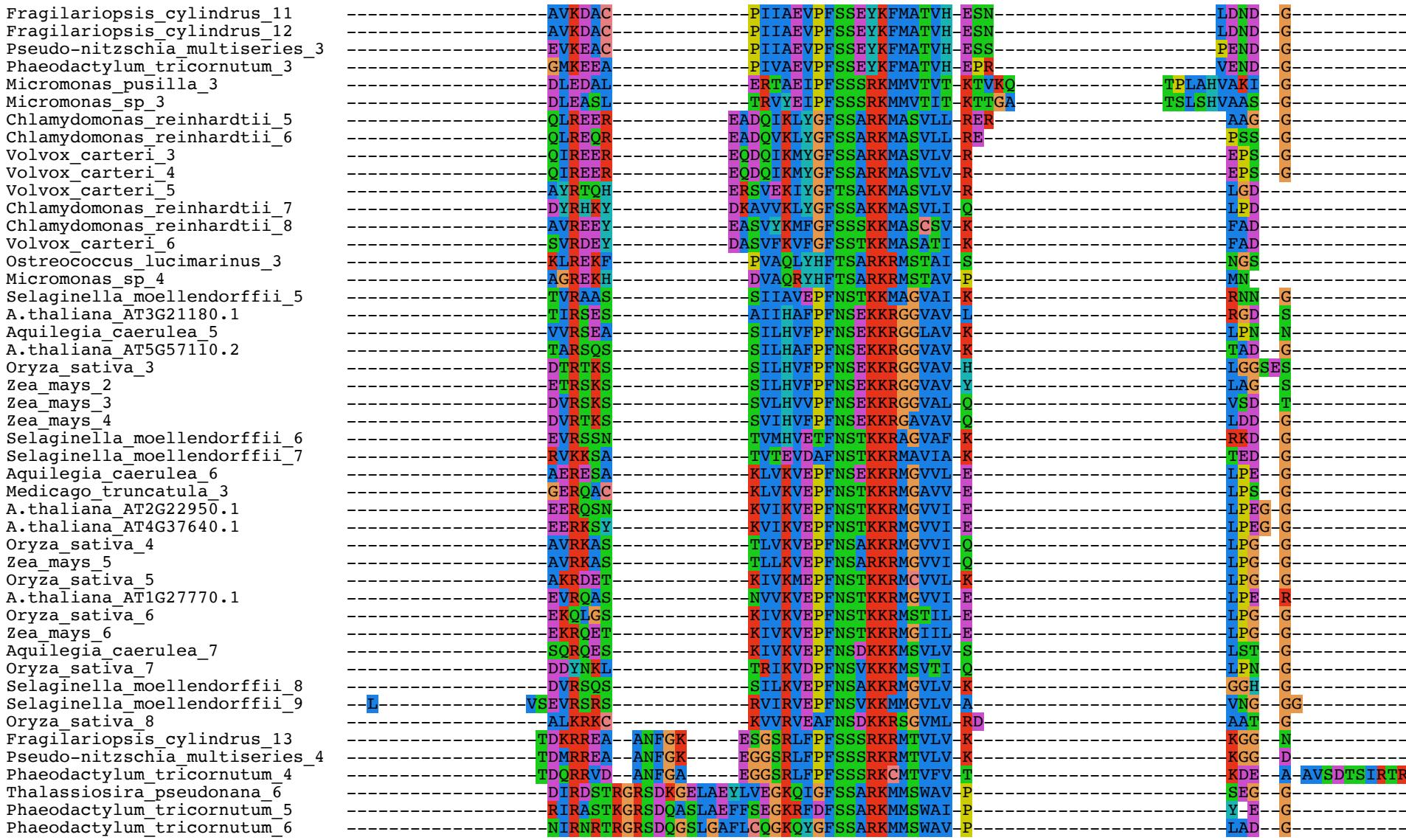


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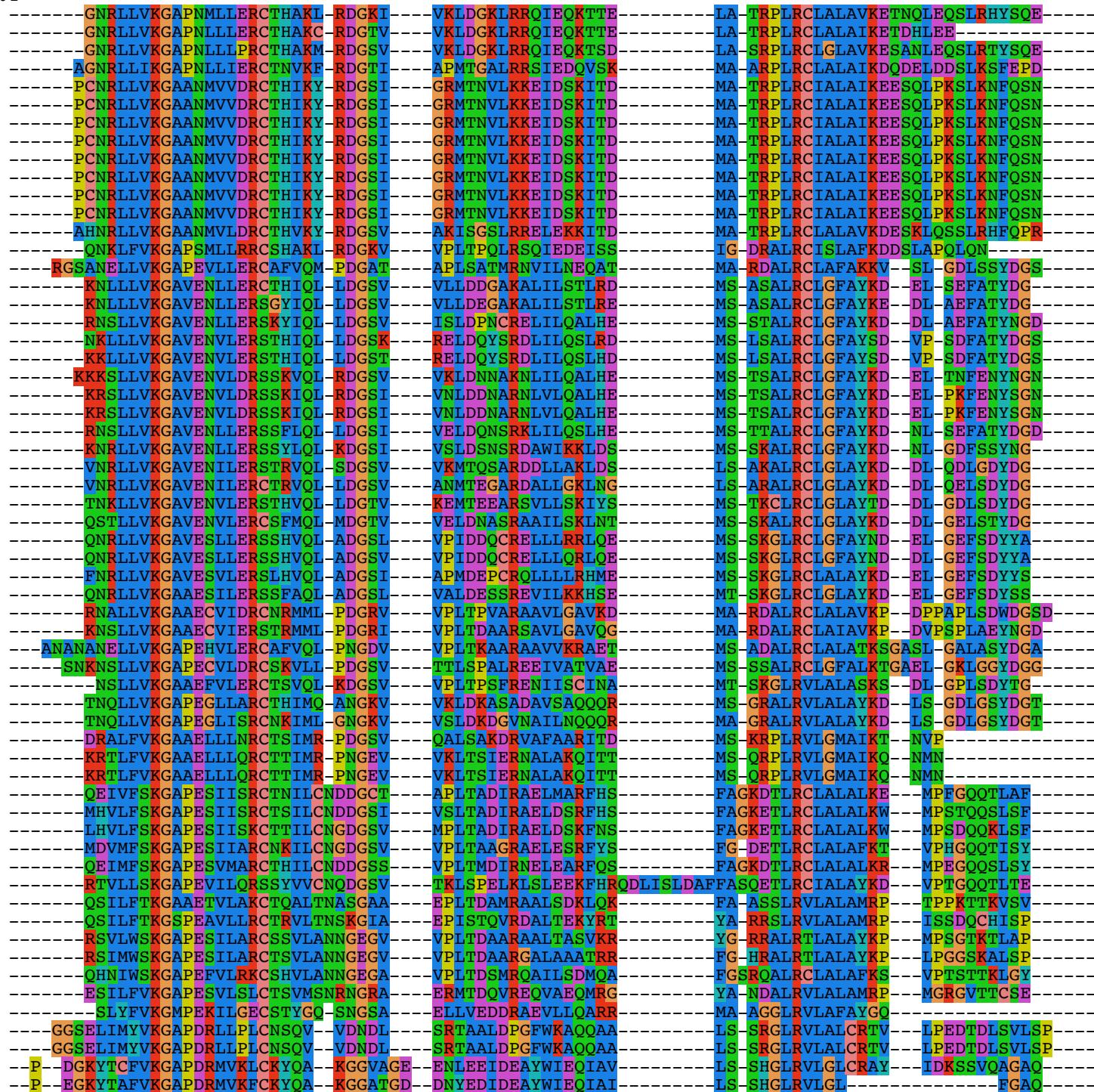
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Chlamydomonas_reinhardtii_3
Chlamydomonas_reinhardtii_4
Thalassiosira_pseudonana_4
Thalassiosira_pseudonana_5





1001

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 Thalassiosira_pseudonana_1
 Thalassiosira_oceanica_1
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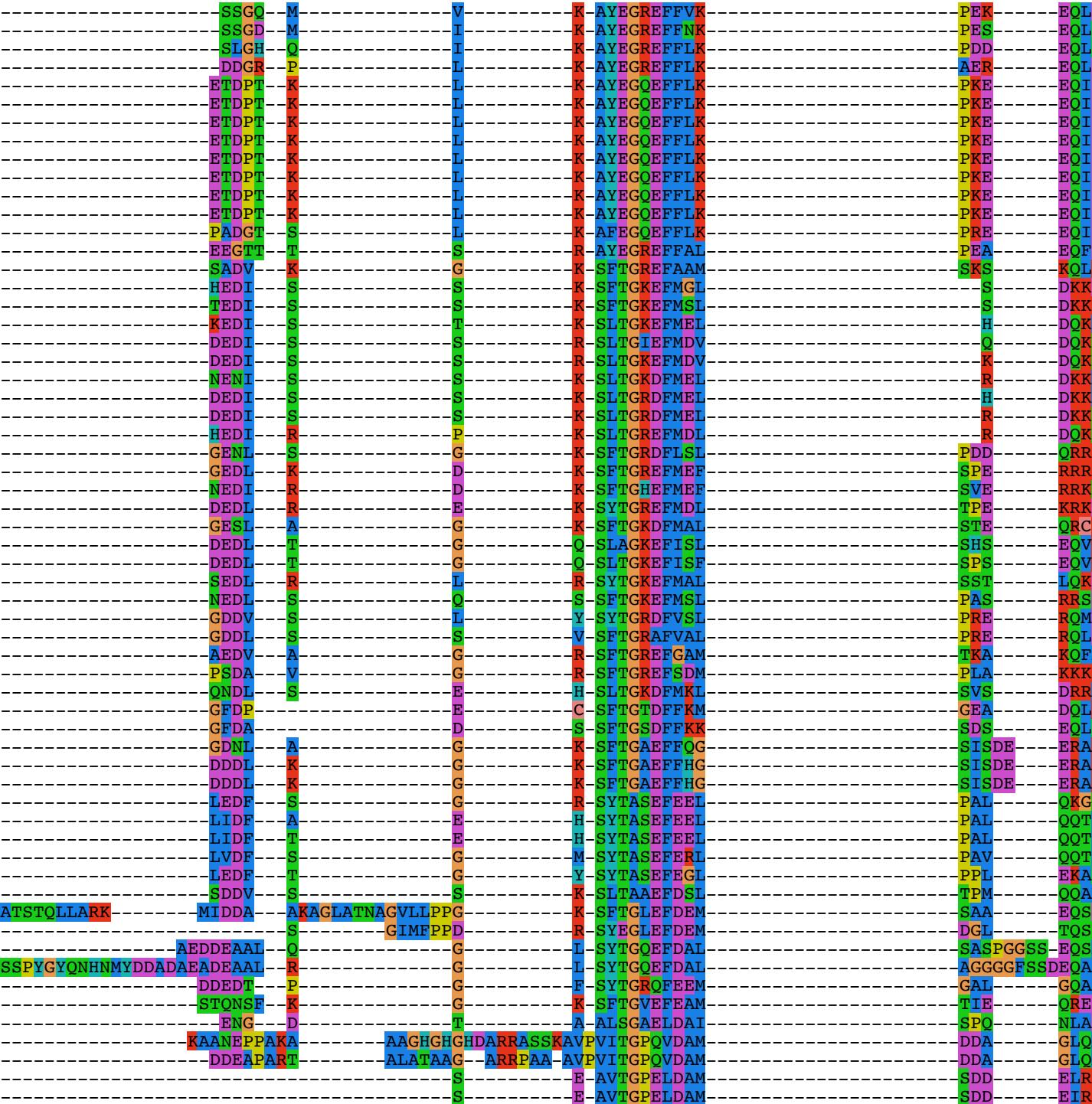
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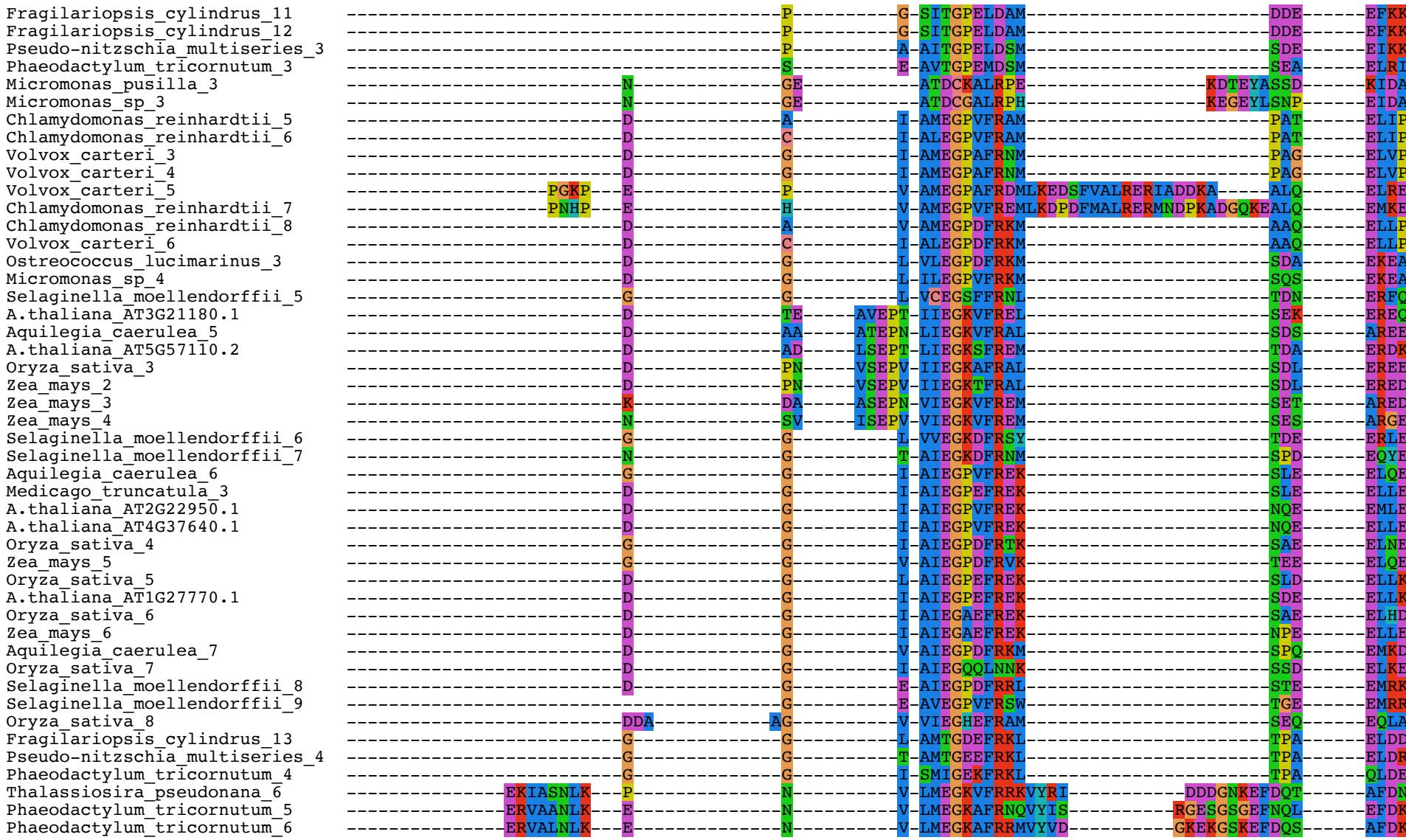
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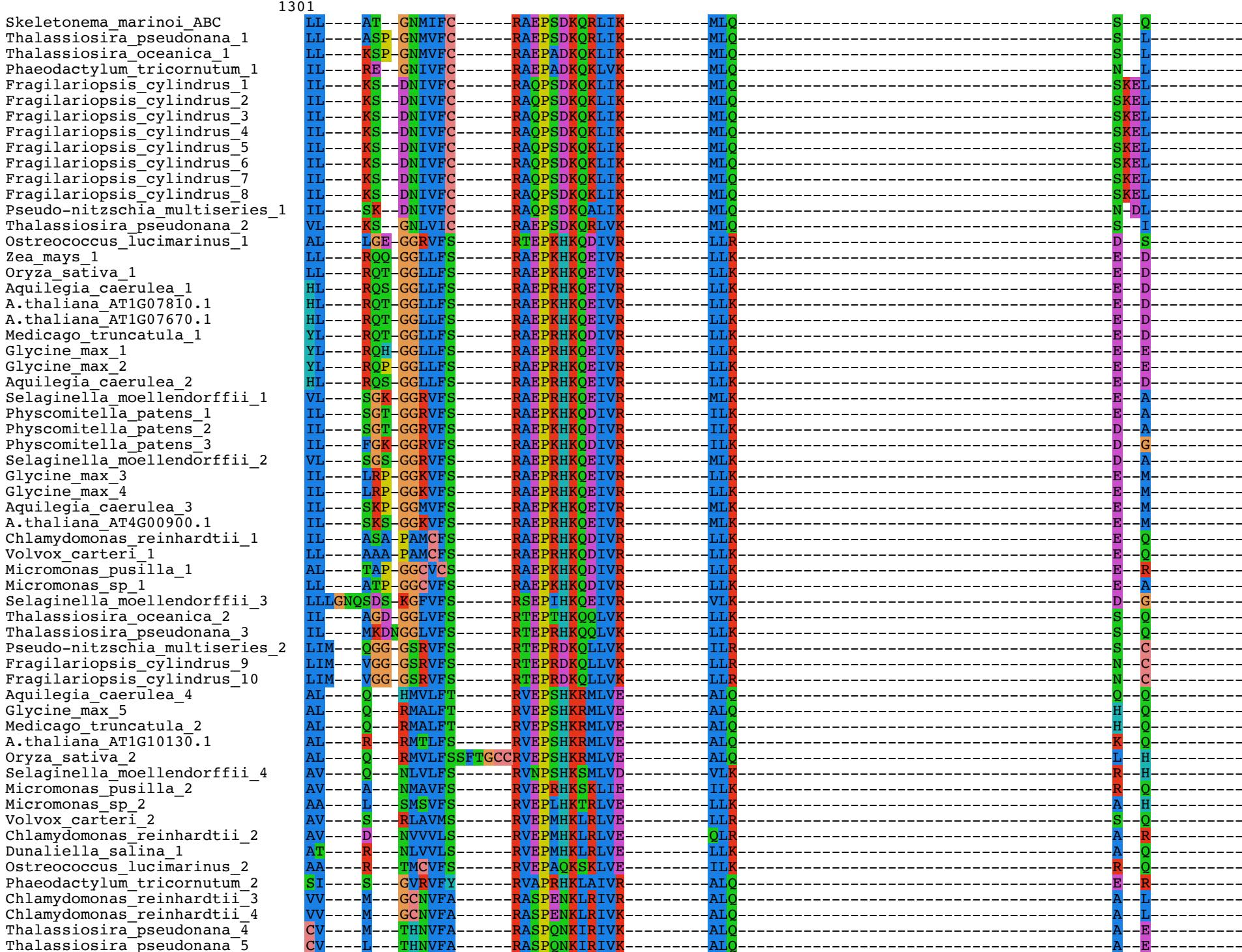
The alignment shows a high degree of conservation in the sequence region starting at position 1101. Many positions are occupied by the same amino acid across all or most of the listed species. For example, positions 1101 through 1110 are mostly occupied by 'P' (Proline) in the first half of the list and by 'D' (Aspartic acid) in the second half. Other positions like 1102, 1103, 1104, 1105, 1106, 1107, 1108, 1109, and 1110 show a mix of 'P', 'D', 'E', 'R', 'S', 'T', 'V', 'W', 'Y', and 'F'. The color-coding highlights the distribution of these amino acids, with darker shades indicating higher frequency in a specific position.

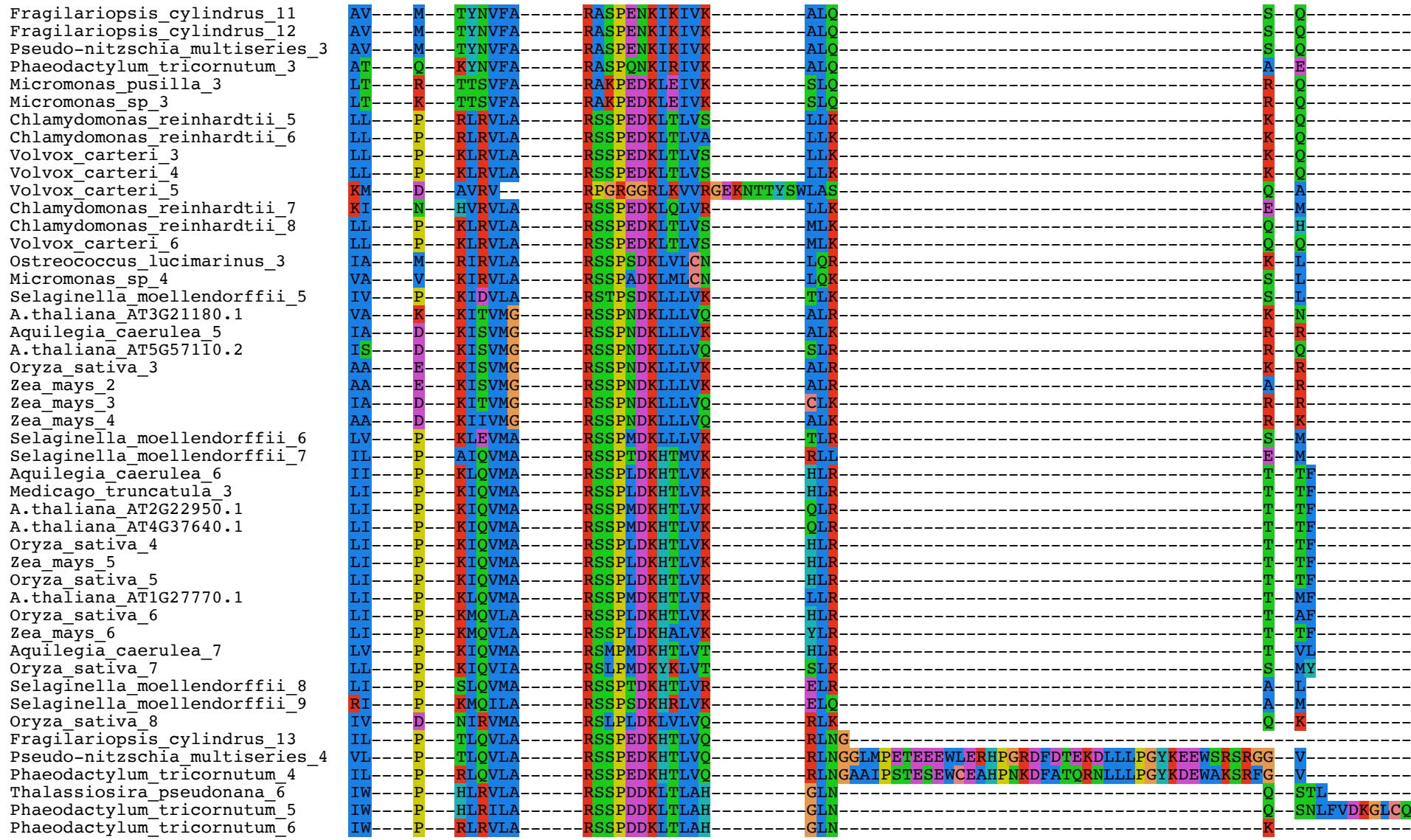
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Phaeodactylum_tricornutum_3	-	R	-GEPWLTI	VGLCAIMDPPRPECQ	QAAEAHAG	AVRVAMITGDHKDTALAIGSMILGLV	DREH	-						
Micromonas_pusilla_3	-	DTDVFKAK	-LVNDLTFCGLCASIDPERDGKVDAVRTSKV	A	AGVVRVVMITGDYI	KTAIAIAKNIGIL	NAKTFTED	-						
Micromonas_sp_3	-	ETDAKFDFK	-IVNGLTICGLCASIDPERDGKVDSVLQSROAQ	G	GRVVRVVMITGDYI	TKTAIAIAKNINIL	NRHTFTEG	-						
Chlamydomonas_reinhardtii_5	-	ARPA	-DFFEDADQ	VDRDLIAAVGIKDPVRKEVPDAVATCQKAGIV	V	VRMVITGDNIHTAQHIARECGLLT	T	E						
Chlamydomonas_reinhardtii_6	-	ARPA	-DFFEDADQ	VDNGLTCLAVGIKDPVRKEVPDAVTCQKAGIV	V	VRMVITGDNIHTAQHISRECGIL	VE	-						
Volvox_carteri_3	-	VRTP	-DFFEDADQ	VDNGLIACAIVGIKDPVRRAEVPAVTCQAA	G	AGIVVRMVITGDNIHTARHIARECGL	VD	-						
Volvox_carteri_4	-	VRTP	-DFFEDADQ	VDNGLIACAIVGIKDPVRRAEVPAVTCQAA	G	AGIVVRMVITGDNIHTARHIARECGL	VD	-						
Volvox_carteri_5	-	KRPA	-DFFEDVDN	VDNDLTCIGIVGIKDPVRTEVPDAVTCRAGIV	T	VRMVITGDNIHTARHIARECGL	YDMG	-						
Chlamydomonas_reinhardtii_7	-	SRPE	-NFFEEADT	VDDNLTCIGIVGIKDPVRRAEVPLAVTCRAGIV	V	VRMVITGDNIHTAQHIARECGL	YDMG	-						
Chlamydomonas_reinhardtii_8	-	SRPA	-DFFEDSDN	LDRNLVALAIVGIKDPVRKEVPEAVRCVQ	A	AGITVRMVITGDNIHTAQHIARECGL	TE	-						
Volvox_carteri_6	-	SRPL	-DFFEDSDY	LDRNLVAMAIVGIKDPVRKEVPEAVRCVQ	A	AGITVRMVITGDNIHTAQHIARECGL	TD	-						
Ostreococcus_lucimarinus_3	-	SLGETP	-PESDITLLGIMGIKDPVRPEA	EAVRLLRGAG	AVRVMVTGDNIHTAEAIAREAGIL	EDGD	-							
Micromonas_sp_4	-	GLSDEP	-PEANITLLGITGIKDPVRPEA	EVRLRQAG	AVRVMVTGDNIHTAEAIAREAGIL	EDGD	-							
Selaginella_moellendorffii_5	-	SMDGRPI	-PTAGLT	TFVALVGIKDPCRPGVREAVRK	CQDAGV	KVRMVTGDNLITARAIASECGL	MP	-						
A.thaliana_AT3G21180.1	-	PKEQ	-EDLDKWAL	PEDELILL	LAIVGIKDPCRPGVREAVRICKTSAGV	KVRMVTGDNLQTAIALECGIL	SS	-						
Aquilegia_caerulea_5	-	PKDE	-DQLEQWVI	PEEDLVLL	LAIVGIKDPCRPGVKNAVQLCTAGV	KVRMVTGDNLQTAIALECGIL	DS	-						
A.thaliana_AT5G57110.2	-	PTGE	-E	LSKWVL	PEDDLLLAIVGIKDPCRPGVKDSV	VLCQAGV	KVRMVTGDNLQTAIALECGIL	SS	-					
Oryza_sativa_3	-	PSED	-R	RADWIL	PEDDLIMI	GIVGIKDPCRPGVKDSVRLCAAAGIK	VVRMVITGDNLQTAIALECGIL	S	-					
Zea_mays_2	-	PDED	-R	REEWQL	PEDDLIMI	GIVGIKDPCRPGVRDSVRLCQAA	GIKVRMVITGDNLQTAIALECGIL	D	-					
Zea_mays_3	-	PMES	-	LDWKWL	PEDDLIMI	GIVGIKDPCRPGVRDSVRLCQAA	GIKVRMVITGDNLQTAIALECGIL	DA	-					
Zea_mays_4	-	PMED	-	ITSWEL	PEDDLIMI	GIVGIKDPCRPGVRDAVRLCTTAGV	KVRMVTGDNLQTAIALECGIL	DA	-					
Selaginella_moellendorffii_6	-	PDES	-E	WESWKI	PEDDLVLL	GIMGIKDPCRPGVDGA	VRLCOKAGV	KVRMVTGDNLQTAIALECGIL	SP	-				
Selaginella_moellendorffii_7	-	PPKKH	-S	TIGPPI	PEDGLTCIAIVGIKDP	PCRPVPEAVHKCQIAGIK	VVRMVITGDNIITAKAI	IAVECGIL	T	-				
Aquilegia_caerulea_6	-	SDKDSI	-	PLKGYTCI	GIVGIKDPVRPGV	KESVAICRSAGITVRMV	ITGDNIINTAKAI	ARECGIL	TD	-				
Medicago_truncatula_3	-	SAEVTI	-	PVTGYTCI	GIVGIKDPVRPGV	KESVALCRSAGITVRMV	ITGDNIINTAKAI	ARECGIL	TD	-				
A.thaliana_AT2G22950.1	-	SADEGI	-	PEKGFTCI	GIVGIKDPVRPGV	RESVELCRRAGIMVRMV	ITGDNIINTAKAI	ARECGIL	TD	-				
A.thaliana_AT4G37640.1	-	SPDDA	-	PASGFTCV	GIVGIKDPVRPGV	KESVELCRRAGITVRMV	ITGDNIINTAKAI	ARECGIL	TD	-				
Oryza_sativa_4	-	SANDQI	-	PEDGYTCI	GIVGIKDPVRPGV	KESVAICRSAGIMVRMV	ITGDNIINTAKAI	ARECGIL	TE	-				
Zea_mays_5	-	SPSEOI	-	PTDGYTCI	CIVVGIKDPVRPGV	KESVAICRSAGITVRMV	ITGDNIINTAKAI	ARECGIL	TD	-				
Oryza_sativa_5	-	SVEEQI	-	PLQGYTCI	GIVGIKDPVRPGV	RESVATCRSAGIMVRMV	ITGDNIINTAKAI	ARECGIL	TE	-				
A.thaliana_AT1G27770.1	-	SLEAPI	-	PSGGYTCI	GIVGIKDPVRPGV	KESVAICKSAGITVRMV	ITGDNIITAKAI	ARECGIL	TD	-				
Oryza_sativa_6	-	STOEQI	-	PLQGYTCI	GIVGIKDPVRPGVRQS	VATCRSAGISVRM	ITGDNIIDTAKAI	ARECGIL	TK	-				
Zea_mays_6	-	SVDEQI	-	PLQGYTCI	GIVGIKDPVRPGVRQS	VATCRSAGIAVRMV	ITGDNIINTAKAI	ARECGIL	TE	-				
Aquilegia_caerulea_7	-	SSGDSL	-	PSEGTYTLIAV	VGIKDPVRPGV	KEAVQTC	CLAAGITVRMV	ITGDNIINTAKAI	ARECGIL	TD	-			
Oryza_sativa_7	-	EFPNDQPI	-	SDDGYTLIAV	FGIKDPVRPGV	DAVRTCMAAGIR	VRMVTGDNI	INTAKAI	ARECGIL	TE	-			
Selaginella_moellendorffii_8	-	PAEDKL	-	PDNGFTCI	GIVGIKDPVRPGV	REAVOLCFAAGIK	VVRMV	ITGDNIINTAVAI	ARECGIL	T	-			
Selaginella_moellendorffii_9	-	QGEEAL	-	PQQGFVCAG	IIVGIKDPVRPGV	EEAVRMCSAGIR	VVRMV	ITGDNLYTAMAI	ARECGIL	TD	-			
Oryza_sativa_8	-	SDNAKI	-	DDEGLTLLG	FGVGLKDPCRPEVK	SAIEACTKAGIAV	KMV	ITGDNLITACAI	AKOAGIL	SGND	-			
Fragilariopsis_cylindrus_13	-	DISTF	-SIADCESI	LEKDMCLDALV	GIADPLRGD	VIEAVTCQRCGIF	VVRMV	ITGDNL	ETACAI	AKOAGIL	TE	-		
Pseudo-nitzschia_multiseries_4	-	DISKV	-VVGDCEST	LEKDMCLDALV	GI	SDPLREDVVDAIA	VCOKAGIF	VVRMV	ITGDNL	ETACAI	AKOAGIL	TK	-	
Phaeodactylum_tricornutum_4	-	DPQTV	-TQQDCEKK	LEKEMCL	DAIA	GIMDPLRPDV	VEAVAI	QRCAGIF	VVRMV	ITGDNL	DTAEAI	AKOAGIL	TE	-
Thalassiosira_pseudonana_6	-	NVLN	-SDGSEALS	VETEL	VF	ALVGIEDPLRPEV	QEA	IKKCYSAGIDV	RLV	TDSPNTAVS	IAYOADIL	ODFHFRNDSD	-	
Phaeodactylum_tricornutum_5	-	SVIKN	-ADGTEANE	IETEMTSYCPHYRRPPSSR	SSR	PGAEV	PGAIKE	YCQAGIDV	RLV	TDSPNTAVS	IAYOADIL	RDFHFLEGSI	-	
Phaeodactylum_tricornutum_6	-	SEHTN	-ADGRDAYV	AETDLIAIALV	GIEDPLRAEV	PGAIKE	YCQAGIDV	RLV	TDSPNTAVS	IAYOADIL	RDFHFLDNTD	-		

Skeletonema_marinoi_ABC
 Thalassiosira_pseudonana_1
 Thalassiosira_oceanica_1
 Phaeodactylum_tricornutum_1
 Fragilaropsis_cylindrus_1
 Fragilaropsis_cylindrus_2
 Fragilaropsis_cylindrus_3
 Fragilaropsis_cylindrus_4
 Fragilaropsis_cylindrus_5
 Fragilaropsis_cylindrus_6
 Fragilaropsis_cylindrus_7
 Fragilaropsis_cylindrus_8
 Pseudo-nitzschia_multiseries_1
 Thalassiosira_pseudonana_2
 Ostreococcus_lucimarinus_1
 Zea_mays_1
 Oryza_sativa_1
 Aquilegia_caerulea_1
 A.thaliana_AT1G07810.1
 A.thaliana_AT1G07670.1
 Medicago_truncatula_1
 Glycine_max_1
 Glycine_max_2
 Aquilegia_caerulea_2
 Selaginella_moellendorffii_1
 Physcomitella_patens_1
 Physcomitella_patens_2
 Physcomitella_patens_3
 Selaginella_moellendorffii_2
 Glycine_max_3
 Glycine_max_4
 Aquilegia_caerulea_3
 A.thaliana_AT4G00900.1
 Chlamydomonas_reinhardtii_1
 Volvox_carteri_1
 Micromonas_pusilla_1
 Micromonas_sp_1
 Selaginella_moellendorffii_3
 Thalassiosira_oceanica_2
 Thalassiosira_pseudonana_3
 Pseudo-nitzschia_multiseries_2
 Fragilaropsis_cylindrus_9
 Fragilaropsis_cylindrus_10
 Aquilegia_caerulea_4
 Glycine_max_5
 Medicago_truncatula_2
 A.thaliana_AT1G10130.1
 Oryza_sativa_2
 Selaginella_moellendorffii_4
 Micromonas_pusilla_2
 Micromonas_sp_2
 Volvox_carteri_2
 Chlamydomonas_reinhardtii_2
 Dunaliella_salina_1
 Ostreococcus_lucimarinus_2
 Phaeodactylum_tricornutum_2
 Chlamydomonas_reinhardtii_3
 Chlamydomonas_reinhardtii_4
 Thalassiosira_pseudonana_4
 Thalassiosira_pseudonana_5

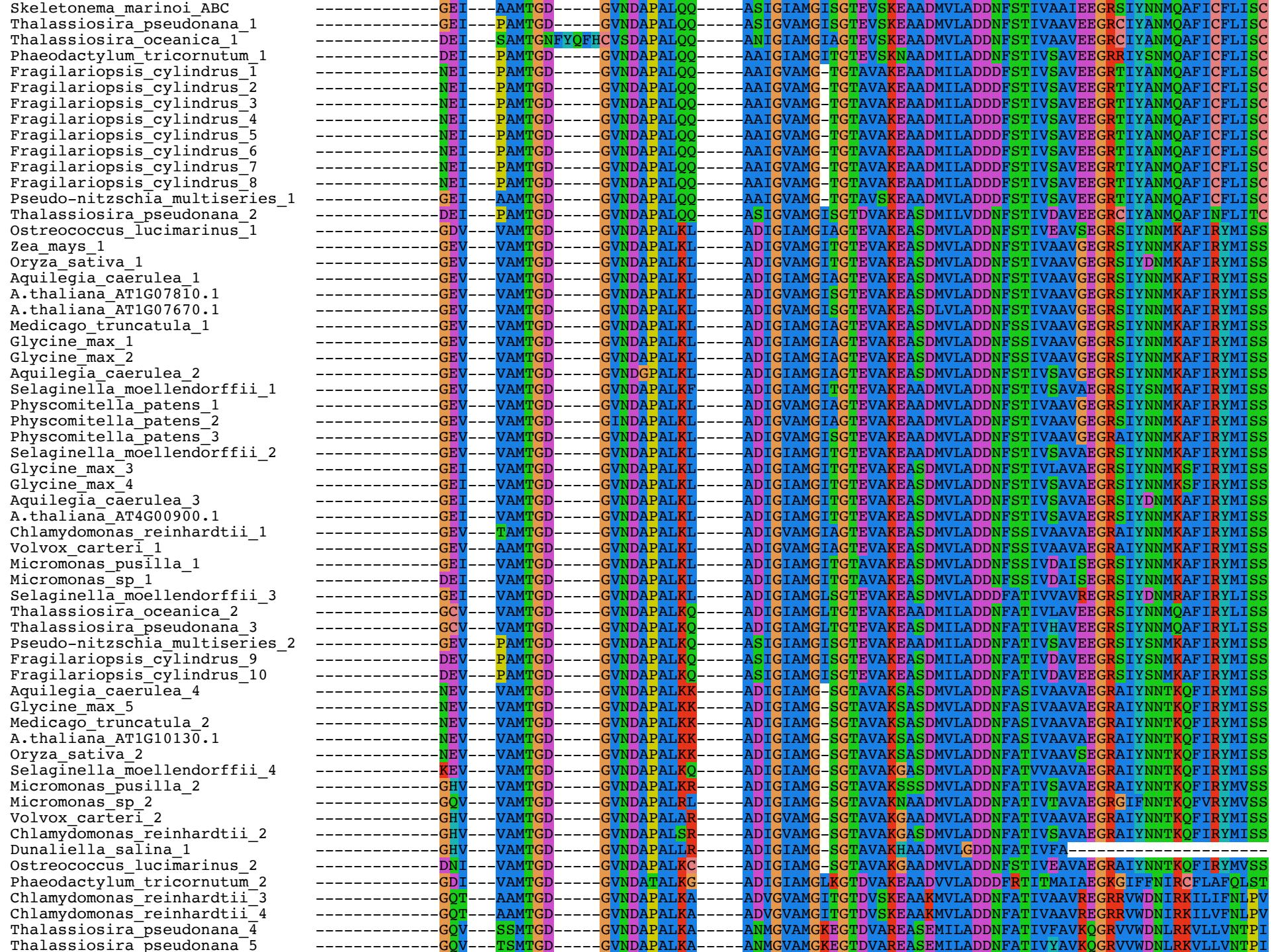








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Fragilariopsis_cylindrus_11	-----	GEV	ASMTGD	GVNDAPALKA	ANMGVAMGKEGTDVAREADMILADDNFATILTAVKEGRVVWDNLRKVLMINTPI
Fragilariopsis_cylindrus_12	-----	GEV	ASMTGD	GVNDAPALKA	ANMGVAMGKEGTDVAREADMILADDNFATILTAVKEGRVVWDNLRKVLMINTPI
Pseudo-nitzschia_multiseries_3	-----	GEV	ASMTGD	GVNDAPALKA	ANMGVAMGLEGTDVAREAAEMILADDNFATILSAVKEGRVVWDNLRKVLMINTPI
Phaeodactylum_tricornutum_3	-----	GEV	CGMTGD	GVNDAPALKA	ADMGVAMGKEGTDVAREAAEMILADDNFATIISAVREGRVVWDNLRKVLLINTPI
Micromonas_pusilla_3	-----	GWV	CAMTGD	GVNDAPALQR	ADIGVAMGLEGTTEVAKGASDMILTDDNFCSIVKAIEKGRIIYAGIQKFVAFIMSV
Micromonas_sp_3	-----	GFV	CAMTGD	GVNDAPALNR	ADIGVAMGLEGTTEVAKGASDMILTDDNFCSIVKAIEKGRITIYAGIQKFVFSFIMSV
Chlamydomonas_reinhardtii_5	-----	GEI	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIIIILDDNFSSIVKSVLWGRAVYMNIRKFLVFQLSI
Chlamydomonas_reinhardtii_6	-----	GEV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIIIILDDNFSSIVKSVLWGRWTYVMNIRKFLVFQLSV
Volvox_carteri_3	-----	GEV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIIIILDDNFSSIVKSVLWGRCVYMNIRKFLVFQLSI
Volvox_carteri_4	-----	GEV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIIIILDDNFSSIVKSVLWGRCVYMNIRKFLVFQLSI
Volvox_carteri_5	-----	PDSTFGVQSNGG	GGPRGERMPEGSTKGDCALGEADGGTEVSKEAADIVILDDNFSSIVKSVLWGRSFANIRKFLQFQLTV		
Chlamydomonas_reinhardtii_7	-----	GDV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIVILDDNFSSIVKSVLWGRSFANIRKFLQFQLTV
Chlamydomonas_reinhardtii_8	-----	GDV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIVILDDNFSSIVKSVLWGRSFVNIRKFLMFQQLTV
Volvox_carteri_6	-----	GEV	VAVTGD	GTNDAPALKE	SDVGLAMGIAGTEVAKEAADIVIMDDNFSSIVKSVLWGRSFVNIRKFLMFQQLTV
Ostreococcus_lucimarinus_3	-----	GEV	VAVTGD	GTNDAPALKD	ADVFGALGIAGTEIAKEACDIVILDDNNIKSMAKAVLWGRNVYQSIRKFLQFQLVV
Micromonas_sp_4	-----	GEV	VSVTGD	GTNDAPALKD	ADVFGALGIAGTEIAKEACDIVILDDNNIKSMAKAVLWGRNVFQSIRKFLQFQLVV
Selaginella_moellendorffii_5	-----	NEI	VAVTGD	GTNDAPALRE	AHIGLSMGIVGTEVAKESSDIIILDDNFASVVVKVHWGRSVYENIQKFQFQLTV
A.thaliana_AT3G21180.1	-----	GDV	VAVTGD	GTNDAPALHE	ADIGLSMGISGTEVAKESSDIIILDDNFASVVVKVHWGRSVYANIQKFQFQLTV
Aquilegia_caerulea_5	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESSDIIILDDNFSSVVVKVVRWGRSVYANIQKFQFQLTV
A.thaliana_AT5G57110.2	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESSDIIILDDNFSSVVVKVVRWGRSVYANIQKFQFQLTV
Oryza_sativa_3	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIQGTEVAKESSDIIILDDNFASVVVKVVRWGRSVYANIQKFQFQLTV
Zea_mays_2	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIQGTEVAKESSDIIILDDNFASVVVKVVRWGRSVYANIQKFQFQLTV
Zea_mays_3	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIQGTEVAKESSDIIILDDFTSVVVKVVRWGRSVYANIQKFQFQLTV
Zea_mays_4	-----	GHV	VAVTGD	GTNDAPALHE	ADIGLAMGIQGTEVAKESSDIIILDDFTSVVVKVVRWGRSVYANIQKFQFQLTV
Selaginella_moellendorffii_6	-----	NDV	VAVTGD	GTNDAPALHE	ADIGLAMGIQGTEVAKESSDIIILDDNFASVVVKVVRWGRSVYANIQKFQFQLTV
Selaginella_moellendorffii_7	-----	GEI	VAVTGD	GTNDAPALHE	ASIGLSMGIACTEVAKESSDIIIMDDDFASIVKVVRWGRAYANIQKFVQFQCTV
Aquilegia_caerulea_6	-----	GEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
Medicago_truncatula_3	-----	GEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
A.thaliana_AT2G22950.1	-----	DEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKEIADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
A.thaliana_AT4G37640.1	-----	DEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
Oryza_sativa_4	-----	DEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
Zea_mays_5	-----	DEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYINIQKFVQFQLTV
Oryza_sativa_5	-----	NEV	VAVTGD	GTNDAPALHE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYVNIQKFVQFQLTV
A.thaliana_AT1G27770.1	-----	QEV	VAVTGD	GTNDAPALHE	ADIGLAMGISGTEVAKESADVIILDDNFSTIVTVAKWGRSVYVNIQKFVQFQLTV
Oryza_sativa_6	-----	NEV	VAVTGD	GTNDAPALRE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYVNIQKFVQFQLTV
Zea_mays_6	-----	NEV	VAVTGD	GTNDAPALRE	ADIGLAMGIAGTEVAKESADVIILDDNFSTIVTVAKWGRSVYVNIQKFVQFQLTV
Aquilegia_caerulea_7	-----	GEV	VAVTGD	GTNDAPALHE	SDIGLAMGIAGTEVAKENADIIMDDNFSTIVNVAKWGRAYINIQKFVQFQLTV
Oryza_sativa_7	-----	QEV	VAVTGD	GTNDAPALHE	SDIGLAMGITGTEVAKESADVIIMDDNFETIVNVARWGRAYLNQKFVQFQLTV
Selaginella_moellendorffii_8	-----	DEV	VSVTGD	GTNDAPALHE	ADVGLAMGIAGTEVAKESADIVILDDRFNTIVVVAKWGRSVYTNQKFVQFQLTV
Selaginella_moellendorffii_9	-----	GEV	VGVTGD	GTNDAPALRE	ADIGISMGIAGTEVAKESSDIIILDDNFASIVNVALWGRSVYTNQKFVQFQATV
Oryza_sativa_8	-----	GHV	VAVTGD	GTNDAPALKE	ADVGLSMGVQGTEVAKESSDIVILNDNFDTVVATRWGRCVYNNQKFQFQLTV
Fragilariopsis_cylindrus_13	-----	GEV	VGVTGD	GTNDGPALKA	ADVGLSMGLSGTDDVAKNASDIIIMDDKFSSIVKAVLWGRSVFDNIRKFLQFQLTV
Pseudo-nitzschia_multiseries_4	-----	GEV	VGVTGD	GTNDGPALKA	ADVGLSMGLSGTDDVAKNASDIIIMDDKFSSIVKAVLWGRSVFDNIRKFLQFQLTV
Phaeodactylum_tricornutum_4	-----	EDNIIIFPDRQV	VAMTGD	GTNDAPALKR	ADIGFAMGIAGTQIAKDAADIILLDDNFASIVTAAKWGRNVYASIQKFQFQLTV
Thalassiosira_pseudonana_6	-----	KYNLEDGINIFPDRQV	IAMTGD	GTNDAPALKR	ADIGFAMGIAGTQIAKDAADIILLDDNFASIVTAAKWGRNIYASIQKFQFQLTV
Phaeodactylum_tricornutum_5	-----	KYNLEDGINIFPDRQV	IAMTGD	GTNDAPALKR	ADIGFAMGIAGTQIAKDAADIILLDDNFASIVTAAKWGRNVYASIQKFQFQLTV
Phaeodactylum_tricornutum_6	-----	EGIKVFPDRQV	IAMTGD	GTNDAPALKR	ADIGFAMGIAGTQIAKDAADIILLDDNFASIVTAAKWGRNVYASIQKFQFQLTV

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Skeletonema_marinoi_ABC	NIGEIAAILFAAVCGFPE	PLSAMHLLWVNLTGDPPATALGFNPPSPDVMTQKPR
Thalassiosira_pseudonana_1	NIGEIAAILISAVCGFPE	PLSAMHLLWVNLTGDPPATALGFNPPADVMSQKPR
Thalassiosira_oceanica_1	NIGEIAILLSTLCGFPE	PLSAMHLLWVNLTGDPPATALGFNPPADVMQKQPR
Phaeodactylum_tricornutum_1	NIGEICAIFFATLAGFPE	PLTAMHLLWVNLTGDPPATALGFNPPAPDLMEQPPR
Fragilariopsis_cylindrus_1	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_2	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_3	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_4	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_5	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_6	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_7	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Fragilariopsis_cylindrus_8	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPLPNIMKQOPPR
Pseudo-nitzschia_multiseries_1	NIGEICAIFFATLFGFPE	PLTAMHLLWVNLTGDPPATALGFNPPSPNSMKEKPR
Thalassiosira_pseudonana_2	NIGEVIGVFLATILGFPQ	LLTPLHLLWVNLTGDPPATALGFNPPGPVMAQKPR
Ostreococcus_lucimarinus_1	NVGEVVSIFLTAALGMPE	GLVPVQLLWVNLTGDPPATALGFNPPDKDIMTKPPR
Zea_mays_1	NIGEVASIFLTSALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Oryza_sativa_1	NIGEVASIFLTSALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Aquilegia_caerulea_1	NIGEVASIFLTAALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
A.thaliana_AT1G07810.1	NIGEVASIFLTAALGIPE	GMIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
A.thaliana_AT1G07670.1	NIGEVASIFLTAALGIPE	GMIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Medicago_truncatula_1	NIGEVASIFLTAALGIPE	GMIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Glycine_max_1	NIGEVASIFLTAALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Glycine_max_2	NIGEVASIFLTAALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Aquilegia_caerulea_2	NIGEVASIFLTAALGIPE	GLIPVQLLWVNLTGDPPATALGFNPPDKDIMKKPPR
Selaginella_moellendorffii_1	NIGEVASIFLTAAVGMPE	GLTPVQLLWVNLTGDPPATALGFNPPDVDIMRKPPR
Physcomitella_patens_1	NIGEVASIFMTAALGMPE	GLVPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Physcomitella_patens_2	NIGEVASIFMTAALGMPE	GLVPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Physcomitella_patens_3	NMGEVASIFLTAALGLPE	GLIPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Selaginella_moellendorffii_2	NIGEVASIFLTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Glycine_max_3	NIGEVASIFLTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Glycine_max_4	NIGEVASIFLTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPPDLDIMQKPPR
Aquilegia_caerulea_3	NIGEVASIFLTAALGMPE	CMISVQLLWVNLTGDPPATALGFNPADVDIMQKPPR
A.thaliana_AT4G00900.1	NIGEVASIFLTAALGMPE	CMIPVQLLWVNLTGDPPATALGFNPADVDIMQKPPR
Chlamydomonas_reinhardtii_1	NIGEVASIFLTAALGLPE	CMIPVQLLWVNLTGDPPATALGFNPADVDIMQKPPR
Volvox_carteri_1	NIGEVASIFLTAALGLPE	GLIPVQLLWVNLTGDPPATALGFNPADVDIMQKPPR
Micromonas_pusilla_1	NVGEVVSIFLTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPADVDIMTKTPR
Micromonas_sp_1	NVGEVVSIFLTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPADVDIMTKTPR
Selaginella_moellendorffii_3	NIGEVVAIFLTAALGMPO	GLIPVQLLWVNLTGDGAATALGFNPPDTDIMDRPPR
Thalassiosira_oceanica_2	NIGEVAAIFFTAALGLPE	GLIPVQLLWVNLTGDPPATALGFNPPDADIMKKLPR
Thalassiosira_pseudonana_3	NIGEVAAIFFTAALGMPE	GLIPVQLLWVNLTGDPPATALGFNPADKDIMKKLPR
Pseudo-nitzschia_multiseries_2	NIGEVVSIFLTSTIIGCPE	GLIPVQLLWVNLTGDPPATALGFNPPDPDVMLKPPR
Fragilariopsis_cylindrus_9	NIGEVVSIFLTSSLGVPE	GLIPVQLLWVNLTGDPPATALGFNPPDPDVMLKPPR
Fragilariopsis_cylindrus_10	NIGEVVSIFLTSSLGVPE	GLIPVQLLWVNLTGDPPATALGFNPPDPDVMLKPPR
Aquilegia_caerulea_4	NIGEVVCIFVAAVLGVPE	TLVPVQLLWVNLTGDGLPATAIGFNKQDSDVMAKPR
Glycine_max_5	NIGEVVCIFVAAVLGVPE	TLAPVQLLWVNLTGDGLPATAIGFNKQDSDVRAKPR
Medicago_truncatula_2	NIGEVVCIFVAAVLGVPE	TLAPVQLLWVNLTGDGLPATAIGFNQDSDVVKVPR
A.thaliana_AT1G10130.1	NIGEVVCIFVAAVLGVPE	TLAPVQLLWVNLTGDGLPATAIGFNQDSDVVKVPR
Oryza_sativa_2	NIGEVVCIFVAAVLGMPD	TLVPVQLLWVNLTGDGLPATAIGFNKPDSNIMTVKPRHLHSATTIIRWLDAHGKOCFGIAFIVDVMGVDDLD
Selaginella_moellendorffii_4	NIGEVVCIFVAAVLGLPE	TLIPVQLLWVNLTGDGLPATAIGFNKPDSNIMTVKPRHLHSATTIIRWLDAHGKOCFGIAFIVDVMGVDDLD
Micromonas_pusilla_2	NIGEVVCIFIAAAALGMPE	TLCPVQLLWVNLTGDGLPATAIGFNKPDRDIMRARP
Micromonas_sp_2	NIGEVVCIFIAAAALGMPE	TLCPVQLLWVNLTGDGLPATAIGFNKPDRDIMRARP
Volvox_carteri_2	NIGEVVAIFVAALLGVPE	VLTVPVQLLWVNLTGDGLPATAIGFNKPDRDIMAVRPR
Chlamydomonas_reinhardtii_2	NIGEVVAIFVAALLGVPE	VLTVPVQLLWVNLTGDGLPATAIGFNKPDRDIMAVRPR
Dunaliella_salina_1	--GGVV--	
Ostreococcus_lucimarinus_2	NIGEVVCIFIAAAALGFPE	TLVPVQLLWVNLTGDGLPATAIGFNRADGDIMRORPR
Phaeodactylum_tricornutum_2	SFAALTMASIATAGLPS	PLNATOIILWINILMDGPPAQSLGVEPVDEKILRAKPR
Chlamydomonas_reinhardtii_3	NLAQGFSVLSYILSLDN	VPLTALQVLLVNLTTSVTLGLALAAEPPEPDIMERQPR
Chlamydomonas_reinhardtii_4	NLAQGFSVLYSYILGFKD	VPLTALQVLLVNLTTSVTLGLALAAEPPEPDIMERQPR
Thalassiosira_pseudonana_4	NNSOGLSVLVGLLVRLPN	TPITTIQILYSNFICAVTGLGFVCAIEPAEDGIMALP
Thalassiosira_pseudonana_5	NNSQGLCVFFGMLVRPN	TPITTIQILYSNFICAVTGLGFVCAVEPAEDGIMALP

Fragilariopsis_cylindrus_11
Fragilariopsis_cylindrus_12
Pseudo-nitzschia_multiseries_3
Phaeodactylum_tricornutum_3
Micromonas_pusilla_3
Micromonas_sp_3
Chlamydomonas_reinhardtii_5
Chlamydomonas_reinhardtii_6
Volvox_carteri_3
Volvox_carteri_4
Volvox_carteri_5
Chlamydomonas_reinhardtii_7
Chlamydomonas_reinhardtii_8
Volvox_carteri_6
Ostreococcus_lucimarinus_3
Micromonas_sp_4
Selaginella_moellendorffii_5
A.thaliana_AT3G21180.1
Aquilegia_caerulea_5
A.thaliana_AT5G57110.2
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Zea_mays_2
Zea_mays_3
Zea_mays_4
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Selaginella_moellendorffii_7
Aquilegia_caerulea_6
Medicago_truncatula_3
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Zea_mays_6
Aquilegia_caerulea_7
Oryza_sativa_7
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Selaginella_moellendorffii_9
Oryza_sativa_8
Fragilariopsis_cylindrus_13
Pseudo-nitzschia_multiseries_4
Phaeodactylum_tricornutum_4
Thalassiosira_pseudonana_6
Phaeodactylum_tricornutum_5
Phaeodactylum_tricornutum_6

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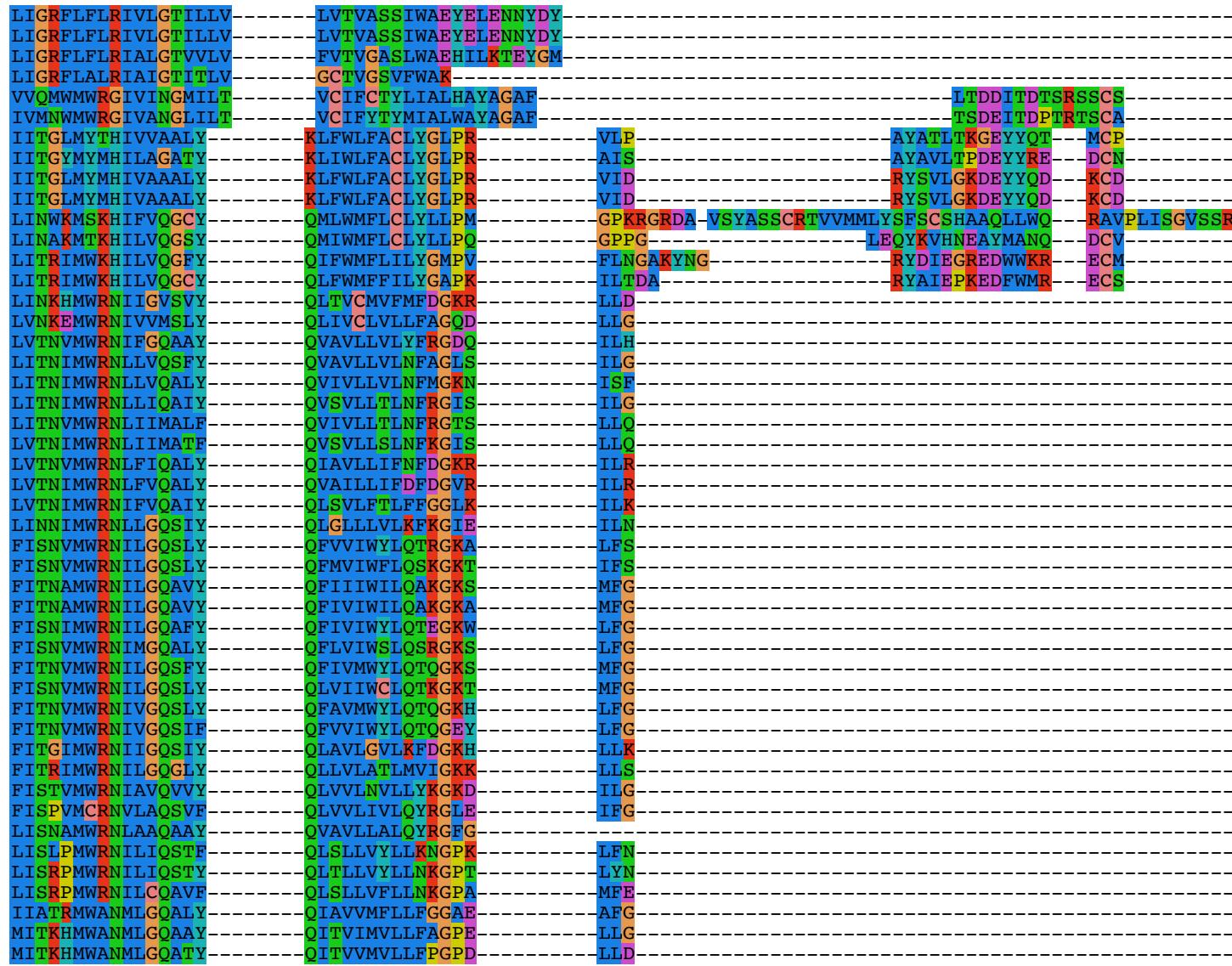
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Thalassiosira_pseudonana_1	PSNEP
Thalassiosira_oceanica_1	PSNEP
Phaeodactylum_tricornutum_1	PSDEP
Fragilaropsis_cylindrus_1	SSSEP
Fragilaropsis_cylindrus_2	SSSEP
Fragilaropsis_cylindrus_3	SSSEP
Fragilaropsis_cylindrus_4	SSSEP
Fragilaropsis_cylindrus_5	SSSEP
Fragilaropsis_cylindrus_6	SSSEP
Fragilaropsis_cylindrus_7	SSSEP
Fragilaropsis_cylindrus_8	SSSEP
Pseudo-nitzschia_multiseries_1	PSNEP
Thalassiosira_pseudonana_2	PTSEE
Ostreococcus_lucimarinus_1	RKDED
Zea_mays_1	RSDDS
Oryza_sativa_1	RSDDS
Aquilegia_caerulea_1	RSDDS
A.thaliana_AT1G07810.1	RSDDS
A.thaliana_AT1G07670.1	RSDDS
Medicago_truncatula_1	RSDDS
Glycine_max_1	HSDDS
Glycine_max_2	HSDDS
Aquilegia_caerulea_2	RSDDS
Selaginella_moellendorffii_1	RSDDK
Physcomitella_patens_1	KSDDV
Physcomitella_patens_2	KSNDV
Physcomitella_patens_3	KSTDV
Selaginella_moellendorffii_2	KSNDK
Glycine_max_3	RSDDP
Glycine_max_4	RNDDP
Aquilegia_caerulea_3	KSND A
A.thaliana_AT4G00900.1	KSDDC
Chlamydomonas_reinhardtii_1	RANDQ
Volvox_carteri_1	RANDQ
Micromonas_pusilla_1	KKDED
Micromonas_sp_1	RKDED
Selaginella_moellendorffii_3	IPTEG
Thalassiosira_oceanica_2	RTDDS
Thalassiosira_pseudonana_3	RADDN
Pseudo-nitzschia_multiseries_2	GKDD S
Fragilaropsis_cylindrus_9	GKNDA
Fragilaropsis_cylindrus_10	GKNDA
Aquilegia_caerulea_4	KVNEA
Glycine_max_5	KVNEA
Medicago_truncatula_2	KVNEA
A.thaliana_AT1G10130.1	KVG E A
Oryza_sativa_2	
Selaginella_moellendorffii_4	QCEHGLLCRLDG GGI SSD E D VSAI Q KI W HCLV QQTA AI T NPLRHCLRLV QCIAAAV SPTFLKLLPIGWTACCA S PSLICEGD T VLAYQQLV LLL E VNEA
Micromonas_pusilla_2	KMDEA
Micromonas_sp_2	RPDES
Volvox_carteri_2	RSDEP
Chlamydomonas_reinhardtii_2	RLDEP
Dunaliella_salina_1	RLDEP
Ostreococcus_lucimarinus_2	DDP
Phaeodactylum_tricornutum_2	SPREQ
Chlamydomonas_reinhardtii_3	KADEP
Chlamydomonas_reinhardtii_4	RRGKR
Thalassiosira_pseudonana_4	RRGKR
Thalassiosira_pseudonana_5	RVG KR

Fragilariopsis_cylindrus_11	RVGKR
Fragilariopsis_cylindrus_12	RVGKR
Pseudo-nitzschia_multiseries_3	RVGKR
Phaeodactylum_tricornutum_3	RVGKR
Micromonas_pusilla_3	PKTQP
Micromonas_sp_3	PKTOP
Chlamydomonas_reinhardtii_5	GRSEA
Chlamydomonas_reinhardtii_6	GRSEP
Volvox_carteri_3	GRTEP
Volvox_carteri_4	GRTEP
Volvox_carteri_5	GRTEH
Chlamydomonas_reinhardtii_7	GRTEQ
Chlamydomonas_reinhardtii_8	GRNEN
Volvox_carteri_6	GRNEN
Ostreococcus_lucimarinus_3	GRTAP
Micromonas_sp_4	GRSAP
Selaginella_moellendorffii_5	GLSEP
A.thaliana_AT3G21180.1	GRREP
Aquilegia_caerulea_5	GRREP
A.thaliana_AT5G57110.2	GRKEP
Oryza_sativa_3	GRREP
Zea_mays_2	GRREP
Zea_mays_3	GRREP
Zea_mays_4	GRREP
Selaginella_moellendorffii_6	GRREP
Selaginella_moellendorffii_7	SKEAP
Aquilegia_caerulea_6	GRKGN
Medicago_truncatula_3	GRKGN
A.thaliana_AT2G22950.1	GRRGN
A.thaliana_AT4G37640.1	GRRGN
Oryza_sativa_4	GRKGN
Zea_mays_5	GRKGN
Oryza_sativa_5	GRTGK
A.thaliana_AT1G27770.1	GRKGN
Oryza_sativa_6	GRKGK
Zea_mays_6	GRTGR
Aquilegia_caerulea_7	GKSVN
Oryza_sativa_7	RRGDN
Selaginella_moellendorffii_8	GRKGS
Selaginella_moellendorffii_9	RRKEN
Oryza_sativa_8	GRAAP
Fragilariopsis_cylindrus_13	KRNSS
Pseudo-nitzschia_multiseries_4	TRNSS
Phaeodactylum_tricornutum_4	RRDSS
Thalassiosira_pseudonana_6	NRSKS
Phaeodactylum_tricornutum_5	NRTES
Phaeodactylum_tricornutum_6	NRTDS

1701

Skeletonema_marinoi_ABC	IMTNWMLFRYLVTLGLYVG	FATVGSFVCHYLIS	
Thalassiosira_pseudonana_1	IMTKFMACRYLVTGLYVG	IATVGSFVGHYRS	
Thalassiosira_oceanica_1	IMTKWMAFRYLIITGLYVG	IATVGSFVSYYLD	
Phaeodactylum_tricornutum_1	IMTRWLLTRYCITGLYVG	LATIGIFQAQHYLS	
Fragilariopsis_cylindrus_1	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_2	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_3	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_4	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_5	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_6	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_7	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Fragilariopsis_cylindrus_8	IMTKWLLIRYMLTGLYVG	IATIGVFAQHYIK	
Pseudo-nitzschia_multiseries_1	IMTKWLLIRYMLTGLYVG	VATIGIFAQHYLO	
Thalassiosira_pseudonana_2	ILTPSLLLRYSTAGLYI	IATVGIYASFYVD	
Ostreococcus_lucimarinus_1	LLSNWVMFRYAVVGLYVG	VATVGAFAIWFTR	SOD
Zea_mays_1	LITPWILFRYMVIGLYVG	VATVGFIIWIYTH	ASD
Oryza_sativa_1	LITPWILFRYMVIGMYVG	IATVGVFIIWIYTH	AGD
Aquilegia_caerulea_1	LISPWILFRYLVIGLYVG	VATVGVFIIWIYTH	SGD
A.thaliana_AT1G07810.1	LITAWILFRYMVIGLYVG	VATVGVFIIWIYTH	SQD
A.thaliana_AT1G07670.1	LITAWILFRYMVIGLYVG	VATVGVFIIWIYTH	SQD
Medicago_truncatula_1	LINLWILFRYLVIGIYVG	LATVGVFIIWIYTH	SSD
Glycine_max_1	LINLWILFRYLVIGIYVG	LATVGVFIIWIYTH	SGD
Glycine_max_2	LINLWILFRYLVIGIYVG	LATVGVFIIWIYTH	SGD
Aquilegia_caerulea_2	LISAWILFRYLVIGLYVG	IATVGVFIIWIWTO	SGD
Selaginella_moellendorffii_1	LINSWVFFRYMVIGLYVG	LATVGI FALWYTH	SGD
Physcomitella_patens_1	LINAWVFFRYLVIGLYVG	VATVGAFAIWFTQ	AAD
Physcomitella_patens_2	LINGWVFFRYLVIGLYVG	IATVGAFAIWFTH	AAD
Physcomitella_patens_3	LINSWVLLRYLIIIGLYVG	IATVGAFSLWFTR	SSD
Selaginella_moellendorffii_2	LINAWVLLRYLIVIGLYVG	VATVGI FALWYTH	VSD
Glycine_max_3	LISSWVLFRYLVIGSYVG	LATVGI FVLYWYTO	VSD
Glycine_max_4	LISSWVLFRYLVIGSYVG	LATVGI FVLYWYTO	VSD
Aquilegia_caerulea_3	LIDSWVLFRYLVIGSYVG	IATVGVFILWYTO	AED
A.thaliana_AT4G00900.1	LIDSWVLIRYLVIGSYVG	VATVGI FVLYWYTO	ISD
Chlamydomonas_reinhardtii_1	LITPWVFFRYMVIGGYVG	VATVGAFWSWYMY	SRD
Volvox_carteri_1	LITPWVFFRYMVIGGYVG	VATVGAFWSWYMF	SRD
Micromonas_pusilla_1	LISAWALVRYLVVGLYVG	AATVGVFAWWYTR	SGD
Micromonas_sp_1	LISTWAMVRYLVVGLYVG	AATVGIFAWWYTK	AKD
Selaginella_moellendorffii_3	FISGWTLFRFLTIGLYVG	LATIGIEGLWLND	SRD
Thalassiosira_oceanica_2	LITPWVFFR YMVVGIYVG	FACVAVFAYWYIY	HEG
Thalassiosira_pseudonana_3	LITPWVFFR YMVVGIYVG	FACVGVFAYWYMY	YES
Pseudo-nitzschia_multiseries_2	LITNWVFFRYMVIGLYVG	FATVGI FVFWYIYEYES	ATFD
Fragilariopsis_cylindrus_9	LINNWVFFRYMVIGLYVG	FATVGI FIEWYIGYEP	STYD
Fragilariopsis_cylindrus_10	LINNWVFFRYMVIGLYVG	FATVGI FFWYIGYEP	STYD
Aquilegia_caerulea_4	VVNGWLFFFRLVIGAYVG	LATVAGFIWWFVYSEN	
Glycine_max_5	VVTGWLFRRYLVIGAYVG	LATVAGFIWWFVYSDS	
Medicago_truncatula_2	VVTGWLFRRYLVIGAYVG	LATVAGFIWWFVYADS	
A.thaliana_AT1G10130.1	VVTGWLFRRYLVIGYVG	LATVAGFIWWFVYSDG	
Oryza_sativa_2	VVNGWLFFFRLVIGAYVG	LATIAGFVWWFVYSED	
Selaginella_moellendorffii_4	IVNGWLFFFRLVVGAYVG	LATVGGFIWWFLYHEE	
Micromonas_pusilla_2	IVDRWLFLVRYLVVGMYVG	FVTVGAFAWWYMSYLD	
Micromonas_sp_2	IVDRWLFLVRYIIVVGIGYVG	LELPDVVFTLIGGFVWWYMFHSG	
Volvox_carteri_2	IVNGWLFLRVLVVGMYVG	LVTVAGFLWWFLGYOG	
Chlamydomonas_reinhardtii_2	IVNGWLFLRVLVVGMYVG	LATVGGFLWWFLAHEG	
Dunaliella_salina_1	IVNGWLFLRVLVVGMYVG	IVTVYGFIIWWYISYPE	
Ostreococcus_lucimarinus_2	IVDRWLLIRYLIIGVYVG	IATVGSFGWWFMSYPG	
Phaeodactylum_tricornutum_2	IVTRALLLRATSSAALIV	FLTILKVS	
Chlamydomonas_reinhardtii_3	LVGKLILLWRCCFFVCN	ALVLGMFYWGGED	
Chlamydomonas_reinhardtii_4	LVGKLILLWRMFFVCHVVV	ALVLGLFYWAGRGRG	
Thalassiosira_pseudonana_4	LIGRFLFLRIIVGTVLLT	CCVVA SAIIVDTI	PRYDY
Thalassiosira_pseudonana_5	LIGRYLFLRIILGTVIL	CCVVGALIVD	

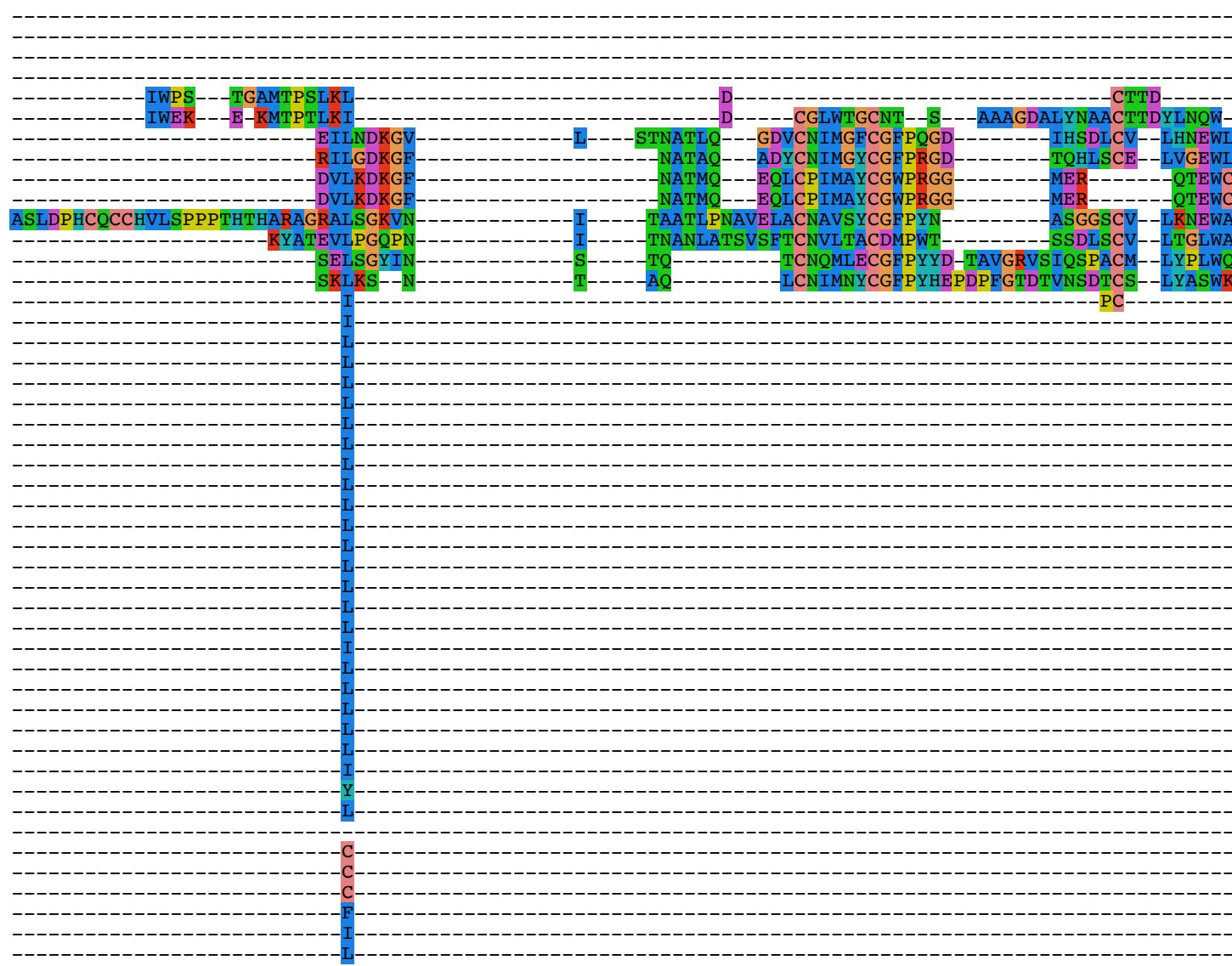
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 Fragilaropsis_cylindrus_12
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 Micromonas_pusilla_3
 Micromonas_sp_3
 Chlamydomonas_reinhardtii_5
 Chlamydomonas_reinhardtii_6
 Volvox_carteri_3
 Volvox_carteri_4
 Volvox_carteri_5
 Chlamydomonas_reinhardtii_7
 Chlamydomonas_reinhardtii_8
 Volvox_carteri_6
 Ostreococcus_lucimarinus_3
 Micromonas_sp_4
 Selaginella_moellendorffii_5
 A.thaliana_AT3G21180.1
 Aquilegia_caerulea_5
 A.thaliana_AT5G57110.2
 Oryza_sativa_3
 Zea_mays_2
 Zea_mays_3
 Zea_mays_4
 Selaginella_moellendorffii_6
 Selaginella_moellendorffii_7
 Aquilegia_caerulea_6
 Medicago_truncatula_3
 A.thaliana_AT2G22950.1
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 Oryza_sativa_4
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 Oryza_sativa_8
 Fragilaropsis_cylindrus_13
 Pseudo-nitzschia_multiseries_4
 Phaeodactylum_tricornutum_4
 Thalassiosira_pseudonana_6
 Phaeodactylum_tricornutum_5
 Phaeodactylum_tricornutum_6

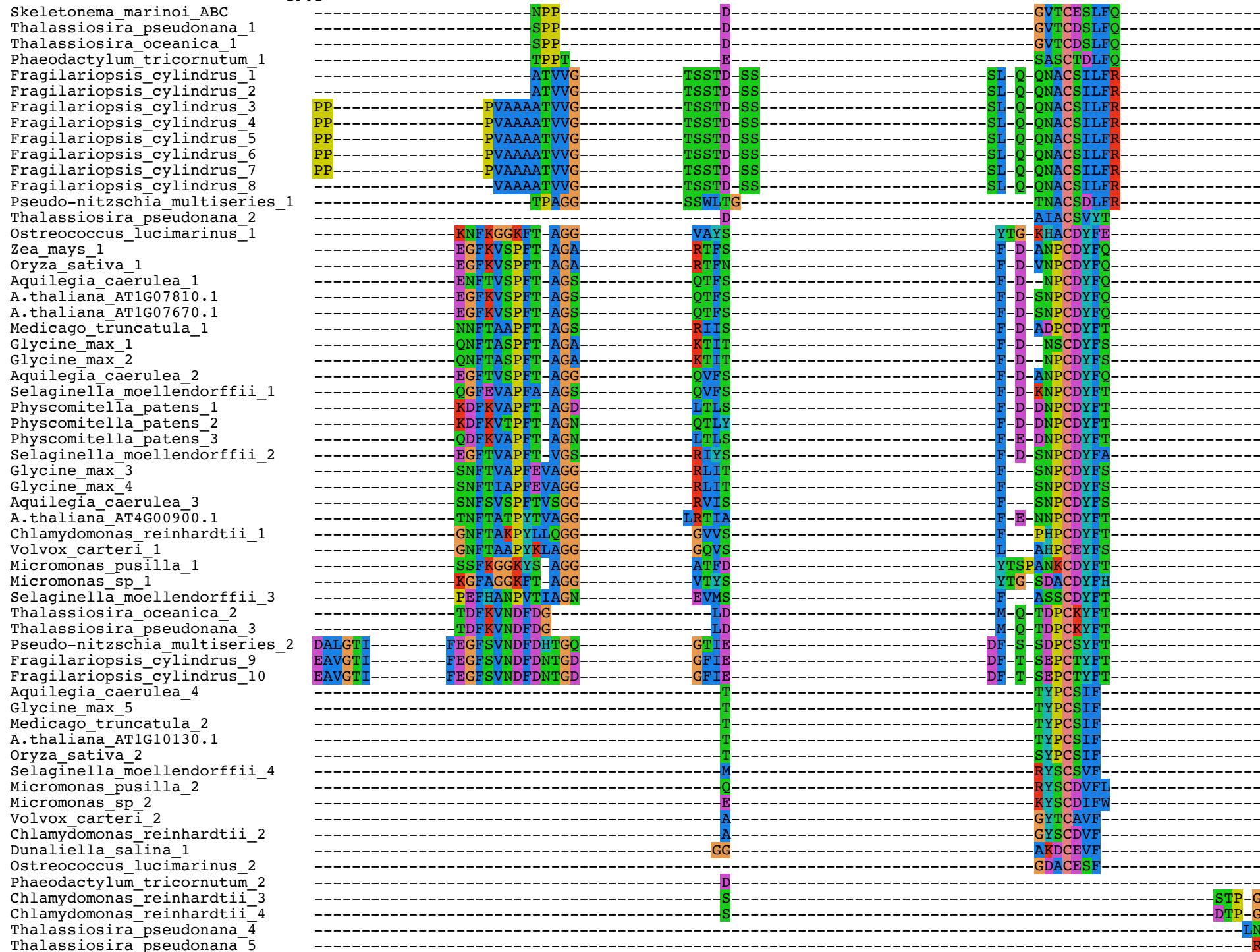


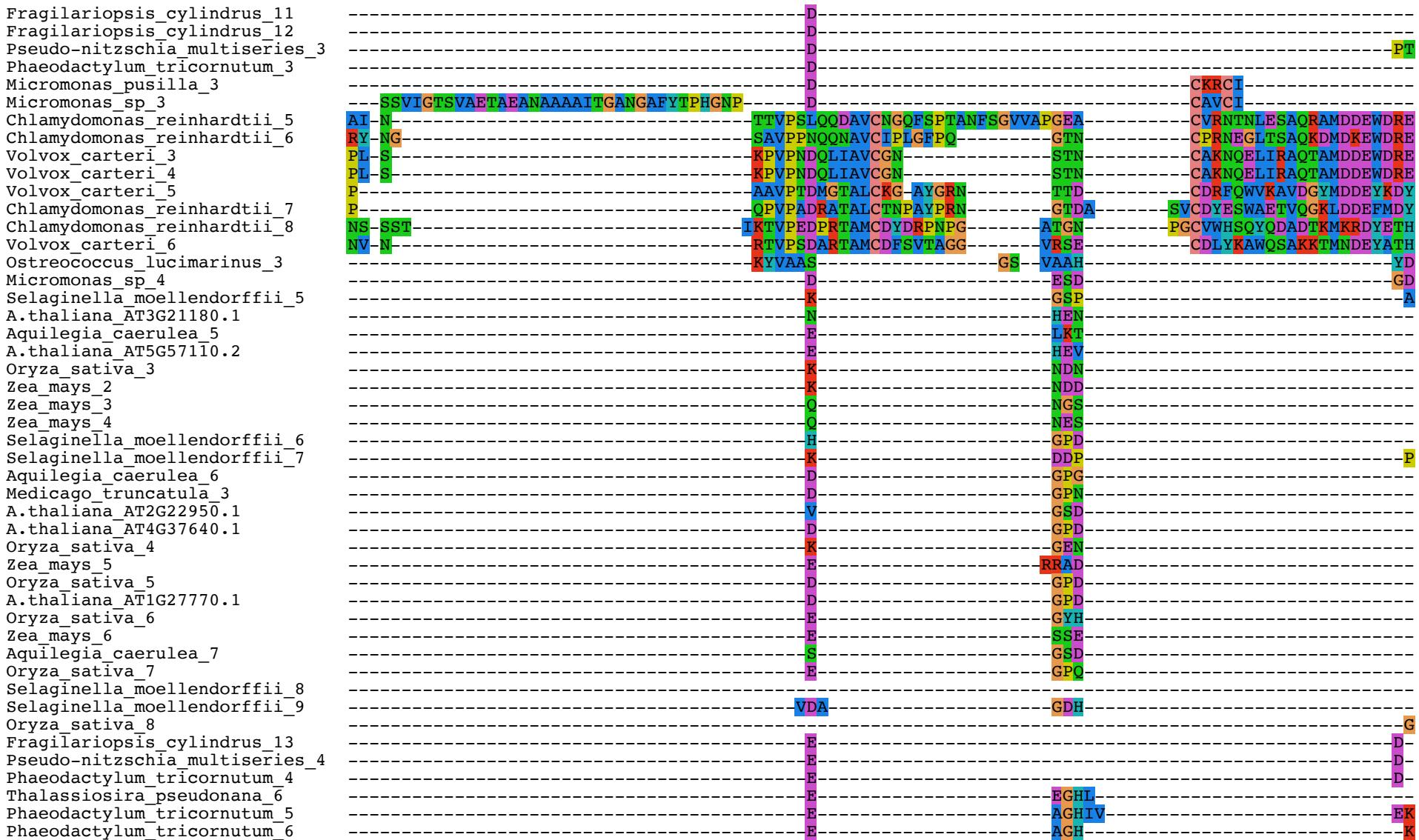
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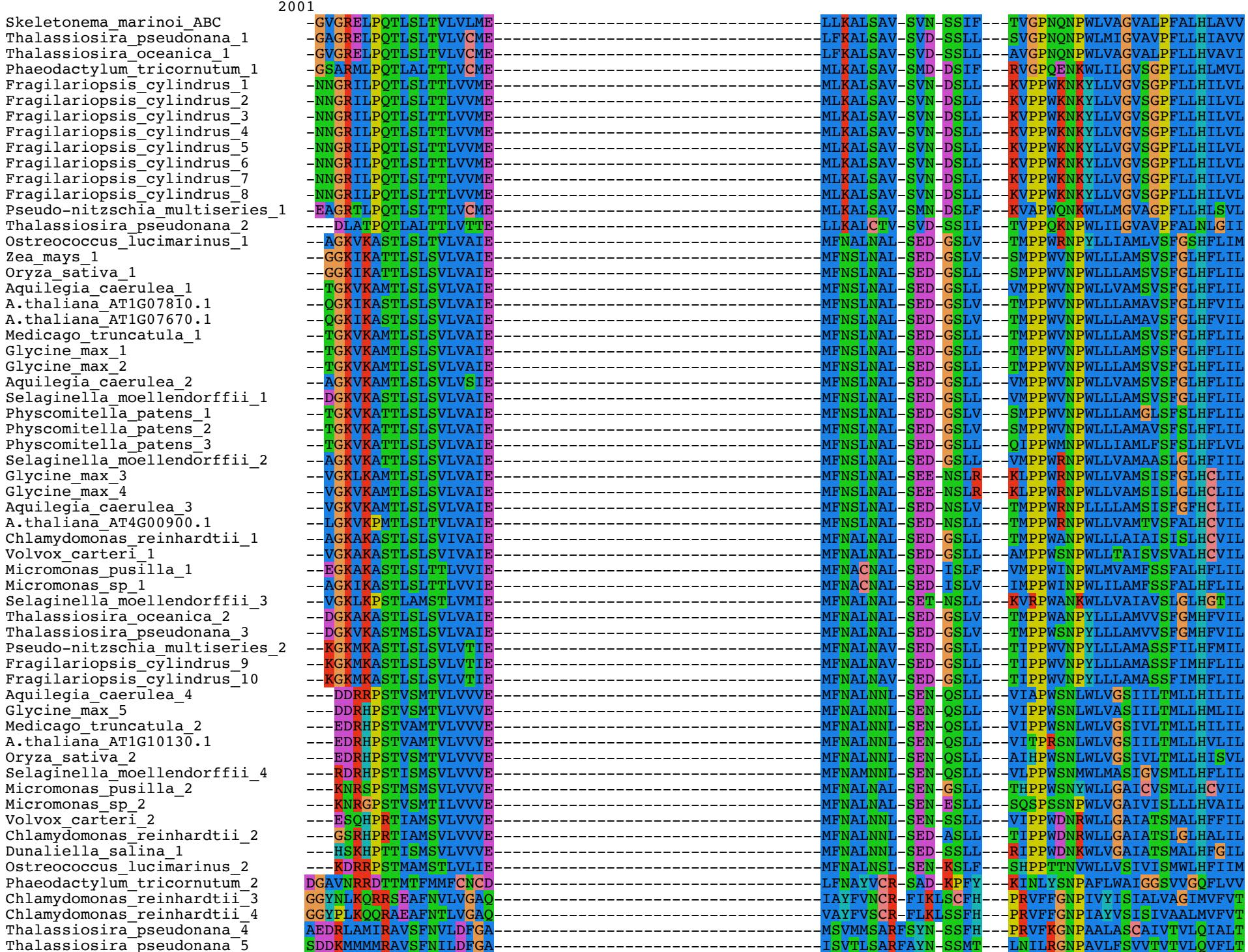
Skeletonema_marinoi_ABC	QGVTLHQL	SSWGKC	NQVW-
Thalassiosira_pseudonana_1	QGLTLRQL	SSWGKC	DQTW-
Thalassiosira_oceanica_1	QGISLKQL	RSGKGC	DQSW-
Phaeodactylum_tricornutum_1	QGITLAAQL	ATWSOC	GFW-
Fragilaropsis_cylindrus_1	QGITLKLQL	SNWSNC	-
Fragilaropsis_cylindrus_2	QGITLKLQL	SNWSNC	-
Fragilaropsis_cylindrus_3	QGITLKLQL	SNWSNC	GTTWM
Fragilaropsis_cylindrus_4	QGITLKLQL	SNWSNC	GTTWM
Fragilaropsis_cylindrus_5	QGITLKLQL	SNWSNC	GTTWM
Fragilaropsis_cylindrus_6	QGITLKLQL	SNWSNC	GTTWM
Fragilaropsis_cylindrus_7	QGITLKLQL	SNWSNC	GTTWM
Fragilaropsis_cylindrus_8	QGITLKLQL	SNWSNC	GTTWM
Pseudo-nitzschia_multiseries_1	QGISLKQL	SSWSSCS	GTDW-
Thalassiosira_pseudonana_2	QGINVQEL	TNWGEC	ASW-
Ostreococcus_lucimarinus_1	GHTPVTFKQL	SNWDKC	SSW-
Zea_mays_1	GHTLVSYSQL	SNWGQC	SSW-
Oryza_sativa_1	GHSLVSYSQL	AHWGQC	SSW-
Aquilegia_caerulea_1	GHTLVTYSQL	AHWGQC	SSW-
A.thaliana_AT1G07810.1	GHSLVSYSQL	AHWGQC	SSW-
A.thaliana_AT1G07670.1	GHTLVTYSQL	ANWGQC	SSW-
Medicago_truncatula_1	GHTLVTYSQL	ANWGQC	SSW-
Glycine_max_1	GHTLVTYTQL	ANWGQC	SSW-
Glycine_max_2	GHTLVTYTQL	ANWGQC	SSW-
Aquilegia_caerulea_2	GHTLVTYSQL	SNWGQC	PSW-
Selaginella_moellendorffii_1	GHTLVTFSQL	TSWGEC	PSW-
Physcomitella_patens_1	GHTLVSFSQL	THWGEC	STW-
Physcomitella_patens_2	GHTLVSFSQL	THWGEC	STW-
Physcomitella_patens_3	GHTLVSFSQL	RNWGDC	PNW-
Selaginella_moellendorffii_2	GHSLVTFSQL	THWGEC	PTW-
Glycine_max_3	GHTIIIELSQL	RNWGEC	PSW-
Glycine_max_4	GHTIIIELSQL	RNWGEC	PSW-
Aquilegia_caerulea_3	GHTLVTLYQL	RTWGQC	PSW-
A.thaliana_AT4G00900.1	GHTLVSFTQL	QNWSEC	SSWG
Chlamydomonas_reinhardtii_1	GHSTVTVWEQL	TNWQSC	REW-
Volvox_carteri_1	GHTTVTVWEQL	TNWQOC	REW-
Micromonas_pusilla_1	GHTTVTWHLQ	SHWGDC	ASWG
Micromonas_sp_1	GHTPVTVHLQ	THWGEC	ETW-
Selaginella_moellendorffii_3	GHTAVSFQQL	SHWGEC	PLW-
Thalassiosira_oceanica_2	DHTNITWEQL	SNWGHC	STW-
Thalassiosira_pseudonana_3	DHTNISWEQL	TGWHHC	STW-
Pseudo-nitzschia_multiseries_2	GHSMVTYGQL	TRWGQC	RTGE
Fragilaropsis_cylindrus_9	GHSMVSYAQL	THWGQC	RTGS
Fragilaropsis_cylindrus_10	GHSMVSYAQL	THWGQC	RTGS
Aquilegia_caerulea_4	GPKLPYYEL	INFDT	PARE
Glycine_max_5	GPKLPYTEL	MNFDT	PTRE
Medicago_truncatula_2	GPKLPYTEL	MNFDT	PTRE
A.thaliana_AT1G10130.1	GPKLTYSSEL	MNFETC	ALRE
Oryza_sativa_2	GPRLPYSSELARKPLLHALLLARKWDKMMWSNNHTSSYQOK	PISLIKEI	PSIGPQVNFDSC
Selaginella_moellendorffii_4	GPRLTWKEI	VSFNEC	KDGT
Micromonas_pusilla_2	GPMLTWSEL	TSFESC	EEGK
Micromonas_sp_2	GPMLTWHLQ	TSFTEC	VEGV
Volvox_carteri_2	GGNLWTSQL	TAFQKCT	EPSAKA
Chlamydomonas_reinhardtii_2	GGGLWTSQL	TSFQKCT	EASAKA
Dunaliella_salina_1	GGNMWTSQL	THFQSC	ASQP
Ostreococcus_lucimarinus_2	GPQMTWAEL	TSASRCI	-
Phaeodactylum_tricornutum_2	NEL		
Chlamydomonas_reinhardtii_3			
Chlamydomonas_reinhardtii_4			
Thalassiosira_pseudonana_4			
Thalassiosira_pseudonana_5			

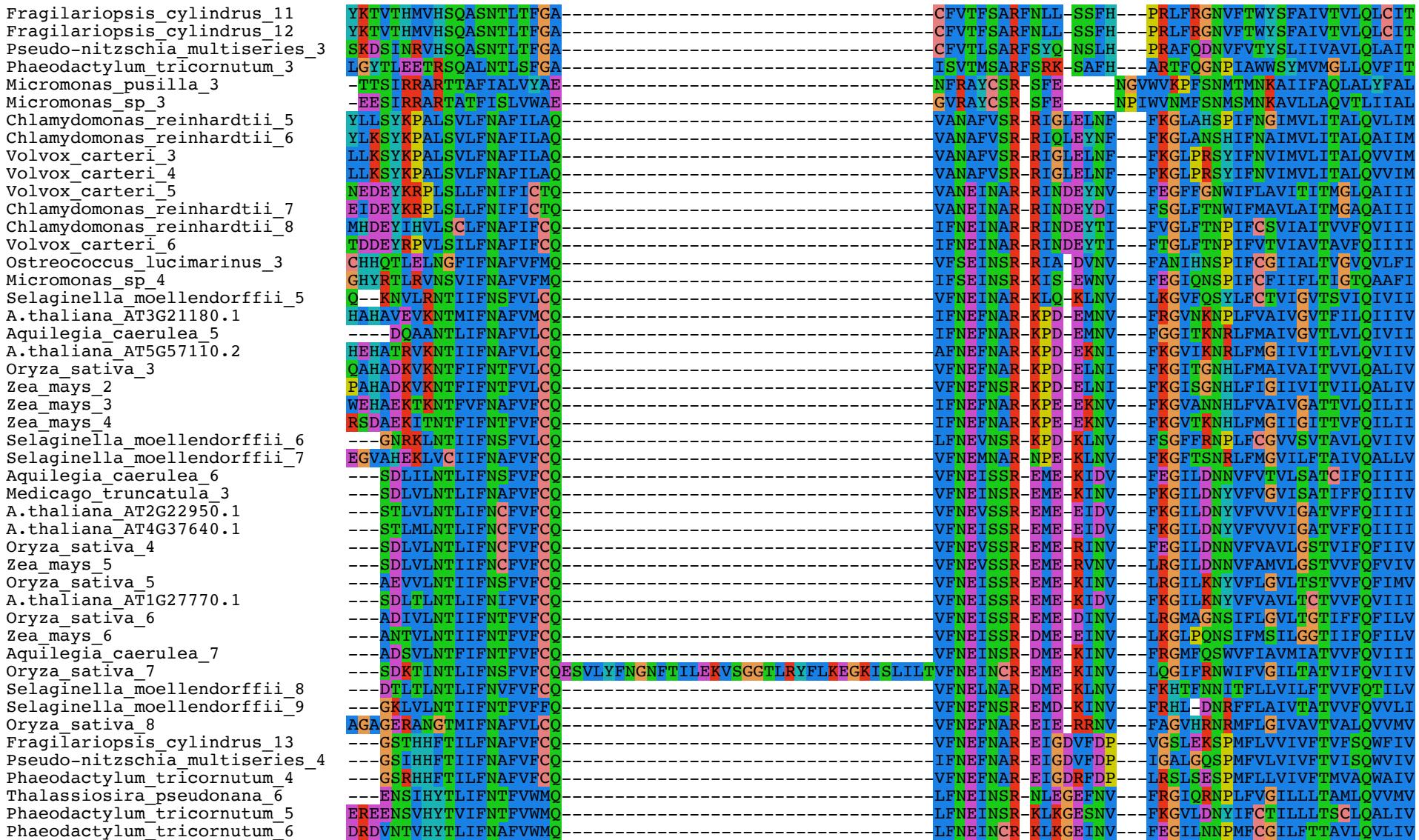
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Pseudo-nitzschia_multiseries_3
Phaeodactylum_tricornutum_3
Micromonas_pusilla_3
Micromonas_sp_3
Chlamydomonas_reinhardtii_5
Chlamydomonas_reinhardtii_6
Volvox_carteri_3
Volvox_carteri_4
Volvox_carteri_5
Chlamydomonas_reinhardtii_7
Chlamydomonas_reinhardtii_8
Volvox_carteri_6
Ostreococcus_lucimarinus_3
Micromonas_sp_4
Selaginella_moellendorffii_5
A.thaliana_AT3G21180.1
Aquilegia_caerulea_5
A.thaliana_AT5G57110.2
Oryza_sativa_3
Zea_mays_2
Zea_mays_3
Zea_mays_4
Selaginella_moellendorffii_6
Selaginella_moellendorffii_7
Aquilegia_caerulea_6
Medicago_truncatula_3
A.thaliana_AT2G22950.1
A.thaliana_AT4G37640.1
Oryza_sativa_4
Zea_mays_5
Oryza_sativa_5
A.thaliana_AT1G27770.1
Oryza_sativa_6
Zea_mays_6
Aquilegia_caerulea_7
Oryza_sativa_7
Selaginella_moellendorffii_8
Selaginella_moellendorffii_9
Oryza_sativa_8
Fragilariopsis_cylindrus_13
Pseudo-nitzschia_multiseries_4
Phaeodactylum_tricornutum_4
Thalassiosira_pseudonana_6
Phaeodactylum_tricornutum_5
Phaeodactylum_tricornutum_6

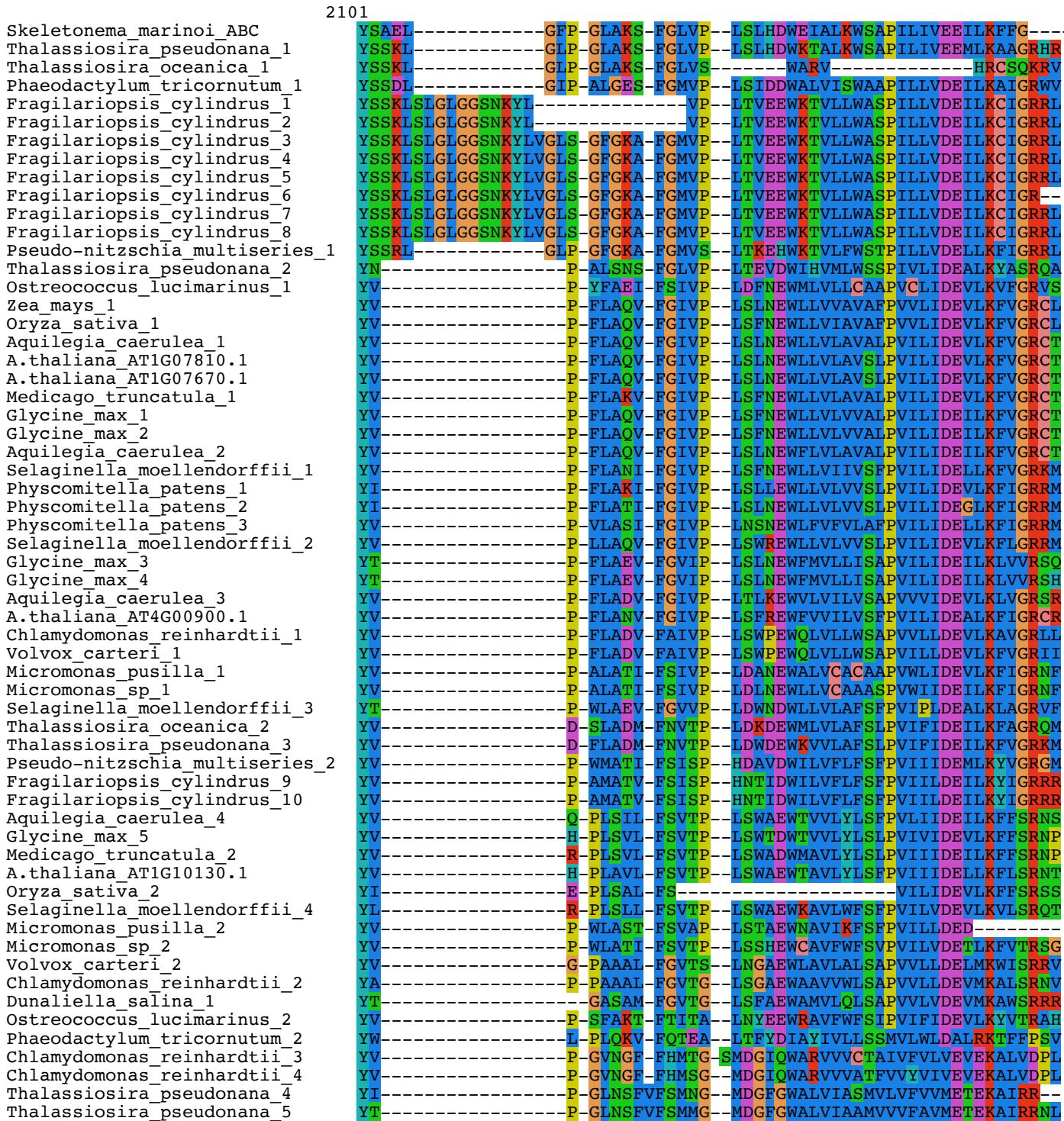




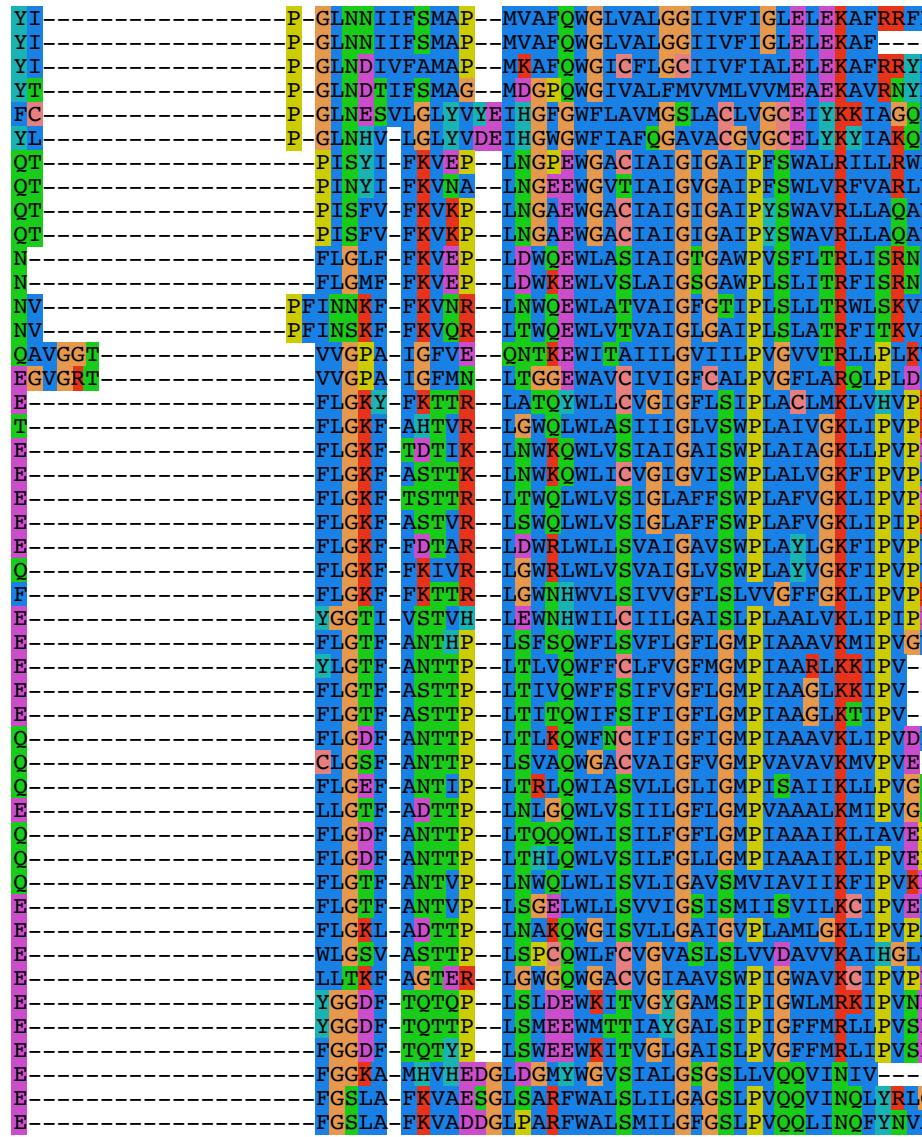








Fragilaropsis_cylindrus_11
 Fragilaropsis_cylindrus_12
 Pseudo-nitzschia_multiseries_3
 Phaeodactylum_tricornutum_3
 Micromonas_pusilla_3
 Micromonas_sp_3
 Chlamydomonas_reinhardtii_5
 Chlamydomonas_reinhardtii_6
 Volvox_carteri_3
 Volvox_carteri_4
 Volvox_carteri_5
 Chlamydomonas_reinhardtii_7
 Chlamydomonas_reinhardtii_8
 Volvox_carteri_6
 Ostreococcus_lucimarinus_3
 Micromonas_sp_4
 Selaginella_moellendorffii_5
 A.thaliana_AT3G21180.1
 Aquilegia_caerulea_5
 A.thaliana_AT5G57110.2
 Oryza_sativa_3
 Zea_mays_2
 Zea_mays_3
 Zea_mays_4
 Selaginella_moellendorffii_6
 Selaginella_moellendorffii_7
 Aquilegia_caerulea_6
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 Fragilaropsis_cylindrus_13
 Pseudo-nitzschia_multiseries_4
 Phaeodactylum_tricornutum_4
 Thalassiosira_pseudonana_6
 Phaeodactylum_tricornutum_5
 Phaeodactylum_tricornutum_6



Supplementary figure 8.

Alignment of the ABC-transporter domains used in the phylogenetic analysis presented in supplementary figure 7. The length of the alignment has been adjusted to the length of the domain identified in *S. marinoi*. Numbers above the sequences indicate the positions in the alignment and the background color of the letters indicates size, shape, solubility, and ionization properties of the different amino acids.