

Revised phylogeny and historical biogeography of the cosmopolitan aquatic plant genus

Typha (Typhaceae)

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Table S1 Ancestral area reconstruction using the S-DIVA and DEC analysis based on the chronogram inferred using BEAST. Node number is indicated in Figure 3. PP is the posterior probability of the node. Letter codes represent biogeographic regions: A - North America, B - Indo-Pacific, C - West Eurasia, D - East Eurasia, E - Africa, F - South America, and G - Australia. Support for specific reconstructions are indicated by the integer to the right of the letter code. Reconstructions with less than 5% support were omitted.

Node	Age (Mya)		Ancestral areas				
	Mean	95% HPD	PP	S-DIVA		DEC	
				Area	Probability	Area	Probability
12	10.01	4.20-19.20	1.00	AC/CE/ACE/CDE/CD/ACD/C/ACDE	12.5/12.5/12.5/12.5/12.5/12.5/12.5	ACDE/C/CD/ACD/CDE	27/22/18/9/9
13	--	--	0.90	ABDE/ABE/ABCD/ABCE/A CDE/BCD/BCDE/BCE/CD/B C/D/ABC	8.3/8.3/8.3/8.3/8.3/8.3/8.3/8.3/8.3/8.3/8.3/8.3	D/ACDE/CD/ABCD/BCDE/C/BCD/BC	29/17/13/7/7/7/6/5
14	3.99	1.38-8.53	1.00	BDEG/DE/DEG/BDE	25/25/25/25	DE/BDEG/BDE /DEG/D	28/18/16/16/8
15	5.73	2.27-11.80	1.00	D	100	D/DE/ACDF/BDEG/CD/AD	52/14/10/9/8/7
16	8.43	3.40-16.85	1.00	D	100	D/CD/AD	73/14/13
17	10.30	4.09-19.84	1.00	D	100	D	100
18	--	--	0.89	D	100	D	100
19	17.20	7.99-30.86	1.00	BCD/ABCD/ABDE/BCDE/A CDE/D	16.7/16.7/16.7/16.7/16.7/16.7	D/ACDE/ABCD/BCDE/CD/BC D	20/16/10/10/7/5
20	11.11	3.78-24.01	1.00	D/BCD/BC	33/33/33	D/BCD/CD/BD/ABCD/BCDE/BCDG/BCDF	28/17/12/12/8/8/8/8
21	39.03	22.64-57.59	1.00	D/BCD/ABCD/ABDE/BCDE/ACDE	16.7/16.7/16.7/16.7/16.7/16.7	D	8

Table S2 The collection information of samples used in this study and their GenBank accession numbers of seven chloroplast DNA regions.

Species	Collection site	Voucher No.	Code	atpB-rbcL	psbA-trnH	psbD-trnT	rpl32-trnL	rps16 intro	rps16-trnK	trnL-trnF
<i>Typha angustifolia</i>	China, Jinlin	Xu0068	<i>T. angustifolia_China1</i>	MG324303	MG430801	MG431016	MG430844	MG430887	MG430930	MG430973
<i>Typha angustifolia</i>	China, Qinghai	Tu519-1	<i>T. angustifolia_China2</i>	MG324304	MG430802	MG431017	MG430845	MG430888	MG430931	MG430974
<i>Typha angustifolia</i>	China, Shaanxi	Tu522	<i>T. angustifolia_China3</i>	MG324305	MG430803	MG431018	MG430846	MG430889	MG430932	MG430975
<i>Typha angustifolia</i>	China, Shaanxi	Tu523	<i>T. angustifolia_China4</i>	MG324306	MG430804	MG431019	MG430847	MG430890	MG430933	MG430976
<i>Typha angustifolia</i>	China, Hainan	Xu2012	<i>T. angustifolia_China5</i>	MG324307	MG430805	MG431020	MG430848	MG430891	MG430934	MG430977
<i>Typha angustifolia</i>	China, Xingjiang	Xu4155	<i>T. angustifolia_China6</i>	MG324318	MG430816	MG431031	MG430859	MG430902	MG430945	MG430988
<i>Typha angustifolia</i>	China, Xingjiang	Xu4450	<i>T.angustifolia_China7</i>	MG324319	MG430817	MG431032	MG430860	MG430903	MG430946	MG430989
<i>Typha angustifolia</i>	Germany, Bavaria	S. Renner s.n.	<i>T. angustifolia_Germany</i>	MG324320	MG430818	MG431033	MG430861	MG430904	MG430947	MG430990
<i>Typha angustifolia</i>	USA, Virginia	Wen9820	<i>T. angustifolia_USA</i>	MG324321	MG430819	MG431034	MG430862	MG430905	MG430948	MG430991
<i>Typha domingensis</i>	China, Yunnan	Xu2212	<i>T. domingensis_China1</i>	MG324308	MG430806	MG431021	MG430849	MG430892	MG430935	MG430978
<i>Typha domingensis</i>	China, Yunnan	Tu507	<i>T. domingensis_China2</i>	MG324309	MG430807	MG431022	MG430850	MG430893	MG430936	MG430979
<i>Typha domingensis</i>	China, Xinjiang	Xu2564	<i>T. domingensis_China3</i>	MG324315	MG430813	MG431028	MG430856	MG430899	MG430942	MG430985
<i>Typha domingensis</i>	Israel, Beersheba	S. Volis s.n.	<i>T. domingensis_Israel</i>	MG324316	MG430814	MG431029	MG430857	MG430900	MG430943	MG430986
<i>Typha domingensis</i>	Indonesia, West Java	Wen10698	<i>T. domingensis_Indonesia</i>	MG324311	MG430809	MG431024	MG430852	MG430895	MG430938	MG430981
<i>Typha domingensis</i>	Pakistan, Khyber Pakhtunkhwa	S.K.Marwat s.n.	<i>T. domingensis_Pakistan</i>	MG324310	MG430808	MG431023	MG430851	MG430894	MG430937	MG430980
<i>Typha domingensis</i>	USA, Texas	Moody1070	<i>T. domingensis_USA</i>	MG324317	MG430815	MG431030	MG430858	MG430901	MG430944	MG430987
<i>Typha latifolia</i>	China, Heilongjiang	Xu107	<i>T. latifolia_China1</i>	MG324324	MG430822	MG431037	MG430865	MG430908	MG430951	MG430994
<i>Typha latifolia</i>	China, Xingjiang	Xu4436	<i>T. latifolia_China2</i>	MG324325	MG430823	MG431038	MG430866	MG430909	MG430952	MG430995
<i>Typha latifolia</i>	USA, Virginia	Wen9821	<i>T. latifolia_USA1</i>	MG324328	MG430826	MG431041	MG430869	MG430912	MG430955	MG430998
<i>Typha latifolia</i>	USA, Alaska	Moody1077	<i>T. latifolia_USA2</i>	MG324329	MG430827	MG431042	MG430870	MG430913	MG430956	MG430999
<i>Typha latifolia</i>	Kenya, Kitale	X. Yan s.n.	<i>T. latifolia_Kenya</i>	MG324323	MG430821	MG431036	MG430864	MG430907	MG430950	MG430993
<i>Typha latifolia</i>	Germany, Bavaria	D/0053	<i>T. latifolia_Germany</i>	MG324327	MG430825	MG431040	MG430868	MG430911	MG430954	MG430997
<i>Typha latifolia</i>	UK, Bramhope	Reader s.n.	<i>T. latifolia_England</i>	MG324326	MG430824	MG431039	MG430867	MG430910	MG430953	MG430996

<i>Typha orientalis</i>	China, Zhejiang	Xu1065	<i>T. orientalis_China1</i>	MG324330	MG430828	MG431043	MG430871	MG430914	MG430957	MG431000
<i>Typha orientalis</i>	China, Hunan	Tu505	<i>T. orientalis_China2</i>	MG324331	MG430829	MG431044	MG430872	MG430915	MG430958	MG431001
<i>Typha orientalis</i>	China, Guangdong	Tu502	<i>T. orientalis_China3</i>	MG324332	MG430830	MG431045	MG430873	MG430916	MG430959	MG431002
<i>Typha laxmannii</i>	China, Guangdong	Tu526	<i>T. laxmannii_China1</i>	MG324333	MG430831	MG431046	MG430874	MG430917	MG430960	MG431003
<i>Typha laxmannii</i>	China, Shaanxi	Xu5239	<i>T. laxmannii_China2</i>	MG324334	MG430832	MG431047	MG430875	MG430918	MG430961	MG431004
<i>Typha laxmannii</i>	China, Jilin	Xu0069	<i>T. laxmannii_China3</i>	MG324335	MG430833	MG431048	MG430876	MG430919	MG430962	MG431005
<i>Typha laxmannii</i>	China, Qinghai	Tu518-1	<i>T. laxmannii_China4</i>	MG324336	MG430834	MG431049	MG430877	MG430920	MG430963	MG431006
<i>Typha laxmannii</i>	Germany, Bavaria	D/0222	<i>T. laxmannii_Germany</i>	MG324337	MG430835	MG431050	MG430878	MG430921	MG430964	MG431007
<i>Typha minima</i>	China, Shaanxi	Xu5235	<i>T. minima_China</i>	MG324338	MG430836	MG431051	MG430879	MG430922	MG430965	MG431008
<i>Typha minima</i>	Germany, Bavaria	D/0054	<i>T. minima_Germany</i>	MG324339	MG430837	MG431052	MG430880	MG430923	MG430966	MG431009
<i>Typha elephantina</i>	China, Yunnan	Tu506	<i>T. elephantina_China1</i>	MG324340	MG430838	MG431053	MG430881	MG430924	MG430967	MG431010
<i>Typha elephantina</i>	China, Yunnan	Tu515	<i>T. elephantina_China2</i>	MG324341	MG430839	MG431054	MG430882	MG430925	MG430968	MG431011
<i>Typha elephantina</i>	Pakistan, Khyber Pakhtunkhwa	S.K.Marwat s.n.	<i>T. elephantina_Pakistan</i>	MG324342	MG430840	MG431055	MG430883	MG430926	MG430969	MG431012
<i>Typha capensis</i>	South Africa, Hogsback	S. Compton s.n	<i>T. capensis_South Africa1</i>	MG324312	MG430810	MG431025	MG430853	MG430896	MG430939	MG430982
<i>Typha capensis</i>	South Africa, Free State	Wen10075	<i>T. capensis_South Africa2</i>	MG324313	MG430811	MG431026	MG430854	MG430897	MG430940	MG430983
<i>Typha capensis</i>	South Africa, KwaZulu-Natal	Wen10085	<i>T. capensis_South Africa3</i>	MG324314	MG430812	MG431027	MG430855	MG430898	MG430941	MG430984
<i>Typha shuttleworthii</i>	Germany, Bavaria	D/0055	<i>T. shuttleworthii_Germany</i>	MG324322	MG430820	MG431035	MG430863	MG430906	MG430949	MG430992
<i>Sparganium stoloniferum</i>	China, Heilongjiang	Xu3874	<i>S. stoloniferum_China</i>	MG324344	MG430842	MG431057	MG430885	MG430928	MG430971	MG431014
<i>Sparganium eurycarpum</i>	USA, Virginia	Strong s.n.	<i>S. eurycarpum_USA</i>	MG324343	MG430841	MG431056	MG430884	MG430927	MG430970	MG431013
<i>Sparganium glomeratum</i>	China, Heilongjiang	Xu3905	<i>S. glomeratum_China</i>	MG324345	MG430843	MG431058	MG430886	MG430929	MG430972	MG431015

Table S3 The primers of seven chloroplast DNA regions and their sources

Regions	Name and sequence (5'-3')	Reference
<i>atpB-rbcL</i>	<i>atpB</i> : GTG GAA ACC CCG GGA CGA GAA GTA GT <i>rbcL</i> : ACT TGC TTT AGT TTC TGT TTG TGG TGA	Hodges and Arnold, 1994
<i>psbA-trnH</i>	<i>trnH</i> ^{GUU} : CGC GCA TGG TGG ATT CAC AAT CC <i>psbA</i> : GTT ATG CAT GAA CGT AAT GCT C	Shaw et al., 2005
<i>psbD-trnT</i>	<i>psbD</i> : CTC CGT ARC CAG TCA TCC ATA <i>trnT</i> ^(GGU) -R: CCC TTT TAA CTC AGT GGT AG	Shaw et al., 2007
<i>rpl32-trnL</i>	<i>trnL</i> ^(UAG) : CTG CTT CCT AAG AGC AGC GT <i>rpl32</i> -F: CAG TTC CAA AA A AAC GTA CTT C	Shaw et al., 2007
<i>rps16 intron</i>	<i>rpS16F</i> : AAA CGA TGT GGT ARA AAG CAA C <i>rpS16R</i> : AAC ATC WAT TGC AAS GAT TCG ATA	Shaw et al., 2005
<i>rps16-trnK</i>	<i>rpS16x2F2</i> : AAA GTG GGT TTT TAT GAT CC <i>trnK</i> ^(UUU) _{x1} : TTA AAA GCC GAG TAC TCT ACC	Shaw et al., 2007
<i>trnL-trnF</i>	<i>e</i> : GGT TCA AGT CCC TCT ATC CC <i>f</i> : ATT TGA ACT GGT GAC ACG AG	Taberlet et al., 1991

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Taberlet P, Gielly L, Pautou G, Bouvet J. 1991. Universal primers for amplification of three non-coding regions of chloroplast DNA. *Plant Molecular Biology* 17:1105-1109.

Table S4 Dispersal matrix among 7 areas used in the DEC analysis. Letters represent biogeographic regions: A - North America, B - Indo-Pacific, C - West Eurasia, D - East Eurasia, E - Africa, F - South America, and G - Australia. Dispersal probabilities among areas are given for 0 - 5 Ma, 5 - 33 Ma, 33 - 49 Ma, respectively.

	B	C	D	E	F	G
A	0.1/0.1/0.25	0.1/0.25/0.5	0.1/0.1/0.25	0.1/0.1/0.25	1.0/0.75/0.5	0.1/0.1/0.1
B	/	1.0/1.0/0.75	1.0/1.0/0.75	0.25/0.25/0.25	0.1/0.1/0.1	0.75/0.5/0.25
C		/	1.0/1.0/1.0	0.75/0.5/0.25	0.1/0.1/0.1	0.25/0.25/0.1
D			/	0.5/0.25/0.1	0.1/0.1/0.1	0.5/0.25/0.1
E				/	0.1/0.1/0.5	0.1/0.1/0.1
F					/	0.1/0.1/0.1