SUPPLEMENTARY FILE

Defining evolutionary conservation units in the Macedonian crested newt, *Triturus macedonicus* (Amphibia; Salamandridae), in a biodiversity hotspot

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Table S1. Sample localities of *Triturus macedonicus* used in this study. Habitat information is given only for populations from Northern Pindos National Park: N: natural habitat (pond, pool, or ditch); A: artificial habitat (cattle pond, reservoir); n/a: not available.

Site	n of sequenced individuals in this study	n of additional sequences from the GenBank	Location	Coordinates	Habitat type	Haplotype	Clade	GenBank Accession #
1	2		Greece: Kerasovo	40.159 / 20.934	N	Tmac01 (2)	1	GU982387
2	2		Greece: Romios	40.131 /20.981	N	Tmac01 (2)	1	GU982387
3	1		Greece: Smolikas	40.120 / 20.880	N	Tmac01 (1)	1	GU982387
4	3		Greece: Samarina	40.105 / 21.073	N	Tmac01 (3)	1	GU982387
5	2		Greece: Lake Distratou	40.055 / 21.070	A	Tmac01 (1) Tmac31 (1)	1	GU982387 OQ877047
6	4		Greece: Avdella	39.996 / 21.106	N	Tmac01 (3) Tmac03 (1)	1	GU982387 JQ240240
7	1		Greece: Vovousa	39.952 / 21.087	A	Tmac01 (1)	1	GU982387
8	2		Greece: Lakos Avgou	39.925 / 21.115	N	Tmac01 (2)	1	GU982387
9	4		Greece: Elafotopos	39.918 / 20.696	A	Tmac02 (4)	2A	GU982388
10	2		Greece: Grabala	39.914 / 20.667	A	Tmac33 (2)	2B	OQ877049
11	2		Greece: Grabala	39.912 / 20.662	A	Tmac33 (2)	2B	OQ877049
12	2		Greece: Grabala	39.910 / 20.673	A	Tmac02 (1) Tmac33 (1)	2A 2B	GU982388 OQ877049
13	1		Greece: Stouros	39.909 / 20.721	A	Tmac02 (1)	2A	GU982388
14	2		Greece: Elafotopos	39.907 / 20.681	A	Tmac02 (2)	2A	GU982388
15	3		Greece: Elafotopos	39.906 / 20.694	A	Tmac02 (3)	2A	GU982388
16	2		Greece: Petrino Dasos	39.901 / 20.742	A	Tmac02 (2)	2A	GU982388
17	1		Greece: Stouros	39.901 / 20.719	A	Tmac02 (1)	2A	GU982388
18	2	4 [28]	Greece: Elafotopos	39.900 / 20.669	A	Tmac02 (5) Tmac28 (1)	2A	GU982388 JQ240265
19	2		Greece: Stouros	39.892 / 20.725	A	Tmac02 (2)	2A	GU982388
20	2		Greece: Stouros	39.891 / 20.721	A	Tmac02 (2)	2A	GU982388

21	3		Greece: Ano Pedina	39.888 / 20.701	A	Tmac02 (3)	2A	GU982388
22	1		Greece: Kato Pedina Plain	39.886 / 20.682	A	Tmac02 (1)	2A	GU982388
23	3		Greece: Kato Pedina Plain	39.885 / 20.676	A	Tmac02 (3)	2A	GU982388
24	2		Greece: Monodendri	39.880 / 20.738	A	Tmac02 (2)	2A	GU982388
25	1		Greece: Ano Pedina - Monodendri	20.722 / 39.880	A	Tmac02 (1)	2A	GU982388
26	4		Greece: Kato Pedina	39.879 / 20.684	A	Tmac02 (4)	2A	GU982388
27	3		Greece: Kato Pedina	39.877 / 20.670	A	Tmac02 (3)	2A	GU982388
28	2		Greece: Tsouka Rossa	39.876 / 21.038	N	Tmac01 (2)	1	GU982387
29	1		Greece: Central Zagori	39.866 / 20.682	A	Tmac02 (1)	2A	GU982388
30	3		Greece: Central Zagori	39.865 / 20.705	A	Tmac02 (2) Tmac32 (1)	2A	GU982388 OQ877048
31	1		Greece: Tsouka Rossa	39.864 / 21.033	N	Tmac01 (1)	1	GU982387
32	1		Greece: Central Zagori	39.863 / 20.721	A	Tmac02 (1)	2A	GU982388
33	3		Greece: Tsouka Rossa	39.863 / 21.040	N	Tmac01 (3)	1	GU982387
34	1		Greece: Central Zagori	39.854 / 20.700	A	Tmac02 (1)	2A	GU982388
35	4	5 [28]	Greece: Zygos-Mikro Koukourelo	39.823 / 21.261	N	Tmac01 (9)	1	GU982387
36	1		Greece: Aoos Springs Lake	39.817 / 21.097	N	Tmac01 (1)	1	GU982387
37	1		Greece: Aoos Springs Lake	39.817 / 21.097	N	Tmac01 (1)	1	GU982387
38	2		Greece: Aoos Springs Lake	39.816 / 21.101	N	Tmac01 (2)	1	GU982387
39	2		Greece: Delvinaki	39.946 / 20.489	A	Tmac33 (2)	2B	OQ877049
40	2		Greece: Lapsista	39.805 / 20.703	N	Tmac02 (2)	2A	GU982388
41	2		Greece: Lapsista	39.771 / 20.700	A	Tmac02 (2)	2A	GU982388
42	1		Greece: Serviana	39.564 / 20.882	A	Tmac34 (1)	2A	OQ877050
43		3 [28]	Greece: Negrades	39.847 / 20.663	A	Tmac02 (3)	2A	GU982388
44		1 [28]	Greece: Elati	39.835 / 20.740	N	Tmac02 (1)	2A	GU982388
45		1 [123]	Greece: Asprangeli	39.830 / 20.730	A	Tmac02 (1)	2A	GU982388
46		11 [28, 124, 125]	Greece: Ano Kaliniki	40.853 / 21.450	n/a	Tmac01 (10 Tmac03 (1)	1	GU982387 JQ240240
47		7 [28]	Greece: Kerameia	39.562 / 22.081	n/a	Tmac29 (7)	3	JQ240266
48		4 [28]	Greece: Corfu Island	39.562 / 19.764	n/a	Tmac16 (1) Tmac17 (1) Tmac18 (1) Tmac19 (1)	1	JQ240253 JQ240254 JQ240255 JQ240256
49		5 [28]	Greece: Paliambela – Amvrakikos Gulf	38.909 / 20.970	n/a	Tmac13 (5)	1	JQ240250
50		4 [28]	Greece: Pefkodasos	41.030 / 22.580	n/a	Tmac01 (4)	1	GU982387
51		5 [28]	Greece: Prespa	40.831 / 21.121	n/a	Tmac01 (4) Tmac20 (1)	1	GU982387 JQ240257
52		5 [28]	Greece: Seli	40.567 / 22.022	n/a	Tmac25 (1) Tmac26 (4)	3	JQ240262 JQ240263
53		5 [28]	Greece: Trilofos	40.358 / 22.477	n/a	Tmac30 (5)	3	JQ240267

54		1 [28]	Greece: Vryta	40.800 / 21.919	n/a	Tmac29 (1)	3	JQ240266
55		5 [28]	Albania: Bajzë	42.303 / 19.441	n/a	Tmac11 (2) Tmac12 (3)	1	JQ240248
56		5 [28]	Albania: Bejar	40.429 / 19.850	n/a	Tmac05 (3) Tmac27 (2)	1 2B	JQ240242 JQ240264
57		5 [28]	Albania: Fushë-Krujë	41.529 / 19.682	n/a	Tmac04 (2) Tmac14 (2) Tmac15 (1)	1	JQ240241 JQ240251 JQ240252
58		5 [28]	Albania: Gracen	41.157 / 19.949	n/a	Tmac21 (5)	1	JQ240258
59		5 [28]	Albania: Kavajë	41.168 / 19.556	n/a	Tmac04 (1) Tmac06 (3) Tmac07 (1)	1	JQ240241 JQ240243 JQ240244
60		5 [28]	Albania: Kolsh	42.072 / 20.326	n/a	Tmac08 (2) Tmac09 (3)	1	JQ240245 JQ240246
61		5 [28]	Albania: Koplik	42.197 / 19.451	n/a	Tmac11 (4) Tmac12 (1)	1	JQ240248 JQ240249
62		5 [28]	Albania: Lushnjë	41.000 /19.664	n/a	Tmac04 (3) Tmac23 (2)	1	JQ240241 JQ240260
63		5 [28]	Albania: Pogradec	40.931 / 20.640	n/a	Tmac22 (4) Tmac24 (1)	1	JQ240259 JQ240261
64		4 [28]	Kosovo: Karagač	42.640 / 20.309	n/a	Tmac08 (4)	1	JQ240245
65		3 [28]	Kosovo: Kijevo	42.568 / 20.728	n/a	Tmac08 (3)	1	JQ240245
66		1 [28]	Kosovo: Xërze	42.341 / 20.573	n/a	Tmac08 (1)	1	JQ240245
67		5 [28]	North Macedonia: Debreste	41.502 / 21.293	n/a	Tmac08 (5)	1	JQ240245
68		5 [28]	North Macedonia: Gorno Srpci	41.072 / 21.219	n/a	Tmac01 (5)	1	GU982387
69		4 [28]	North Macedonia: Gostivar	41.817 / 20.899	n/a	Tmac01 (4)	1	GU982387
70		5 [28]	North Macedonia: Kolibari	41.580 / 20.951	n/a	Tmac08 (3) Tmac10 (2)	1	JQ240245 JQ240247
71		5 [28]	North Macedonia: Leskoec	40.962 / 20.885	n/a	Tmac20 (1) Tmac22 (4)	1	JQ240257 JQ240259
72		5 [28]	North Macedonia: Loznani	41.221 / 21.446	n/a	Tmac03 (5)	1	JQ240240
73		2 [28]	North Macedonia: Prilep	41.419 / 21.611	n/a	Tmac03 (2)	1	JQ240240
74		5 [28]	North Macedonia: Vrbjani	41.413 / 20.816	n/a	Tmac08 (5)	1	JQ240245
75		2 [28]	Montenegro: Bjeloši	42.374 / 18.907	n/a	Tmac11 (2)	1	JQ240248
76		5 [28]	Montenegro: Donji Lokanj	41.900 / 19.267	n/a	Tmac11 (5)	1	JQ240248
77		1 [28]	Montenegro: Kućišta	42.460 / 18.854	n/a	Tmac11 (1)	1	JQ240248

Table S2. Primer pairs used in this study for the amplification of seven microsatellite loci [42]. The fluorescent dyes used, the allele range and the number of detected alleles in the present study are given.

Microsatellite loci	Microsatellite Primers Sequence (5′–3′)	Repeat motif	Allele length (bp)	Number of alleles
TCM-83	F: FAM-CTGGCCATACAGTGGTTGTC R: TGGCCAGCAAGTAAGGAGT	$(TG)_9$	105-107	2
TCM-96	F: ROX-GCTCATTTCTTACACCACATCA R: GGAGGAACTGAGCCCTATCC	$(CA)_{10}$	121-143	7
TCM-273	F: HEX-CGCCAAGTTCATGTAGATGCT R: GAGACGAGACCGTATTCATCG	$(GT)_{11}$	136-150	3
TCM-414	F: FAM-TTTTGGCACCAATCAACGTAT R: CACGTAGTGGTGCAGAGTGAG	$(CA)_2CG(CA)_{11}$	130-138	4
TCM-417	F: ROX-GGCTTTATTTCAGCTGTCTTCC R: GGCTGACTTCATATGCCTGAC	$(CA)_7(CGCA)_3(CA)_3(CGCA)_3(CTCA)3(CT)_2C_4AC_5$	172-186	3
TCM-496	F: FAM-TGTGCAAACTGCACCTTTAG R: TTAAACCCATTGGCTGTTGT	$(AC)_{14}$	184-200	9
TCM-517	F: HEX-CCCACCCATACAGTGACTCTT R: GTGAGTGAGCAGGTGTGTGAG	$(TCAC)_{25}$	184-264	18

F: Forward; R: Reverse.

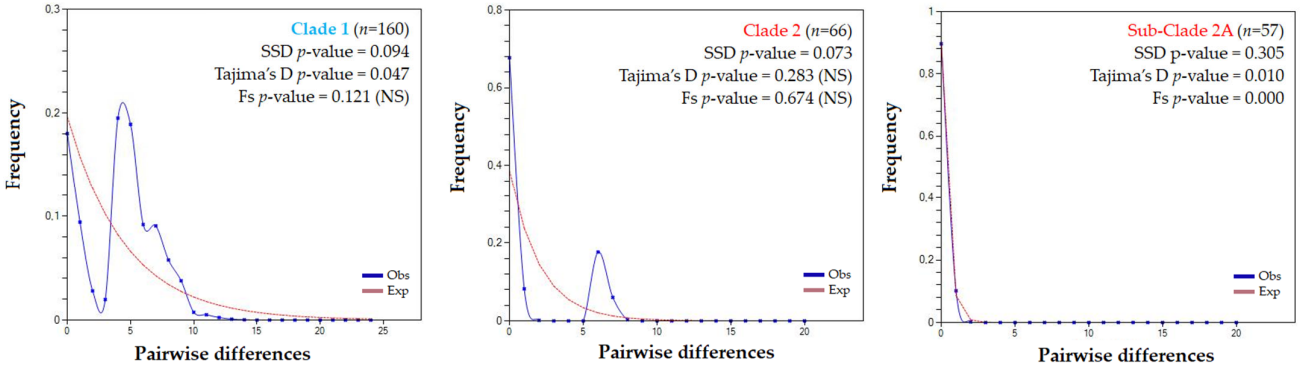

Figure S1. Mismatch distributions and neutrality tests for Clades 1, 2 and Sub-Clade 2A. *n*: number of sequences, NS: non-significant, Obs: Observed distribution of frequencies, Exp: Expected distribution of frequencies.

Table S3. Mismatch analysis and neutrality tests for clades and sub-clades with sufficient number of sequences (Clades 1 and 2, Sub-Clade 2A).

	Clade 1	**Clade 2**	**Sub-Clade 2A**
	n=160	*n=66*	*n=57*
Tau (τ)	6.027	3.000	3.000
Theta 0 (θ_0)	0.000	0.000	0.000
Theta 1 (θ_1)	8.740	0.341	0.120
SSD	0.031	0.532	0.000
SSD *p*-value	**0.094**	**0.073**	**0.305**
Tajima's *D*	-1.424	-0.669	-1.682
***D p*-value**	**0.047**	0.283 (NS)	**0.010**
Fu's *Fs*	-4.385	0.695	-4.555
***Fs p*-value**	0.121 (NS)	0.674 (NS)	**0.000**

n: number of sequences; NS: non-significant.

Table S4. Distribution of haplotypes and diversity measures of each clade and sub-clade identified in the phylogenetic analysis.

Haplotype	Clade 1	Clade 2	Sub-Clade 2A	Sub-Clade 2B	Clade 3	TOTAL
Tmac01	61					61
Tmac02		54	54			54
Tmac03	9					9
Tmac04	6					6
Tmac05	3					3
Tmac06	3					3
Tmac07	1					1
Tmac08	24					24
Tmac09	3					3
Tmac10	2					2
Tmac11	14					14
Tmac12	3					3
Tmac13	5					5
Tmac14	2					2
Tmac15	1					1
Tmac16	1					1
Tmac17	1					1
Tmac18	1					1
Tmac19	1					1
Tmac20	2					2
Tmac21	5					5
Tmac22	8					8
Tmac23	2					2
Tmac24	1					1
Tmac25					1	1
Tmac26					4	4
Tmac27		2		2		2
Tmac28		1	1			1

Tmac29					8	8
Tmac30					5	5
Tmac31	1					1
Tmac32		1	1			1
Tmac33		7		7		7
Tmac34		1	1			1
n	160	66	57	9	18	244
Hd	0.820	0.323	0.103	0.389	0.712	0.872
SD	± 0.025	±0.071	±0.055	±0.164	±0.065	±0.013
Pi	0.00622	0.00241	0.00016	0.00059	0.00146	0.01692
SD	±0.0004	±0.00059	±0.00009	±0.00025	±0.00025	±0.00065

n: number of sequences (sample size); *Hd*: haplotype diversity (±SD); *Pi*: nucleotide diversity (±SD).

Table S5. Inbreeding coefficient (F_{IS}) and percentage of randomizations that gave smaller (F_{IS} 95% CI+) or larger (F_{IS} 95% CI-) values than observed. Confidence Interval: 95% CI= 0.00036, adjusted for 120,000 randomizations.

Population	F_{IS}	F_{IS} 95% CI+	F_{IS} 95% CI-
4	0.183	0.032	0.985
5	-0.071	0.773	0.389
6	0.045	0.377	0.661
9	0.048	0.222	0.786
11	0.198	0.057	0.976
15	-0.054	0.812	0.266
18	-0.069	0.832	0.283
21	-0.019	0.610	0.392
22	0.066	0.235	0.799
23	0.131	0.026	0.979
24	-0.139	0.917	0.100
25	0.364	0.005	0.997
26	0.055	0.190	0.810
27	0.105	0.019	0.981
29	-0.099	0.794	0.253
30	0.105	0.010	0.990
33	-0.110	0.906	0.158
34	0.242	0.047	0.968
35	-0.031	0.675	0.434
38	0.122	0.202	0.889

Table S6. Hierarchical Analysis of Molecular Variance (AMOVA) among the two geographic areas in the park. Only the breeding sites with a sample size greater than six individuals were analyzed.

Source of variation	df	Sum of squares	Variance components	Percentage variation	Fixation Indices	*p*
Among areas	1	78.306	0.42733	17.82063	F_{ST}= 0.22984	0.0000
Among sites within areas	18	90.350	0.12383	5.16386	F_{SC}= 0.06284	0.0000
Within sites	248	935.584	1.84680	77.0155	F_{CT}= 0.17821	0.0000
Total		1104.24	2.39796			

Table S7. Posterior distributions of recent migration rates among breeding sites. The mean and 95 % CI in cases of insufficient data were 0.833 (0.675–0.992) for the non-migration rate and 0.0105 (1.43E-19–0.0933) for the migration rate. Migration rates >0.05 are shown in bold. The mean migration rates outside the expected CIs in the case of insufficient data are underlined. "Sink" populations are listed in the columns, and "source" populations are listed in the rows.

Population	4	5	6	9	11	15	18	21	22	23	24	25	26	27	29	30	33	34	35	38
4	**0.969** (0.89-0.99)	**0.104** (0.01-0.26)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.010 (0.00-0.07)	0.003 (0.00-0.03)	0.007 (0.00-0.05)	0.005 (0.00-0.03)	0.009 (0.00-0.06)	0.005 (0.00-0.03)	0.009 (0.00-0.06)	0.007 (0.00-0.05)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.007 (0.00-0.05)	0.001 (0.00-0.01)	0.007 (0.00-0.06)	0.007 (0.00-0.06)	**0.097** (0.00-0.27)	0.011 (0.00-0.08)
5	0.002 (0.00-0.02)	**0.710** (0.67-0.8)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.009 (0.00-0.06)	0.002 (0.00-0.02)	0.008 (0.00-0.05)	0.004 (0.00-0.03)	0.008 (0.00-0.06)	0.005 (0.00-0.03)	0.008 (0.00-0.05)	0.008 (0.00-0.07)	0.004 (0.00-0.03)	0.001 (0.00-0.01)	0.008 (0.00-0.06)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.009 (0.00-0.07)	0.006 (0.00-0.05)	0.007 (0.00-0.06)
6	0.002 (0.00-0.019)	0.028 (0.00-0.15)	**0.969** (0.89-0.99)	0.001 (0.00-0.01)	0.010 (0.00-0.07)	0.003 (0.00-0.02)	0.008 (0.00-0.05)	0.005 (0.00-0.03)	0.008 (0.00-0.05)	0.005 (0.00-0.03)	0.008 (0.00-0.06)	0.007 (0.00-0.05)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.06)	0.001 (0.00-0.01)	0.008 (0.00-0.08)	0.006 (0.00-0.05)	0.039 (0.00-0.18)	0.011 (0.00-0.08)
9	0.002 (0.00-0.02)	0.010 (0.00-0.06)	0.002 (0.00-0.02)	**0.982** (0.93-0.99)	0.015 (0.00-0.1)	0.004 (0.00-0.05)	0.009 (0.00-0.06)	**0.098** (0.00-0.26)	0.037 (0.00-0.15)	0.009 (0.00-0.06)	**0.079** (0.00-0.22)	0.024 (0.00-0.15)	0.010 (0.00-0.06)	0.001 (0.00-0.01)	0.050 (0.00-0.18)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.010 (0.00-0.08)	0.006 (0.00-0.04)	0.007 (0.00-0.06)
11	0.002 (0.00-0.01)	0.010 (0.00-0.067)	0.001 (0.00-0.02)	0.001 (0.00-0.01)	**0.720** (0.67-0.83)	0.001 (0.00-0.01)	0.009 (0.00-0.06)	0.004 (0.00-0.03)	0.006 (0.00-0.05)	0.005 (0.00-0.03)	0.008 (0.00-0.05)	0.007 (0.00-0.07)	0.003 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.009 (0.00-0.07)	0.008 (0.00-0.05)	0.004 (0.00-0.04)
15	0.002 (0.00-0.02)	0.010 (0.00-0.07)	0.002 (0.00-0.02)	0.001 (0.00-0.02)	0.027 (0.00-0.13)	**0.956** (0.83-0.99)	0.008 (0.00-0.06)	0.009 (0.00-0.05)	0.015 (0.00-0.09)	0.018 (0.00-0.09)	0.011 (0.00-0.07)	0.008 (0.00-0.06)	0.008 (0.00-0.05)	0.001 (0.00-0.01)	0.012 (0.00-0.07)	0.002 (0.00-0.02)	0.002 (0.00-0.02)	0.036 (0.00-0.15)	0.007 (0.00-0.05)	0.007 (0.00-0.05)
18	0.002 (0.00-0.017)	0.010 (0.00-0.07)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.002 (0.00-0.02)	**0.697** (0.667-0.77)	0.005 (0.00-0.04)	0.008 (0.00-0.06)	0.005 (0.00-0.04)	0.009 (0.00-0.07)	0.008 (0.00-0.06)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.001 (0.00-0.01)	0.002 (0.00-0.02)	0.011 (0.00-0.08)	0.007 (0.00-0.05)	0.007 (0.00-0.05)
21	0.002 (0.00-0.02)	0.010 (0.00-0.06)	0.001 (0.00-0.01)	0.001 (0.00-0.01)	0.024 (0.00-0.13)	0.003 (0.00-0.08)	0.011 (0.00-0.07)	**0.693** (0.67-0.77)	0.014 (0.00-0.08)	0.008 (0.00-0.05)	0.016 (0.00-0.10)	0.013 (0.00-0.10)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.023 (0.00-0.14)	0.001 (0.00-0.01)	0.002 (0.00-0.02)	0.026 (0.00-0.15)	0.007 (0.00-0.06)	0.007 (0.00-0.05)
22	0.002 (0.00-0.02)	0.010 (0.00-0.07)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.008 (0.00-0.06)	0.002 (0.00-0.02)	0.009 (0.00-0.06)	0.006 (0.00-0.03)	**0.702** (0.67-0.78)	0.004 (0.00-0.03)	0.008 (0.00-0.06)	0.009 (0.00-0.06)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.008 (0.00-0.05)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.011 (0.00-0.09)	0.007 (0.00-0.05)	0.007 (0.00-0.06)
23	0.002 (0.00-0.02)	0.012 (0.00-0.08)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.013 (0.00-0.09)	0.002 (0.00-0.02)	0.010 (0.00-0.06)	0.006 (0.00-0.04)	0.021 (0.00-0.12)	**0.711** (0.67-0.83)	0.012 (0.00-0.08)	0.022 (0.00-0.14)	0.006 (0.00-0.04)	0.001 (0.00-0.01)	0.010 (0.00-0.07)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.008 (0.00-0.07)	0.007 (0.00-0.05)	0.007 (0.00-0.05)
24	0.002 (0.00-0.02)	0.009 (0.00-0.05)	0.002 (0.00-0.023)	0.001 (0.00-0.02)	0.010 (0.00-0.08)	0.002 (0.00-0.02)	0.009 (0.00-0.06)	0.006 (0.00-0.04)	0.010 (0.00-0.07)	0.006 (0.00-0.04)	**0.703** (0.67-0.80)	0.011 (0.00-0.08)	0.005 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.012 (0.00-0.09)	0.007 (0.00-0.05)	0.006 (0.00-0.05)

25	0.001 (0.00-0.01)	0.011 (0.00-0.08)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.003 (0.00-0.02)	0.010 (0.00-0.06)	0.006 (0.00-0.04)	0.010 (0.00-0.07)	0.006 (0.00-0.04)	0.010 (0.00-0.08)	**0.777** (0.67-0.99)	0.004 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.010 (0.00-0.08)	0.007 (0.00-0.05)	0.007 (0.00-0.06)
26	0.002 (0.00-0.02)	0.010 (0.00-0.07)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.050 (0.00-0.19)	0.002 (0.00-0.02)	0.010 (0.00-0.07)	0.005 (0.00-0.03)	0.015 (0.00-0.12)	0.015 (0.00-0.1)	0.012 (0.00-0.09)	0.014 (0.00-0.11)	0.689 (0.67-0.75)	0.001 (0.00-0.01)	0.023 (0.00-0.14)	0.001 (0.00-0.01)	0.002 (0.00-0.02)	0.013 (0.00-0.1)	0.007 (0.00-0.05)	0.007 (0.00-0.05)
27	0.002 (0.00-0.02)	0.009 (0.00-0.07)	0.002 (0.00-0.01)	0.001 (0.00-0.01)	0.030 (0.00-0.16)	0.006 (0.00-0.05)	0.118 (0.01-0.26)	0.127 (0.00-0.28)	0.080 (0.00-0.22)	0.136 (0.02-0.27)	0.034 (0.00-0.15)	0.018 (0.00-0.11)	0.172 (0.04-0.30)	0.983 (0.93-0.99)	0.054 (0.00-0.19)	0.002 (0.00-0.02)	0.003 (0.00-0.03)	0.026 (0.00-0.14)	0.007 (0.00-0.05)	0.006 (0.00-0.05)
29	0.002 (0.00-0.02)	0.011 (0.00-0.07)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.002 (0.00-0.02)	0.007 (0.00-0.04)	0.005 (0.00-0.03)	0.009 (0.00-0.06)	0.006 (0.00-0.04)	0.008 (0.00-0.05)	0.007 (0.00-0.06)	0.003 (0.00-0.03)	0.001 (0.00-0.01)	0.709 (0.67-0.81)	0.001 (0.00-0.01)	0.002 (0.00-0.02)	0.009 (0.00-0.07)	0.007 (0.00-0.04)	0.003 (0.00-0.03)
30	0.002 (0.00-0.02)	0.010 (0.00-0.06)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.016 (0.00-0.1)	0.005 (0.00-0.04)	0.061 (0.00-0.21)	0.011 (0.00-0.06)	0.039 (0.00-0.16)	0.051 (0.00-0.18)	0.057 (0.00-0.19)	0.048 (0.00-0.19)	0.065 (0.00-0.19)	0.003 (0.00-0.02)	0.024 (0.00-0.13)	0.983 (0.94-0.99)	0.003 (0.00-0.03)	0.016 (0.00-0.10)	0.006 (0.00-0.05)	0.007 (0.00-0.05)
33	0.003 (0.00-0.03)	0.018 (0.00-0.11)	0.002 (0.00-0.03)	0.001 (0.00-0.01)	0.009 (0.00-0.07)	0.002 (0.00-0.02)	0.008 (0.00-0.049)	0.005 (0.00-0.03)	0.008 (0.00-0.06)	0.005 (0.00-0.04)	0.008 (0.00-0.05)	0.008 (0.00-0.06)	0.004 (0.00-0.03)	0.001 (0.00-0.01)	0.008 (0.00-0.07)	0.001 (0.00-0.09)	0.947 (0.76-0.99)	0.008 (0.00-0.06)	0.026 (0.00-0.13)	0.068 (0.00-0.25)
34	0.002 (0.00-0.02)	0.009 (0.00-0.06)	0.001 (0.00-0.02)	0.001 (0.00-0.01)	0.011 (0.00-0.08)	0.002 (0.00-0.02)	0.007 (0.00-0.04)	0.006 (0.00-0.04)	0.008 (0.00-0.06)	0.005 (0.00-0.06)	0.008 (0.00-0.06)	0.007 (0.00-0.05)	0.003 (0.00 0.02)	0.001 (0.00-0.01)	0.011 (0.00-0.07)	0.002 (0.00-0.01)	0.003 (0.00-0.03)	0.763 (0.00-0.98)	0.007 (0.00-0.05)	0.003 (0.00-0.04)
35	0.003 (0.00-0.02)	0.016 (0.00-0.1)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.010 (0.00-0.08)	0.002 (0.00-0.02)	0.008 (0.00-0.05)	0.005 (0.00-0.03)	0.008 (0.00-0.05)	0.005 (0.00-0.03)	0.008 (0.00-0.05)	0.008 (0.00-0.06)	0.004 (0.00-0.03)	0.001 (0.00-0.01)	0.008 (0.00-0.06)	0.001 (0.00-0.01)	0.003 (0.00-0.03)	0.008 (0.00-0.06)	0.748 (0.67-0.99)	0.008 (0.00-0.07)
38	0.002 (0.00-0.02)	0.012 (0.00-0.08)	0.002 (0.00-0.02)	0.001 (0.00-0.01)	0.007 (0.00-0.05)	0.001 (0.00-0.01)	0.009 (0.00-0.05)	0.004 (0.00-0.03)	0.008 (0.00-0.06)	0.005 (0.00-0.04)	0.008 (0.00-0.05)	0.009 (0.00-0.06)	0.005 (0.00-0.04)	0.001 (0.00-0.01)	0.008 (0.00-0.06)	0.001 (0.00-0.01)	0.004 (0.00-0.04)	0.007 (0.00-0.05)	0.009 (0.00-0.06)	0.822 (0.67-0.99)

off-diagonal: migration rates; on-diagonal: non-migration rates; (95 % CI).

Table S8. List of private alleles detected in each of the two gene pools.

Gene pool 1 (Eastern group)		Gene pool 2 (Western group)	
Microsatellite locus	**Private allele (bp)**	**Microsatellite locus**	**Private allele (bp)**
TCM-96	129	TCM-96	123
TCM-96	143	TCM-273	136
TCM-414	132	TCM-414	136
TCM-496	184	TCM-414	138
TCM-496	190	TCM-417	184
TCM-496	200	TCM-496	186
TCM-517	200	TCM-496	188
TCM-517	204	TCM-496	194
TCM-517	208	TCM-517	184
TCM-517	244	TCM-517	220
TCM-517	248	TCM-517	228
TCM-517	252		
TCM-517	256		
TCM-517	264		

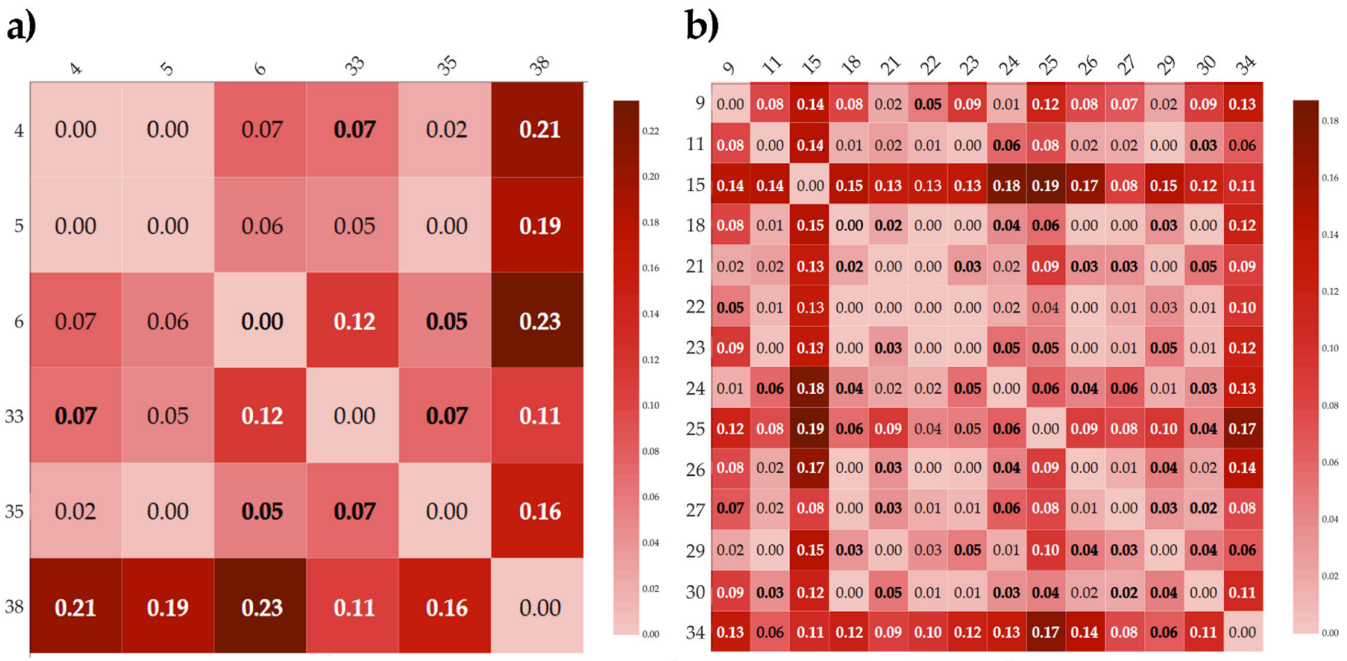

Figure S2. Heatmap of pairwise F_{ST} values among breeding sites (n ≥6, Table 1), within (**a**) Eastern and (**b**) Western genetic group. The 95% statistically significant values were obtained after 10,000 permutations and are given in bold. Site numbers correspond to Table S1 and Figure 2.

Table S9. Indices of genetic diversity per analyzed breeding site. Listed in order are the site ID, sample size (n), mean number of alleles per locus (A), corrected allelic richness (Ar), number of private alleles (Ap), expected (He) and observed (Ho) heterozygosity, with corresponding ±SE in parentheses. For site locations see Table S1 and Figure 3.

Site	n	A	Ar	Ap	He	Ho
4	11	3.429	2.971	1	0.451	0.390
		(1.152)	(0.842)		(0.096)	(0.106)
5	6	3.000	3.000	0	0.431	0.500
		(0.756)	(0.756)		(0.090)	(0.115)
6	10	3.000	2.565	0	0.404	0.408
		(0.655)	(0.457)		(0.081)	(0.089)
9	20	3.000	2.652	0	0.489	0.478
		(0.436)	(0.347)		(0.077)	(0.088)
11	6	3.286	3.286	0	0.508	0.452
		(0.808)	(0.808)		(0.106)	(0.144)
15	16	3.571	3.072	0	0.493	0.536
		(0.896)	(0.612)		(0.114)	(0.123)
18	9	3.571	3.350	0	0.535	0.603
		(0.719)	(0.620)		(0.088)	(0.108)
21	21	3.714	3.128	0	0.563	0.588
		(0.606)	(0.407)		(0.068)	(0.101)
22	8	3.571	3.373	0	0.537	0.538
		(0.719)	(0.629)		(0.096)	(0.123)
23	17	3.714	3.177	0	0.517	0.466
		(0.865)	(0.627)		(0.108)	(0.125)
24	8	3.143	2.999	0	0.507	0.612
		(0.553)	(0.499)		(0.078)	(0.121)
25	6	2.571	2.571	0	0.464	0.333
		(0.297)	(0.297)		(0.069)	(0.096)
26	18	3.429	3.040	0	0.498	0.485
		(0.612)	(0.508)		(0.096)	(0.139)
27	33	3.571	2.976	0	0.535	0.487
		(0.896)	(0.538)		(0.102)	(0.121)
29	6	2.857	2.857	0	0.487	0.581
		(0.595)	(0.595)		(0.096)	(0.123)
30	40	4.000	3.090	0	0.541	0.491
		(0.816)	(0.534)		(0.093)	(0.115)
33	10	2.857	2.742	0	0.467	0.543
		(0.595)	(0.527)		(0.081)	(0.092)
34	6	3.286	3.286	0	0.558	0.486
		(0.522)	(0.522)		(0.069)	(0.132)
35	10	3.143	2.803	1	0.488	0.529
		(0.738)	(0.560)		(0.094)	(0.113)
38	7	2.857	2.753	0	0.449	0.429
		(0.553)	(0.500)		(0.071)	(0.094)