



Table S1. List of individuals of the four broodstocks used in this study. Name of the stock, origin of the stock, genetic identification (ID), microchip number, length and weight of adult individuals, number of microsatellite loci, fragment length of mtDNA D-Loop and Cyt b are detailed.

Stock	Origin	ID	Microchip-02	Length (mm)	Weight (g)	N Loci	D-Loop	Cyt b
S1	Fattoria del Pesce	Hh01	311802	1117	10,480	13	1023 bp	
	Fattoria del Pesce	Hh02	311873	980	5720	13	1023 bp	
	Fattoria del Pesce	Hh03	311834	1260	12,000	13	1023 bp	
	Fattoria del Pesce	Hh04	311805	1230	11,700	11	1023 bp	
	Fattoria del Pesce	Hh05	311843	1150	8760	13	1023 bp	
	Fattoria del Pesce	Hh06	311887	1170	7220	13	1023 bp	
	Fattoria del Pesce	Hh07	311900	1180	10,100	13		
	Fattoria del Pesce	Hh08	311846	840	3160	13		
	Fattoria del Pesce	Hh09	311809	1260	12,600	13		
	Fattoria del Pesce	Hh10	311891	1130	7980	13	1023 bp	
	Fattoria del Pesce	Hh11	311881	1080	6880	13		
	Fattoria del Pesce	Hh12	311827	1060	7420	9		
	Fattoria del Pesce	Hh13	311874	1110	6620	13	1023 bp	
	Fattoria del Pesce	Hh14	311865	1080	6260	13	1023 bp	
	Fattoria del Pesce	Hh15	311831	1157	8340	12	1023 bp	
	Fattoria del Pesce	Hh16	100717222	1020	7000	13		
	Fattoria del Pesce	Hh17	311819	1030	7960	13	1023 bp	
	Fattoria del Pesce	Hh18	311864	1250	12,400	13	1023 bp	
	Fattoria del Pesce	Hh19	311810	1130	8460	10		
	Fattoria del Pesce	Hh20	311854	1135	9740	13		
	Fattoria del Pesce	Hh21	311818	1140	10,880	13		
	Fattoria del Pesce	Hh22	311812	1110	9240	13		
	Fattoria del Pesce	Hh23	311829	1060	7140	13		
	Fattoria del Pesce	Hh24	311856	1130	8900	13		
	Fattoria del Pesce	Hh25	311897	1170	10,740	13	1023 bp	
	Fattoria del Pesce	Hh26	311851	1210	10,240	13		
	Fattoria del Pesce	Hh27	311857	1160	9200	13		
	Fattoria del Pesce	Hh28	311878	970	9700	13		
	Fattoria del Pesce	Hh29	311880	1140	8600	13		
	Fattoria del Pesce	Hh30	311830	960	6280	13	1023 bp	
	Fattoria del Pesce	Hh31	311859	1120	9260	13	1023 bp	
	Fattoria del Pesce	Hh32	311836	1190	9820	13		
	Fattoria del Pesce	Hh33	311876	1060	6820	13	1023 bp	1118bp
	Fattoria del Pesce	Hh34	311847	1230	10,660	13	1023 bp	
	Fattoria del Pesce	Hh35	311893	920	3600	13		1118bp
	Fattoria del Pesce	Hh36	311860	1010	5460	13		
	Fattoria del Pesce	Hh37	311869	1160	8680	13	1023 bp	
	Fattoria del Pesce	Hh38	311886	1210	9840	13		
	Fattoria del Pesce	Hh39	311804	1160	8500	13	1023 bp	
	Fattoria del Pesce	Hh40	311883	1210	10,500	13	1023 bp	
	Fattoria del Pesce	Hh41	311801	1180	9630	13	1023 bp	
	Fattoria del Pesce	Hh42	311862	1190	9720	13	1023 bp	
	Fattoria del Pesce	Hh43	311896	1020	5820	13	1023 bp	
	Fattoria del Pesce	Hh44	311803	1040	6700	13	1023 bp	
	Fattoria del Pesce	Hh45	311822	980	4420	13	1023 bp	
	Fattoria del Pesce	Hh46	311866	1120	8680	13	1023 bp	
	Fattoria del Pesce	Hh47	311895	1110	7260	13	1023 bp	
S2	Piano Dossi	A01	/	/	/	13	1023 bp	1118bp
	Piano Dossi	A02	/	/	/	13	1023 bp	1118bp

	Piano Dossi	A03	/	/	/	13	1023 bp	1118bp
	Piano Dossi	A04	/	/	/	13	1023 bp	1118bp
	Piano Dossi	A05	/	/	/	13	1023 bp	1118bp
S2	Piano Dossi	A06	/	/	/	13	1023 bp	
	Piano Dossi	A07	/	/	/	13	1023 bp	
	Piano Dossi	A08	/	/	/	13	1023 bp	
	Piano Dossi	A09	/	/	/	13	1023 bp	
	Piano Dossi	A10	/	/	/	13	1023 bp	
	Piano Dossi	A11	/	/	/	13	1023 bp	
	Piano Dossi	A12	/	/	/	13	1023 bp	
S3	Neptum BT	HhHU1	982126055970712	/	/	13	1023 bp	1118bp
	Neptum BT	HhHU2	982126055970762	/	/	13	1023 bp	1118bp
	Neptum BT	HhHU3	982126055970752	/	/	13		
	Neptum BT	HhHU4	982126055970738	/	/	13		1118bp
	Neptum BT	HhHU5	982126055970742	/	/	13	1023 bp	
	Neptum BT	HhHU6	982126055970703	/	/	13	1023 bp	
	Neptum BT	HhHU7	982126055970673	/	/	13		
	Neptum BT	HhHU9	982126055970699	/	/	13	1023 bp	
	Neptum BT	HhHU10	982126055970669	/	/	13	1023 bp	
S4	L'Avannotto	MAN01	98201061212414	/	/	13	1023 bp	1118bp
	L'Avannotto	MAN02	98201061211839	/	/	13	1024 bp	1118bp
	L'Avannotto	MAN03	98201061212399	/	/	13	1025 bp	1118bp
	L'Avannotto	MAN04	98201061212479	/	/	13	1026 bp	
	L'Avannotto	MAN05	98201061212419	/	/	13	1027 bp	
	L'Avannotto	MAN06	98201061212413	/	/	13	1028 bp	
	L'Avannotto	MAN09	98201061212400	/	/	13	1029 bp	
	L'Avannotto	MAN12	98201061212445	/	/	13	1030 bp	
	L'Avannotto	MAN13	98201061212406	/	/	13	1031 bp	
	L'Avannotto	MAN14	98201061212451	/	/	13	1032 bp	
	L'Avannotto	MAN15	98201061212395	/	/	13	1033 bp	
	L'Avannotto	MAN16	98201061212429	/	/	13	1034 bp	
	L'Avannotto	MAN17	98201061212400	/	/	13	1035 bp	

Table S2. Microsatellite loci used in this study; name of the locus, primer sequence, repeat motif, annealing temperature, fluorescent probe and reference are listed.

Multiplex	Locus		Primer Sequence	Repeat motif	(°C)	Label	Reference
Mix 1	Anac_tag71527	F:	TGATCAACACCCACAGGCAT	(TA)6	60	FAM	[1]
		R:	CGCCATTCATCTCCCAAGAACAT				
	Anac_c15214	F:	ACAGCATTTTGGAGGGAACAA	(AC)13		PET	[1]
		R:	ACCTGTTTACTGACCAGTAGGT				
	Anac_c31601	F:	AAAGCATGGCAAAACATAGG	(TAA)10		VIC	[1]
		R:	AATCTCATGTTGGTGTCTTAC				
Mix 2	Anac_c6179	F:	GGAGACATGCAGGCCTTTCTA	(TTTA)7	60	FAM	[1]
		R:	TCCAAACTTGTGAAAAGTGCTGC				
	Anac_c16902	F:	CTACTTTGTGCCTGGGTAAGG	(AAT)7		PET	[1]
		R:	ATTGCTGTAAGTTGGGGTGG				
	Anac_c12159	F:	GCTGGATAACAATAATGCTAC	(AAT)11		VIC	[1]
		R:	TGGGAAACGTTATATTGACTTG				
Mix 3	An20	F:	AATAACAATCATTACATGAGGCT	(ATCT)10(TG)5	58	VIC	[2]
		R:	TGGTCAGTTGTTTTTTTATTGAT				
	Afug51	F:	ATAATAATGAGCGTGCTTTCTGTT	(AAAC)6(AC)2(AAAC)8		NED	[3]
		R:	ATTCCGCTTGCGACTTATTTA				
	AoxD165	F:	TTTGACAGCTCCTAAGTGATACC	(CTAT)13CTAC(CTAT)2		PET	[4]
		R:	AAAGCCCTACAACAAATGTCAC				
	AoxD161	F:	GTTTGAAATGATTGAGAAAATGC	(CTAT)15		FAM	[4]
		R:	TGAGACAGACACTCTAGTTAAACAGC				
Mix 4	Ag09	F:	CAGTCGATTTTCATTCAAAGCA	(TATG)15	60	FAM	[5]
		R:	CCTGCCCCCTAAACTGATTGA				
	Ag22	F:	TTTCCACACCTCCTCACTCA	(CT)12		NED	[5]
		R:	GGCAGTTTTTAACAGGGTTGC				
	Spl101	F:	CCCTCCACTGGAAATTTGAC	(TCTA)7		VIC	[6]
		R:	GCAATCAACAAGGTCTCTTTCA	(TA)6			

Table S3. References sequences retrieved from GenBank. Species, origin, GenBank accession number (code), and reference are details for both mitochondrial control region (D-Loop) and Cytochrome b gene.

D-Loop			
Species	Origin	Code	Reference
<i>Huso huso maeoticus</i>	Azov Sea	AY846666-67	[7]
<i>H. h. ponticus</i>	Black Sea	AY846668-79	[7]
<i>H. h. caspicus</i>	Caspian Sea	AY846640-65	[7]
<i>H. h. ponticus</i>	Danube Delta (Black Sea)	KF431840-46	[8]
Cytochrome b			
Species	Origin	Code	Reference
<i>H. huso</i>	Black Sea	AY846702-03; GU647228; KC30110-117	[7,9,10]
<i>H. huso</i>	Azov Sea	AY846708	[7]
<i>H. huso</i>	Caspian Sea	AY846704-707; AY846709-712; AJ245840	[7,11]
<i>H. huso</i>	Unknown	FJ974044	[9]
<i>Acipenser gueldenstaedtii</i>	-	KC130090-102	[10]
<i>A. stellatus</i>	-	KC130103-109	[10]
<i>A. ruthenus</i>	-	KF153104; GU647225; AJ249694	[9,11,12]

Table S4. Matrix of the observed allelic dimensions of each analysed locus for each individual. Genetic identification (ID), name of the stock and locus number are detailed.

Stock	ID	Anac_tag 71527	Anac_c15214	Anac_c31601	Anac_c6179	Anac_c16902	Anac_c12159	AoxD161	An20	AoxD165	Afug51	Ag22	Ag09	Sp101
S1	Hh01	201	205	259	259	297	300	122	126	340	340	242	245	98
	Hh02	201	205	269	269	300	309	118	122	334	340	242	248	98
	Hh03	201	209	267	267	294	300	122	126	334	340	242	248	98
	Hh04	201	201	269	269	294	306	122	126	334	340	0	0	0
	Hh05	201	201	267	267	297	306	118	118	334	334	248	248	98
	Hh06	201	205	267	267	297	300	118	122	334	340	248	248	98
	Hh07	205	209	267	267	297	300	118	122	334	340	0	0	98
	Hh08	201	201	269	269	300	309	118	118	334	334	251	251	98
	Hh09	199	201	261	261	300	309	118	122	340	340	239	248	98
	Hh10	199	205	267	267	300	303	118	122	334	334	242	248	98
	Hh11	201	201	0	0	297	300	118	122	340	340	242	248	98
	Hh12	201	205	0	0	297	306	118	122	0	0	0	0	98
	Hh13	201	209	269	269	297	306	118	122	334	334	248	248	98
	Hh14	199	205	267	267	300	303	118	118	334	340	233	236	98
	Hh15	201	203	265	265	294	294	118	118	334	340	248	248	102
	Hh16	201	209	269	269	294	300	118	122	334	340	242	248	98
	Hh17	199	205	267	267	297	306	118	122	340	340	248	248	98
	Hh18	205	209	267	267	300	303	118	122	334	340	242	248	98
	Hh19	201	201	269	269	297	300	118	122	334	334	0	0	0
	Hh20	199	203	0	0	297	297	118	122	334	334	0	0	98
	Hh21	201	201	265	267	297	300	118	118	334	334	242	248	98
	Hh22	201	205	267	267	297	300	118	122	334	340	248	248	98
	Hh23	199	205	265	265	297	306	118	122	334	340	242	248	98
	Hh24	205	209	267	267	297	306	118	118	334	340	242	248	98
	Hh25	205	209	267	267	297	306	118	118	334	334	251	251	98
	Hh26	201	201	259	259	297	306	118	122	340	340	239	242	98
	Hh27	199	205	259	267	297	300	118	118	340	340	239	248	98
	Hh28	205	209	267	267	294	300	118	122	334	340	0	0	98
	Hh29	199	199	267	267	300	309	118	118	334	334	242	248	98
	Hh30	201	205	269	269	300	309	118	122	340	340	0	0	98
	Hh31	201	205	267	267	297	300	118	118	340	340	248	248	98
	Hh32	201	205	261	261	300	309	118	118	340	340	239	242	98
	Hh33	201	205	267	269	300	309	118	122	340	340	242	248	98
	Hh34	199	205	267	267	297	300	118	122	340	340	239	242	98
	Hh35	199	205	267	267	297	306	118	118	334	340	233	242	98
	Hh36	199	205	269	269	300	309	118	118	334	334	233	248	98
	Hh37	201	205	269	269	297	300	118	122	340	340	242	248	98
	Hh38	201	205	267	267	294	306	118	118	334	334	242	248	98
	Hh39	201	205	261	261	300	309	118	122	340	340	239	242	98
	Hh40	201	201	269	269	297	306	118	122	334	340	248	248	90
	Hh41	201	201	269	269	297	300	118	118	334	340	242	242	98
	Hh42	201	205	267	267	294	306	118	122	334	340	242	242	98
	Hh43	199	205	267	267	297	306	118	122	334	340	248	248	98
	Hh44	201	201	269	269	300	303	118	122	340	340	242	248	98
	Hh45	199	199	269	269	300	309	118	122	340	340	233	242	98
	Hh46	199	205	269	269	300	303	118	122	334	334	233	242	98
	Hh47	199	205	267	267	297	306	118	118	334	340	242	248	98
S2	A01	199	209	269	269	300	312	118	118	328	334	239	239	98
	A02	201	205	267	267	306	318	118	118	331	334	257	257	98

Stock	ID	Anac_tag 71527	Anac_c15214	Anac_c31601	Anac_c6179	Anac_c16902	Anac_c12159	AoxD161	An20	AoxD165	Afug51	Ag22	Ag09	Sp101
S2	A03	201	209	267	267	306	306	118	118	331	334	236	257	98
	A04	199	205	267	267	306	306	118	118	328	334	236	257	98
	A05	201	205	269	269	300	312	118	118	328	331	236	236	98
	A06	201	209	267	267	294	306	118	118	331	334	236	257	98
	A07	199	205	0	0	306	318	118	118	331	334	236	257	98
	A08	201	205	269	269	300	324	118	118	328	334	236	257	98
	A09	201	205	267	267	306	306	118	118	328	334	236	257	98
	A10	201	209	269	269	300	324	118	118	328	331	236	257	98
	A11	199	209	267	267	306	318	118	118	328	328	257	257	98
	A12	201	209	267	267	294	318	118	118	328	331	257	257	98
S3	HhHU1	199	209	267	269	297	306	118	118	334	340	239	248	98
	HhHU2	199	209	267	267	297	306	118	118	334	340	242	245	98
	HhHU3	201	209	267	269	294	297	118	118	328	331	239	248	98
	HhHU4	201	209	267	267	294	297	118	118	328	334	239	242	98
	HhHU5	201	201	269	269	294	306	118	118	328	334	239	248	98
	HhHU6	201	201	267	269	297	306	118	118	334	340	242	245	98
	HhHU7	201	209	269	269	294	297	118	118	328	334	242	245	98
	HhHU9	199	209	269	269	294	306	118	118	331	340	239	242	98
	HhHU10	201	205	267	267	306	306	118	118	328	334	233	245	98
S4	MAN01	205	207	267	267	300	300	118	122	328	328	233	245	98
	MAN02	205	207	267	267	297	303	118	122	328	340	233	245	98
	MAN03	199	207	267	267	303	306	118	122	328	328	245	254	98
	MAN04	205	207	269	269	303	306	118	118	328	328	236	236	98
	MAN05	205	207	267	269	303	306	118	118	328	328	236	236	98
	MAN06	205	207	267	269	300	306	118	122	328	340	236	236	98
	MAN09	199	207	267	267	300	300	118	122	328	328	245	248	98
	MAN12	205	207	267	267	303	303	118	122	328	340	233	245	98
	MAN13	201	205	263	267	297	303	118	118	328	328	245	254	98
	MAN14	205	207	267	269	300	306	118	122	328	328	236	236	98
	MAN15	199	201	263	267	300	306	118	122	328	340	245	248	98
	MAN16	205	207	267	267	297	303	118	118	328	328	233	245	98
	MAN17	199	207	0	0	300	300	118	122	328	328	245	248	98
	MAN18	199	201	267	267	300	306	118	122	328	328	245	248	98

Table S5. Descriptive statistics of genetic diversity for all individuals and each broodstock (*cf* Table S1). Number of different alleles (Na), number of effective alleles, observed heterozygosity (H_o), expected heterozygosity (H_e), fixation index (F), and allele range of each locus are detailed.

	All					S1					S2					S3					S4				
Locus	Na	Ne	H_o	H_e	F	Na	Ne	H_o	H_e	F	Na	Ne	H_o	H_e	F	Na	Ne	H_o	H_e	F	Na	Ne	H_o	H_e	F
Anac_tag71527	6	4.051	0.829	0.753	−0.101	5	3.344	0.745	0.701	−0.062	4	3.789	1.000	0.736	−0.358	4	2.945	0.778	0.660	−0.178	4	3.322	1.000	0.699	−0.431
Anac_c15214	6	2.209	0.333	0.547	0.391	5	2.750	0.273	0.636	0.571	0	0.000	0.000	0.000	0.000	2	2.000	0.333	0.500	0.333	3	1.733	0.385	0.423	0.091
Anac_c31601	9	5.172	0.878	0.807	−0.088	6	4.440	0.957	0.775	−0.236	6	4.000	0.750	0.750	0.000	3	2.945	0.889	0.660	−0.346	4	3.532	0.714	0.717	0.004
Anac_c6179	3	1.672	0.500	0.402	−0.244	3	1.933	0.660	0.483	−0.366	1	1.000	0.000	0.000	0.000	1	1.000	0.000	0.000	0.000	2	1.849	0.714	0.459	−0.556
Anac_c16902	4	3.225	0.531	0.690	0.231	2	1.992	0.413	0.498	0.170	3	2.969	0.917	0.663	−0.382	4	3.447	1.000	0.710	−0.409	2	1.324	0.286	0.245	−0.167
Anac_c12159	9	6.203	0.757	0.839	0.097	7	3.249	0.771	0.692	−0.114	3	2.268	0.583	0.559	−0.043	5	4.263	1.000	0.765	−0.306	5	3.920	0.714	0.745	0.041
AoxD161	4	1.916	0.722	0.478	−0.509	4	1.892	0.682	0.472	−0.446	2	2.000	1.000	0.500	−1.000	3	2.160	0.889	0.537	−0.655	2	1.600	0.500	0.375	−0.333
An20	5	1.500	0.298	0.333	0.107	3	1.164	0.150	0.141	−0.062	2	2.000	1.000	0.500	−1.000	2	1.528	0.444	0.346	−0.286	2	1.237	0.214	0.191	−0.120
AoxD165	2	1.536	0.325	0.349	0.068	2	1.167	0.156	0.143	−0.084	2	1.946	0.500	0.486	−0.029	2	1.670	0.556	0.401	−0.385	2	1.849	0.571	0.459	−0.244
Afug51	3	2.209	0.714	0.547	−0.305	3	2.048	0.690	0.512	−0.350	2	1.280	0.250	0.219	−0.143	3	2.793	1.000	0.642	−0.558	2	2.000	1.000	0.500	−1.000
Ag22	3	2.097	0.451	0.523	0.137	2	1.693	0.319	0.409	0.221	3	2.667	0.333	0.625	0.467	3	1.742	0.556	0.426	−0.304	2	1.990	0.929	0.497	−0.867
Ag09	3	1.235	0.159	0.190	0.167	1	1.000	0.000	0.000	0	1	1.000	0.000	0.000	0.000	1	1.000	0.000	0.000	0.000	3	2.703	0.929	0.630	−0.474
Spl101	2	1.089	0.085	0.082	−0.045	2	1.022	0.021	0.021	−0.011	2	1.600	0.500	0.375	−0.333	1	1.000	0.000	0.000	0.000	1	1.000	0.000	0.000	0.000

References

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