

Table S1. Details of 22 sampling sites in New Zealand between 2018 and 2020.

Climate zone	Island	Sampling site code ^a	Sampling date	Location of sampling site	Latitude, Longitude	Sampling depth (m)
Subtropical	North Island	1	2018 or 2019, and 2020 January 14	Kerikeri, Far North, Northland	-35.1986, 174.0602	0–9 (0+3+6+9)
		2	2018 November 27 and 2020 June 22	Tapeka Point, Russell, Far North, Northland	-35.2451, 174.1179	0
		3	2020 July 20	Bream Bay, Whangārei, Northland	-35.8838, 174.4697	7
		4	2020 July 13	Ōpōtiki, Ōpōtiki, Bay of Plenty	-37.9079, 177.2161	0, 5, 10, and 15
Warm temperate	South Island	5	2018 or 2019, and 2020 June 21	Collingwood Farms, Tasman, Tasman	-40.6550, 172.7172	0–12
		6	2020 June 21	Tākaka, Tasman, Tasman	-40.8162, 172.8126	0–12
		7	2020 June 21	Bomber Port, Motueka, Tasman, Tasman	-41.1193, 173.0447	0–12
		8	2020 June 23 and 30	Oyster Bay, Marlborough, Marlborough	-41.0458, 173.7196	0–15
		9	2020 June 30	Hallam Cove, Marlborough, Marlborough	-40.9958, 173.8246	0–15
		10	2019 April 16 and 2020 July 20	Pukatea Bay, Tasman, Tasman	-40.9564, 173.8967	0–15
		11	2020 June 23	Cannon Hill, Marlborough, Marlborough	-40.9475, 173.9738	0–15
		12	2020 July 07	Brightlands Bay, Marlborough, Marlborough	-41.0612, 173.8544	0–15
		13	2020 January	Richmond Bay, Marlborough, Marlborough	-41.0102, 173.9665	0–15
		14	2020 January and 2020 July 13	Crail Bay, Marlborough, Marlborough	-41.1147, 173.9656	0–15
		15	2019 October and 2020 July 13	Beatrix Bay, Marlborough, Marlborough	-41.0306, 174.0054	0–15
		16	2020 July 13	Laverique Bay, Marlborough, Marlborough	-41.0530, 174.0455	0–15
		17	2019 October	Waitaria Bay, Marlborough, Marlborough	-41.1677, 174.0330	0–15
		18	2020 June 24	Horahora, Marlborough, Marlborough	-41.3307, 174.1360	0–15
		19	2019 May 01	Opihi Bay, Marlborough, Marlborough	-41.2938, 174.1452	0–15
		20	2019 May 01 and 2020 June 24	Whangakoko Bay, Marlborough, Marlborough	-41.2961, 174.1656	0–15
Cold temperate	Stewart Island	21	2020 July 06, 19, and November 22	Big Glory Bay, Southland, Southland	-46.9700, 168.1172	0–10
		22	2019 March 17 and 2020 November 22	Big Glory Bay, Southland, Southland	-46.98, 168.11	0–10

^a Sampling site code shown in Figure 4.

Table S2. Details of 130 clonal strains of *Pseudo-nitzschia* established from New Zealand between 2018 and 2020. NR: sequence determined but not registered as it was identical to the other registered sequences of each species/clade/subclade. ND: not detected. -: sequence not determined or toxin production not tested.

Species	Clade/ subclade	Sequence type (ST) ^a of the ITS region	Strain name in CICCMB ^b	Strain name	Sampling site code ^c	Sampling date	Isolater	DDBJ ^d accession number for the ITS region	DDBJ ^d accession number for the LSU rDNA D1–D3	DA and/or DA isomers production ^e
<i>P. americana</i>				PB024Ps03	3	2020 July 20	T. Nishimura	LC636495	-	ND
<i>P. americana</i>				PB024Ps04	3	2020 July 20	T. Nishimura	NR	-	-
<i>P. americana</i>				PB024Ps09	3	2020 July 20	T. Nishimura	NR	-	-
<i>P. americana</i>				G005Ps02	7	2020 June 21	T. Nishimura	LC636496	-	ND
<i>P. americana</i>			CAWB138	G018Ps01	10	2020 July 20	T. Nishimura	LC636497	LC636556	ND
<i>P. americana</i>				G031Ps01	16	2020 July 13	H. Bowers and T. Nishimura	NR	-	ND
<i>P. arenysensis</i>			CAWB139	PA030Ps01	2	2020 June 22	T. Nishimura	LC636498	-	ND
<i>P. arenysensis</i>				PA030Ps04	2	2020 June 22	T. Nishimura	LC636499	-	ND
<i>P. arenysensis</i>				PA030Ps05	2	2020 June 22	T. Nishimura	NR	-	-
<i>P. arenysensis</i>			CAWB131	Psd-5	10	2019 April 16	M. Balci	LC636500	LC636557	ND
<i>P. arenysensis</i>			CAWB132	Psd-6	10	2019 April 16	M. Balci	LC636501	LC636558	ND
<i>P. arenysensis</i>			CAWB133	Psd-7	10	2019 April 16	M. Balci	-	LC636559	ND
<i>P. arenysensis</i>			CAWB134	Psd-8	10	2019 April 16	M. Balci	-	LC636560	ND
<i>P. australis</i>		ST1		PD009Ps05	4	2020 July 13	T. Nishimura	LC636502	-	Produced
<i>P. australis</i>		ST1	CAWB140	G007Ps03	6	2020 June 21	T. Nishimura	LC636503	LC636561	Produced
<i>P. australis</i>		ST1		G010Ps02	9	2020 June 30	T. Nishimura	NR	-	-
<i>P. australis</i>		ST1		G010Ps03	9	2020 June 30	T. Nishimura	NR	-	-
<i>P. australis</i>		ST1		G027Ps01	12	2020 July 07	T. Nishimura	LC636504	-	Produced
<i>P. australis</i>		ST1		G027Ps02	12	2020 July 07	T. Nishimura	NR	-	-
<i>P. australis</i>		ST1		G015Ps04	14	2020 July 13	H. Bowers and T. Nishimura	LC636505	LC636562	Produced
<i>P. australis</i>		ST1		G037Ps01	16	2020 July 13	H. Bowers and T. Nishimura	LC636506	-	Produced
<i>P. australis</i>		ST1		G037Ps02	16	2020 July 13	H. Bowers and T. Nishimura	NR	-	-
<i>P. australis</i>		ST1		J328Ps20	21	2020 November 22	T. Nishimura	LC636507	-	Produced
<i>P. australis</i>		ST1		J237Ps07	22	2020 November 22	T. Nishimura	LC636508	-	Produced
<i>P. australis</i>		ST2		G018Ps02	10	2020 July 20	T. Nishimura	LC636509	LC636563	Produced
<i>P. australis</i>		ST2		G018Ps03	10	2020 July 20	T. Nishimura	NR	-	-
<i>P. australis</i>		ST2		G018Ps04	10	2020 July 20	T. Nishimura	LC636510	-	Produced
<i>P. calliantha</i>		ST1	CAWB135	Crail Bay	14	2020 January	L. L. Rhodes	LC636511	LC636564	ND
<i>P. calliantha</i>		ST2		Waitaria Bay	17	2019 October	S. Challenger	LC636512	LC636565	ND
<i>P. calliantha</i>		ST3		Richmond Bay	13	2020 January	L. L. Rhodes	LC636513	LC636566	ND
<i>P. cuspidata</i>	clade Ia		CAWB141	PA030Ps10	2	2020 June 22	T. Nishimura	LC636514	LC636567	Produced (trace)

<i>P. delicatissima</i>	subclade I	CAWB136	PA014Ps04	1	2020 January 14	T. Nishimura	LC636515	LC636568	ND
<i>P. delicatissima</i>	subclade I		PA014Ps05	1	2020 January 14	T. Nishimura	LC636516	LC636569	ND
<i>P. delicatissima</i>	subclade I		PA014Ps11	1	2020 January 14	T. Nishimura	LC636517	LC636570	ND
<i>P. delicatissima</i>	subclade I	CAWB142	PA030Ps06	2	2020 June 22	T. Nishimura	LC636518	-	ND
<i>P. delicatissima</i>	subclade I		PA030Ps11	2	2020 June 22	T. Nishimura	LC636519	-	ND
<i>P. delicatissima</i>	subclade II	CAWB143	G013Ps03	8	2020 June 23	T. Nishimura	LC636520	LC636571	ND
<i>P. delicatissima</i>	subclade II		G013Ps04	8	2020 June 23	T. Nishimura	LC636521	-	ND
<i>P. delicatissima</i>	subclade II		G013Ps05	8	2020 June 23	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps08	8	2020 June 23	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps10	8	2020 June 23	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps12	8	2020 June 23	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps13	8	2020 June 30	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps14	8	2020 June 30	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps15	8	2020 June 30	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		G013Ps16	8	2020 June 30	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II	CAWB144	J328Ps01	21	2020 July 06	H. Bowers and T. Nishimura	LC636522	LC636572	ND
<i>P. delicatissima</i>	subclade II		J328Ps02	21	2020 July 06	H. Bowers and T. Nishimura	LC636523	-	ND
<i>P. delicatissima</i>	subclade II		J328Ps03	21	2020 July 06	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps04	21	2020 July 06	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps05	21	2020 July 06	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps06	21	2020 July 06	H. Bowers and T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps07	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps08	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps09	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps10	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps11	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps14	21	2020 November 22	T. Nishimura	NR	-	-
<i>P. delicatissima</i>	subclade II		J328Ps17	21	2020 November 22	T. Nishimura	NR	-	-
<i>P. fraudulent</i>			828	1	2018 or 2019	S. Challenger	-	LC636573	-
<i>P. fraudulent</i>			G013Ps02	8	2020 June 23	T. Nishimura	NR	-	ND
<i>P. fraudulent</i>			G013Ps09	8	2020 June 23	T. Nishimura	NR	-	-
<i>P. fraudulent</i>		CAWB145	G015Ps01	14	2020 July 13	H. Bowers	LC636524	LC636574	ND
<i>P. fraudulent</i>			G015Ps03	14	2020 July 13	H. Bowers	NR	-	-
<i>P. fraudulent</i>			G011Ps04	20	2020 June 24	T. Nishimura	NR	-	ND
<i>P. fraudulent</i>			J328Ps12	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. fraudulent</i>			J328Ps13	21	2020 July 19	T. Nishimura	NR	-	-
<i>P. fraudulent</i>			J328Ps16	21	2020 November 22	T. Nishimura	LC636525	-	ND

<i>P. fraudulenta</i>			J328Ps18	21	2020 November 22	T. Nishimura	NR	-	-	
<i>P. fraudulenta</i>			J328Ps19	21	2020 November 22	T. Nishimura	NR	-	-	
<i>P. fraudulenta</i>		CAWB128	Kiwa1	22	2019 March 17	S. Challenger	-	LC636575	ND	
<i>P. fraudulenta</i>			Kiwa2	22	2019 March 17	S. Challenger	-	LC636576	-	
<i>P. galaxiae</i>		CAWB153	PA014Ps01	1	2020 January 14	T. Nishimura	LC636526	LC636577	ND	
<i>P. galaxiae</i>		CAWB154	PA014Ps07	1	2020 January 14	T. Nishimura	LC636527	LC636578	ND	
<i>P. hasleana</i>	clade I	CAWB146	G007Ps01	6	2020 June 21	T. Nishimura	LC636528	LC636579	ND	
<i>P. hasleana</i>	clade I	CAWB147	G007Ps02	6	2020 June 21	T. Nishimura	LC636529	-	ND	
<i>P. hasleana</i>	clade II	CAWB156	J328Ps15	21	2020 November 22	T. Nishimura	LC636530	LC636580	ND	
<i>P. hasleana</i>	clade II		J237Ps01	22	2020 November 22	T. Nishimura	LC636531	-	-	
<i>P. hasleana</i>	clade II		J237Ps02	22	2020 November 22	T. Nishimura	LC636532	-	-	
<i>P. hasleana</i>	clade II	CAWB155	J237Ps05	22	2020 November 22	T. Nishimura	LC636533	LC636581	ND	
<i>P. multiseriis</i>		CAWB149	G001Ps03	5	2020 June 21	T. Nishimura	LC636534	LC636582	Produced	
<i>P. multiseriis</i>			G001Ps04	5	2020 June 21	T. Nishimura	LC636535	-	Produced	
<i>P. multiseriis</i>			G007Ps04	6	2020 June 21	T. Nishimura	LC636536	-	Produced	
<i>P. multiseriis</i>			G010Ps10	9	2020 June 30	T. Nishimura	LC636537	-	Produced	
<i>P. multistriata</i>	ST1	CAWB137	PA014Ps09	1	2020 January 14	T. Nishimura	LC636538	LC636583	ND	
<i>P. multistriata</i>	ST1		PA014Ps10	1	2020 January 14	T. Nishimura	LC636539	LC636584	ND	
<i>P. multistriata</i>	ST1		G013Ps21	8	2020 June 30	T. Nishimura	LC636540	-	ND	
<i>P. multistriata</i>	ST1		G013Ps22	8	2020 June 30	T. Nishimura	NR	-	-	
<i>P. multistriata</i>	ST1		G027Ps06	12	2020 July 07	T. Nishimura	LC636541	-	Produced (trace)	
<i>P. multistriata</i>	ST2	CAWB127	827	2	2018 November 27	S. Challenger	LC636542	LC636585	ND	
<i>P. multistriata</i>	ST2	CAWB150	PB024Ps01	3	2020 July 20	T. Nishimura	LC636543	-	ND	
<i>P. multistriata</i>	ST2		PB024Ps02	3	2020 July 20	T. Nishimura	NR	-	-	
<i>P. multistriata</i>	ST2		PB024Ps05	3	2020 July 20	T. Nishimura	NR	-	-	
<i>P. multistriata</i>	ST2		PB024Ps07	3	2020 July 20	T. Nishimura	NR	-	-	
<i>P. multistriata</i>	ST2		PB024Ps08	3	2020 July 20	T. Nishimura	NR	-	-	
<i>P. multistriata</i>	ST2		G001Ps01	5	2020 June 21	T. Nishimura	LC636544	-	Produced (trace)	
<i>P. multistriata</i>	ST2		G001Ps02	5	2020 June 21	T. Nishimura	LC636545	LC636586	ND	
<i>P. plurisecta</i>		CAWB151	G027Ps03	12	2020 July 07	T. Nishimura	LC636546	LC636587	ND	
<i>P. pungens</i>	clade I	ST1	CAWB152	PA030Ps03	2	2020 June 22	T. Nishimura	LC636547	LC636588	ND
<i>P. pungens</i>	clade I	ST1		PA030Ps08	2	2020 June 22	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1		G007Ps06	6	2020 June 21	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1		G005Ps01	7	2020 June 21	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1		G005Ps04	7	2020 June 21	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1		G010Ps01	9	2020 June 30	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1		G010Ps04	9	2020 June 30	T. Nishimura	NR	-	-

<i>P. pungens</i>	clade I	ST1	G010Ps06	9	2020 June 30	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G010Ps07	9	2020 June 30	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G010Ps08	9	2020 June 30	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G018Ps05	10	2020 July 20	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1	Beatrix Bay	15	2019 October	S. Challenger	LC636548	-	ND
<i>P. pungens</i>	clade I	ST1	G011Ps01	20	2020 June 24	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1	G011Ps02	20	2020 June 24	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G011Ps05	20	2020 June 24	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G011Ps07	20	2020 June 24	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G011Ps08	20	2020 June 24	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	G011Ps09	20	2020 June 24	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	J328Ps21	21	2020 November 22	T. Nishimura	LC636549	-	ND
<i>P. pungens</i>	clade I	ST1	J237Ps03	22	2020 November 22	T. Nishimura	NR	-	ND
<i>P. pungens</i>	clade I	ST1	J237Ps04	22	2020 November 22	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	J237Ps06	22	2020 November 22	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST1	J237Ps08	22	2020 November 22	T. Nishimura	NR	-	-
<i>P. pungens</i>	clade I	ST2	G027Ps05	12	2020 July 07	T. Nishimura	LC636550	-	ND
<i>P. pungens</i>	clade I	ST2	G015Ps02	14	2020 July 13	H. Bowers	LC636551	-	ND
<i>P. pungens</i>	clade I	-	CAWB130	Psd-4	19	2019 May 01	M. Balci	-	LC636589
<i>P. pungens</i>	clade I	-	CAWB129	Psd-3	20	2019 May 01	M. Balci	-	LC636590
<i>P. cf. subpaci</i>			PA030Ps02	2	2020 June 22	T. Nishimura	LC636552	-	ND
<i>P. cf. subpaci</i>			PA030Ps07	2	2020 June 22	T. Nishimura	LC636553	LC636591	ND
<i>P. cf. subpaci</i>		CAWB148	PD009Ps04	4	2020 July 13	T. Nishimura	LC636554	LC636592	ND
<i>P. cf. subpaci</i>			826	5	2018 or 2019	S. Challenger	-	LC636593	-
<i>P. cf. subpaci</i>			G011Ps03	20	2020 June 24	T. Nishimura	LC636555	-	ND

^a Sequences that were not identical but were not a clade/subclade level divergence were assigned as sequence type (ST).

^b CICC: the Cawthron Institute Culture Collection of Microalgae.

^c Sampling site code shown in Figure 4.

^d DDBJ: the DNA Data Bank of Japan.

^e Details of the DA and DA isomers analysis are shown in Table S5.

Table S3. Average, SD, minimum, and maximum uncorrected *p* distance of the ITS region of selected combinations between *Pseudo-nitzschia* species/clades/subclades. NA: data not available.

Species/clade/subclade	Species/clade/subclade	Number of positions ^a	Number of combinations	<i>p</i> distance			
				Average	SD	Minimum	Maximum
<i>P. arenysensis</i> clade I	<i>P. arenysensis</i> clade I	690	36	0	0	0	0
	<i>P. arenysensis</i> clade II	690	18	0.045	0.005	0.040	0.052
	<i>P. cf. arenysensis</i>	690	27	0.040	0.000	0.040	0.040
<i>P. arenysensis</i> clade II	<i>P. arenysensis</i> clade II	690	1	NA	NA	NA	0.013
	<i>P. cf. arenysensis</i>	690	6	0.041	0.000	0.040	0.041
<i>P. cf. arenysensis</i>	<i>P. cf. arenysensis</i>	690	3	0	0	0	0
<i>P. cuspidata</i> clade I ^b	<i>P. cuspidata</i> clade I ^b	724	21	0.009	0.006	0	0.021
	<i>P. cuspidata</i> clade II	724	35	0.030	0.004	0.024	0.037
	<i>P. pseudodelicatissima</i> ^c	724	63	0.031	0.003	0.026	0.037
	<i>P. plurisecta</i>	724	35	0.031	0.002	0.026	0.034
<i>P. cuspidata</i> clade Ia	<i>P. cuspidata</i> clade Ia	724	3	0.001	0.001	0	0.001
	<i>P. cuspidata</i> clade Ib	724	12	0.013	0.004	0.010	0.021
<i>P. cuspidata</i> clade Ib	<i>P. cuspidata</i> clade Ib	724	6	0.006	0.003	0	0.010
<i>P. cuspidata</i> clade II	<i>P. cuspidata</i> clade II	724	10	0.012	0.008	0.003	0.024
	<i>P. pseudodelicatissima</i> ^c	724	45	0.021	0.005	0.015	0.031
	<i>P. plurisecta</i>	724	25	0.036	0.003	0.030	0.041
<i>P. pseudodelicatissima</i> ^c	<i>P. pseudodelicatissima</i> ^c	724	36	0.004	0.003	0	0.009
	<i>P. plurisecta</i>	724	45	0.034	0.004	0.027	0.039
<i>P. plurisecta</i>	<i>P. plurisecta</i>	724	10	0.004	0.002	0	0.007
<i>P. decipiens</i> clade A	<i>P. decipiens</i> clade A	644	1	NA	NA	NA	0.005
	<i>P. decipiens</i> clade B	644	4	0.027	0.002	0.023	0.030
<i>P. decipiens</i> clade B	<i>P. decipiens</i> clade B	644	1	NA	NA	NA	0.014
<i>P. delicatissima</i> subclade I	<i>P. delicatissima</i> subclade I	671	45	0.002	0.002	0	0.005
	<i>P. delicatissima</i> subclade II	671	80	0.042	0.001	0.041	0.046
	<i>P. delicatissima</i> subclade III	671	60	0.044	0.001	0.043	0.046
<i>P. delicatissima</i> subclade II	<i>P. delicatissima</i> subclade II	671	28	0.001	0.001	0	0.003
	<i>P. delicatissima</i> subclade III	671	48	0.033	0.001	0.031	0.036
<i>P. delicatissima</i> subclade III	<i>P. delicatissima</i> subclade III	671	15	0.001	0.001	0	0.003
Four New Zealand strains of <i>P. delicatissima</i> subclade II	Four other strains of <i>P. delicatissima</i> subclade II	671	16	0.002	0.001	0.002	0.003
<i>P. galaxiae</i> clade A ^d	<i>P. galaxiae</i> clade A ^d	593	1	NA	NA	NA	0.012
	<i>P. galaxiae</i> clade B ^d	593	12	0.047	0.002	0.045	0.050
	<i>P. galaxiae</i> clade C	593	6	0.068	0.004	0.064	0.072
<i>P. galaxiae</i> clade B ^d	<i>P. galaxiae</i> clade B ^d	593	15	0.010	0.009	0	0.024
	<i>P. galaxiae</i> clade C	593	18	0.056	0.005	0.053	0.067
<i>P. galaxiae</i> clade C	<i>P. galaxiae</i> clade C	593	3	0	0	0	0
<i>P. hasleana</i> clade I	<i>P. hasleana</i> clade I	620	36	0.002	0.001	0	0.005
	<i>P. hasleana</i> clade II	620	36	0.012	0.001	0.011	0.015
<i>P. hasleana</i> clade II	<i>P. hasleana</i> clade II	620	6	0	0	0	0
<i>P. pungens</i> clade I	<i>P. pungens</i> clade I	624	28	0.001	0.001	0	0.003
	<i>P. pungens</i> clade II	624	24	0.012	0.001	0.011	0.014
	<i>P. pungens</i> clade III ^e	624	72	0.033	0.005	0.026	0.047
<i>P. pungens</i> clade II	<i>P. pungens</i> clade II	624	3	0.001	0.001	0	0.002
	<i>P. pungens</i> clade III ^e	624	27	0.035	0.005	0.027	0.048
<i>P. pungens</i> clade III ^e	<i>P. pungens</i> clade III ^e	624	36	0.014	0.006	0	0.027
<i>P. pungens</i> clade IIIaa	<i>P. pungens</i> clade IIIaa	624	3	0.006	0.001	0.005	0.008
	<i>P. pungens</i> clade IIIab	624	12	0.015	0.005	0.011	0.024
	<i>P. pungens</i> clade IIIb	624	6	0.013	0.001	0.011	0.014
<i>P. pungens</i> clade IIIab	<i>P. pungens</i> clade IIIab	624	6	0.010	0.005	0.005	0.016
	<i>P. pungens</i> clade IIIb	624	8	0.019	0.005	0.016	0.027
<i>P. pungens</i> clade IIIb	<i>P. pungens</i> clade IIIb	624	1	NA	NA	NA	0
<i>P. simulans</i> subclade I	<i>P. simulans</i> subclade I	703	1	NA	NA	NA	0
	<i>P. simulans</i> subclade II	703	4	0.039	0.000	0.039	0.039
<i>P. simulans</i> subclade II	<i>P. simulans</i> subclade II	703	1	NA	NA	NA	0.001

^a Number of positions used for the final data set for the calculation of uncorrected *p* distances.^b *Pseudo-nitzschia cuspidata* clade I contains clades Ia and Ib. Clade Ib includes two *P. cuspidata* var. *manzanillensis* strains (Ps116 and Ps142) and two *P. cuspidata* strains (MC3041 and MC3207).^c *Pseudo-nitzschia pseudodelicatissima* includes two *P. cuspidata* strains (Tenerife8 and NWFSC194).^d *Pseudo-nitzschia galaxiae* clades A and B correspond to clades I and II reported by McDonald et al. (2007), respectively.^e *Pseudo-nitzschia pungens* clade III contains clades IIIaa, IIIab, and IIIb.

Table S4. Average, SD, minimum, and maximum uncorrected *p* distance of the LSU rDNA D1–D3 of selected combinations between *Pseudo-nitzschia hasleana* and several related species. NA: data not available.

Species/subclade	Species/subclade	Number of positions ^a	Number of combinations	<i>p</i> distance			
				Average	SD	Minimum	Maximum
<i>P. hasleana</i> clade I	<i>P. hasleana</i> clade I	659	10	0	0	0	0
	<i>P. hasleana</i> clade II	659	10	0.006	0	0.006	0.006
	' <i>P. calliantha</i> ' ^b	659	5	0.014	< 0.001	0.014	0.014
	<i>P. calliantha</i>	659	25	0.018	0.001	0.015	0.020
	<i>P. mannii</i>	659	10	0.025	0.001	0.024	0.026
	<i>P. limii</i>	659	10	0.007	0.001	0.006	0.008
	<i>P. kodamae</i>	659	10	0.018	0	0.018	0.018
<i>P. hasleana</i> clade II	<i>P. hasleana</i> clade II	659	1	NA	NA	NA	0
	' <i>P. calliantha</i> ' ^b	659	2	0.008	0	0.008	0.008
	<i>P. calliantha</i>	659	10	0.019	0.001	0.017	0.020
	<i>P. mannii</i>	659	4	0.027	0.001	0.026	0.027
	<i>P. limii</i>	659	4	0.008	0.001	0.008	0.009
	<i>P. kodamae</i>	659	4	0.021	0	0.021	0.021
' <i>P. calliantha</i> ' ^b	' <i>P. calliantha</i> ' ^b	659	NA	NA	NA	NA	NA
	<i>P. calliantha</i>	659	5	0.027	0.001	0.025	0.028
	<i>P. mannii</i>	659	2	0.035	0.001	0.034	0.035
	<i>P. limii</i>	659	2	0.016	0.001	0.015	0.017
	<i>P. kodamae</i>	659	2	0.027	< 0.001	0.027	0.027
<i>P. calliantha</i>	<i>P. calliantha</i>	659	10	0.001	0.001	0	0.003
	<i>P. mannii</i>	659	10	0.016	0.001	0.014	0.018
	<i>P. limii</i>	659	10	0.014	0.002	0.011	0.017
	<i>P. kodamae</i>	659	10	0.027	0.001	0.024	0.029
<i>P. mannii</i>	<i>P. mannii</i>	659	1	NA	NA	NA	0.002
	<i>P. limii</i>	659	4	0.018	0.001	0.017	0.020
	<i>P. kodamae</i>	659	4	0.014	0.001	0.014	0.015
<i>P. limii</i>	<i>P. limii</i>	659	1	NA	NA	NA	0.002
	<i>P. kodamae</i>	659	4	0.014	0.001	0.014	0.015
<i>P. kodamae</i>	<i>P. kodamae</i>	659	1	NA	NA	NA	0.003

^a Number of positions used for the final data set for the calculation of uncorrected *p* distances.^b '*Pseudo-nitzschia calliantha*' indicates strain CAWB114 from New Zealand (Rhodes et al. 2013).

Table S5. Details of 73 clonal strains of *Pseudo-nitzschia* from New Zealand tested for domoic acid (DA) and DA isomers analysis in the present study. 0: no toxins were detected at the limit of detection (LoD). trace: toxin cell quota was detected between LoD and lower limit of quantitation (LLOQ). NA: data not available. NC: data not calculated. Bold letter: detected or quantified value.

Species	Clade/ subclade	Sequence type (ST) ^a of the ITS	Strain name in CICCM ^b	Strain name	Medium	Harvested cell number	Number of		LoD	LLOQ	Cell quota (pg cell ⁻¹)								Proportion (%)							
							Day	months after isolation ^c			DA	epi-DA	iso-DA A	iso-DA B	iso-DA C	iso-DA D	iso-DA E	DA	epi-DA	iso-DA A	iso-DA B	iso-DA C	iso-DA D	iso-DA E		
<i>P. americana</i>				PB024Ps03	f/2-Si	9,653,333	9	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. americana</i>				G005Ps02	f/2-Si	7,056,000	10	8	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. americana</i>			CAWB138	G018Ps01	f/2-Si	8,576,000	9	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. americana</i>				G031Ps01	f/2-Si	8,224,000	10	7	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I		CAWB139	PA030Ps01	f/2-Si	13,813,333	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	2,288,000	43	5	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I			PA030Ps04	f/2-Si	9,226,667	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	1,981,333	43	5	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I		CAWB131	Psd-5	f/2	16,350,000	10	14	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I		CAWB132	Psd-6	f/2	37,050,000	10	14	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	18,144,000	10	22	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I		CAWB133	Psd-7	f/2	55,500,000	10	14	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. arenysensis</i>	clade I		CAWB134	Psd-8	f/2	20,550,000	10	14	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. australis</i>		ST1		PD009Ps05	f/2-Si	1,260,000	9	3	0.00008	0.0008	0.27	0	0.009	trace	0.119	0	0	68.1	NA	2.3	NC	29.7	NA	NA	NA	
<i>P. australis</i>		ST1	CAWB140	G007Ps03	f/2-Si	420,000	9	4	0.0002	0.002	0.06	0	0.003	trace	0.021	0	0	72.6	NA	3.0	NC	24.4	NA	NA	NA	
<i>P. australis</i>		ST1		G027Ps01	f/2-Si	573,333	9	3	0.0002	0.002	0.73	0	0.024	trace	0.884	0	0	44.4	NA	1.5	NC	54.1	NA	NA	NA	
<i>P. australis</i>		ST1		G015Ps04	f/2-Si	304,000	9	3	0.0003	0.003	0.05	0	0	trace	0.010	0	0	83.2	NA	NA	NC	16.8	NA	NA	NA	
					f/2-Si	1,506,667	10	7	0.00007	0.0007	0.004	0	trace	trace	0.002	0	0	67.1	NA	NC	NC	32.9	NA	NA	NA	
<i>P. australis</i>	ST1			G037Ps01	f/2-Si	966,667	9	3	0.0001	0.001	0.09	0	0.003	trace	0.051	0	0	62.1	NA	2.2	NC	35.8	NA	NA	NA	
<i>P. australis</i>	ST1			J328Ps20	f/2-Si	580,000	10	3	0.0002	0.002	0.14	trace	0.006	trace	0.007	trace	trace	91.5	NC	3.7	NC	4.8	NC	NC	NC	
<i>P. australis</i>	ST1			J237Ps07	f/2-Si	372,000	10	3	0.0003	0.003	2.02	0.010	0.060	trace	0.118	0.003	trace	91.3	0.4	2.7	NC	5.4	0.1	NC	NC	
<i>P. australis</i>	ST2			G018Ps02	f/2-Si	1,173,333	9	3	0.00009	0.0009	0.04	0	0.001	trace	0.006	0	0	83.8	NA	2.7	NC	13.5	NA	NA	NA	
<i>P. australis</i>	ST2			G018Ps04	f/2-Si	392,000	10	7	0.0003	0.003	0.11	0	0.003	trace	0.050	trace	0	67.9	NA	2.0	NC	30.1	NC	NA	NA	
<i>P. calliantha</i>	ST1		CAWB135	Crail Bay	f/2	41,400,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	6,240,000	9	9	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. calliantha</i>	ST2			Waitaria Bay	f/2	39,300,000	10	8	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	11,386,667	9	12	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. calliantha</i>	ST3			Richmond Bay	f/2	40,800,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	13,216,000	10	13	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. cuspidata</i>	clade Ia		CAWB141	PA030Ps10	f/2-Si	12,026,667	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	1,317,333	43	5	0.00008	0.0008	trace	0	0	0	0	0	0	0	NC	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade I		CAWB136	PA014Ps04	f/2	120,900,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	29,333,333	9	9	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade I			PA014Ps05	f/2	52,125,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	4,213,333	9	9	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade I			PA014Ps11	f/2	102,000,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade I		CAWB142	PA030Ps06	f/2-Si	5,093,333	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade I			PA030Ps11	f/2-Si	13,066,667	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade II		CAWB143	G013Ps03	f/2-Si	12,880,000	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	853,333	43	5	0.0001	0.001	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade II			G013Ps04	f/2-Si	17,386,667	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	829,333	43	5	0.0001	0.001	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
<i>P. delicatissima</i>	subclade II		CAWB144	J328Ps01	f/2-Si	10,666,667	9	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	
					f/2-Si	3,280,000	43	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA	

<i>P. delicatissima</i>	subclade II	J328Ps02	f/2-Si	15,680,000	9	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	3,440,000	43	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. fraudulenta</i>		G013Ps02	f/2-Si	1,328,000	10	8	0.00008	0.0008	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. fraudulenta</i>	CAWB145	G015Ps01	f/2-Si	1,306,667	9	3	0.00008	0.0008	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. fraudulenta</i>		G011Ps04	f/2-Si	4,640,000	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. fraudulenta</i>		J328Ps16	f/2-Si	1,200,000	10	3	0.00008	0.0008	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. fraudulenta</i>	CAWB128	Kiwa1	f/2	82,200,000	10	15	0.0005	0.0025	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. galaxiae</i>	clade B	CAWB153	PA014Ps01	f/2	121,500,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	70,666,667	9	9	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	4,560,000	43	10	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. galaxiae</i>	clade B	CAWB154	PA014Ps07	f/2	213,750,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	39,733,333	9	9	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	3,040,000	43	10	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. hasleana</i>	clade I	CAWB146	G007Ps01	f/2-Si	426,667	9	4	0.0002	0.002	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	304,000	43	5	0.0003	0.003	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. hasleana</i>	clade I	CAWB147	G007Ps02	f/2-Si	1,866,667	9	4	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	955,200	43	5	0.0001	0.001	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. hasleana</i>	clade II	CAWB156	J328Ps15	f/2-Si	8,026,667	10	3	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	6,032,000	13	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	6,336,000	43	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			K	23,386,667	12	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. hasleana</i>	clade II	CAWB155	J237Ps05	f/2-Si	11,792,000	10	3	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	3,008,000	13	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			K	8,016,000	12	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multiseris</i>		CAWB149	G001Ps03	f/2-Si	2,896,000	9	4	0.00005	0.0005	1.65	0.012	0.045	0	0	0.020	0.003	95.4	0.7	2.6	NA	NA	1.2	0.2
<i>P. multiseris</i>			G001Ps04	f/2-Si	3,024,000	10	8	0.00005	0.0005	0.47	0.002	0.014	trace	0.012	0.003	trace	94.1	0.3	2.8	NC	2.3	0.5	NC
<i>P. multiseris</i>			G007Ps04	f/2-Si	1,306,667	9	4	0.00008	0.0008	2.05	NC ^d	0.059	0	NC ^d	0.012	0.002	96.5	NC	2.8	NA	NC	0.6	0.1
<i>P. multiseris</i>			G010Ps10	f/2-Si	1,006,667	9	4	0.0001	0.001	7.20	0	0.206	0	0.239	0.019	0.005	93.9	NA	2.7	NA	3.1	0.3	0.1
<i>P. multistriata</i>	ST1	CAWB137	PA014Ps09	f/2	51,150,000	10	5	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multistriata</i>	ST1		PA014Ps10	f/2	51,562,500	10	5	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	10,506,667	8	11	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multistriata</i>	ST1	CAWB150	PB024Ps01	f/2-Si	10,240,000	9	4	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multistriata</i>	ST1		G013Ps21	f/2-Si	7,013,333	9	4	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multistriata</i>	ST1		G027Ps06	f/2-Si	3,040,000	8	5	0.00005	0.0005	trace	0	trace	0	0	0	0	NC	NA	NC	NA	NA	NA	NA
<i>P. multistriata</i>	ST2	CAWB127	827	f/2	50,700,000	10	19	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. multistriata</i>	ST2		G001Ps01	f/2-Si	4,213,333	8	6	0.00005	0.0005	trace	0	trace	0	0	0	0	NC	NA	NC	NA	NA	NA	NA
<i>P. multistriata</i>	ST2		G001Ps02	f/2-Si	11,216,000	10	8	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. plurisecta</i>		CAWB151	G027Ps03	f/2-Si	7,733,333	9	3	0.00005	0.0005	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
			f/2-Si	962,667	43	4	0.0001	0.001	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1	CAWB152	PA030Ps03	f/2-Si	953,333	9	4	0.0001	0.001	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		G007Ps06	f/2-Si	3,536,000	10	8	0.00005	0.0005	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		G005Ps01	f/2-Si	3,216,000	10	8	0.00005	0.0005	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		G010Ps01	f/2-Si	3,744,000	10	8	0.00005	0.0005	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		G018Ps05	f/2-Si	1,664,000	10	7	0.00006	0.0006	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1	Beatrix Bay	f/2	8,100,000	10	8	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		G011Ps01	f/2-Si	1,368,000	9	4	0.00007	0.0007	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		J328Ps21	f/2-Si	380,000	10	3	0.0003	0.003	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST1		J237Ps03	f/2-Si	555,000	10	3	0.0002	0.002	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	ST2		G027Ps05	f/2-Si	1,216,000	10	7	0.00008	0.0008	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA

<i>P. pungens</i>	clade I	ST2	G015Ps02	<i>t</i> /2–Si	1,824,000	10	7	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	Unknown	CAWB129	Ps0-3	<i>t</i> /2	16,020,000	10	13	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. pungens</i>	clade I	Unknown	CAWB130	Ps0-4	<i>t</i> /2	8,700,000	10	13	0.0005	0.0025	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. cf. subpaci</i>			PA030Ps02	<i>t</i> /2–Si	3,453,333	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. cf. subpaci</i>			PA030Ps07	<i>t</i> /2–Si	3,440,000	10	8	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. cf. subpaci</i>		CAWB148	PD009Ps04	<i>t</i> /2–Si	3,813,333	9	3	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA
<i>P. cf. subpaci</i>			G011Ps03	<i>t</i> /2–Si	4,080,000	9	4	0.00005	0.0005	0	0	0	0	0	0	0	0	NA	NA	NA	NA	NA	NA	NA

^a Sequences that were not identical but were not a clade/subclade level divergence were assigned as sequence type (ST).

^b CICCM: the Cawthron Institute Culture Collection of Microalgae.

^c The number of months took after isolation at the time of cultivation for DA and DA isomers analysis.

^d Possible mix of epi-DA and iso-DA C was detected. These analogues co-eluted and could not be differentiated from each other for this sample.