

S2 Table. Genetic and geographic distances between localities separated by fronts and no-fronts for the 176 datasets. Species are classified by taxonomic group and the type of molecular marker used is specified. For each dataset we give the name of the localities and the genetic distance between them, as given in the References, for localities separated by a front and control localities without front between them. We have estimated approximate geographic distance between localities following the coastline with goggle earth.

Taxonomic group	Species	Marker	Front	FRONT			NO-FRONT			References
				Localities	Genetic distance	KM	Localities	Genetic distance	KM	
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	GS	Ca-TC	0.229	183	TC-R	0.285	163	[1]
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	AOF	R-U	0.142	253	TC-R	0.285	80	[1]
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	IC	U-D	0.156	190	TC-R	0.285	80	[1]
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	BF	F-Fa	0.174	259	TC-R	0.285	80	[1]
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	ADR	CR-SM	0.423	1289	TC-R	0.285	80	[1]
Angiosperm	<i>Cymodocea nodosa</i>	Microsatellites	AEG	AN-CY	0.183	977	TC-R	0.285	80	[1]
Porifera	<i>Crambe crambe</i>	COI	IC	GAT-TOS	0.140	705	TOS-BNY	0.915	131	[2]
Porifera	<i>Crambe crambe</i>	COI	BF	BAL-TOS	0.060	313	TOS-BNY	0.915	131	[2]
Porifera	<i>Ircinia fasciculata</i>	Microsatellites	AOF	GRA-ALI	0.083	335	CAL-BLA	0.016	215	[3]
Porifera	<i>Ircinia fasciculata</i>	Microsatellites	IC	ALI-CAL	0.020	315	CAL-BLA	0.016	215	[3]
Porifera	<i>Ircinia fasciculata</i>	Microsatellites	BF	CAB-CAL	0.034	260	CAL-BLA	0.016	215	[3]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Porifera	<i>Paraleucilla magna</i>	Microsatellites	GS	SGR-LHE	0.060	518	CGA-CPA	0.038	137	[4]
Porifera	<i>Paraleucilla magna</i>	Microsatellites	AOF	LHE-CGA	0.048	150	CGA-CPA	0.038	137	[4]
Porifera	<i>Paraleucilla magna</i>	Microsatellites	IC	BLN-CPA	0.057	536	CGA-CPA	0.038	137	[4]
Porifera	<i>Scopalina lophyropoda</i>	Microsatellites	IC	CG-MU	0.195	830	GA-MU	0.105	577	[5]
Porifera	<i>Spongia lamella</i>	Microsatellites	BF	Are-Cab	0.114	267	Are-Cr	0.047	130	[6]
Porifera	<i>Spongia officinalis</i>	Microsatellites	AEG	ECR-CYC	0.031	257	CYC-SPO	0.025	276	[7]
Cnidaria	<i>Astroides calycularis</i>	Microsatellites	AOF	GRA-PM	0.363	319	GRA-GAT	0.078	153	[8]
Cnidaria	<i>Aurelia aurita</i>	Allozymes	SC	BIZ-BB	0.025	577	SA-BB	0.003	217	[9]
Cnidaria	<i>Cladocora caespitosa</i>	Microsatellites	IC	IC-CP	0.028	273	LA-CI	0.027	110	[10]
Cnidaria	<i>Cladocora caespitosa</i>	Microsatellites	BF	IC-CG	0.051	267	LA-CI	0.027	110	[10]
Cnidaria	<i>Corallium rubrum</i>	Microsatellites	BF	FOR-XIC	0.072	237	FOR-LPP	0.034	83	[11]
Cnidaria	<i>Paramuricea clavata</i>	Microsatellites	AOF	TYL-CPS	0.115	501	MOU-ALT	0.124	473	[12]
Cnidaria	<i>Paramuricea clavata</i>	Microsatellites	IC	CPS-CLB	0.129	310	MOU-ALT	0.124	473	[12]
Mollusca	<i>Donax trunculus</i>	Microsatellites	GS	Don-Cala	0.018	250	Gan-SCRap	0.005	200	[13]
Mollusca	<i>Haliotis tuberculata</i>	Microsatellites	IC	CAR-BYS	0.031	624	AST-GAL	0.050	394	[14]
Mollusca	<i>Haliotis tuberculata</i>	Microsatellites	BF	BAL-BYS	0.037	273	AST-GAL	0.050	394	[14]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Mollusca	<i>Mytilus galloprovincialis</i>	Microsatellites	GS	ATdo-ALma	0.021	294	ALma-ALsa	0.018	95	[15]
Mollusca	<i>Mytilus galloprovincialis</i>	Microsatellites	AOF	ALsa-MEse	0.027	190	ALma-ALsa	0.018	95	[15]
Mollusca	<i>Mytilus galloprovincialis</i>	Microsatellites	IC	MEse-MEcu	0.017	170	ALma-ALsa	0.018	95	[15]
Mollusca	<i>Octopus vulgaris</i>	COI	SC	GI-ZA	0.664	460	LI-ZA	0.743	230	[17]
Mollusca	<i>Octopus vulgaris</i>	COI	ADR	CRZ-PCS	0.005	500	PCS-STM	0.091	300	[16]
Mollusca	<i>Patella rustica</i>	Allozymes	GS	EST-CAS	0.050	439	PES-SAV	0.020	211	[18]
Mollusca	<i>Patella rustica</i>	COI	ADR	PES-CRT	0.350	736	PES-SAV	0.260	214	[18]
Mollusca	<i>Patella ulyssiponensis</i>	COI	GS	EST-CAD	0.375	230	CAD-CAS	0.161	230	[19]
Mollusca	<i>Patella ulyssiponensis</i>	COI	SC	CGA-PSE	-0.075	330	CAD-CAS	0.161	230	[19]
Mollusca	<i>Pinna nobilis</i>	COI	SC	N-M	-0.084	383	S-B	-0.087	161	[20]
Mollusca	<i>Pinna nobilis</i>	COI	ADR	VEN-PAC	0.400	1171	S-B	-0.087	161	[20]
Mollusca	<i>Pinna nobilis</i>	COI	AEG	XI-PAC	0.400	1173	S-B	-0.087	161	[20]
Mollusca	<i>Ruditapes decussatus</i>	COI	SC	MZJ-SFAX	0.238	350	SFAX-BIBEN	0.116	170	[21]
Mollusca	<i>Sepia officinalis</i>	Microsatellites	GS	Rmar-Faro	0.010	603	Lisb-Faro	0.000	327	[22]
Mollusca	<i>Sepia officinalis</i>	Microsatellites	AOF	Rmar-Alic	0.069	335	Lisb-Faro	0.000	327	[22]
Mollusca	<i>Sepia officinalis</i>	Microsatellites	IC	Alic-Vgel	0.009	490	Lisb-Faro	0.000	327	[22]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Mollusca	<i>Sepia officinalis</i>	COI	BF	PAMA-VIGE	0.912	236	VIGE-SETE	0.095	392	[23]
Mollusca	<i>Sepia officinalis</i>	COI	SC	ALGI-GUHA	0.816	994	FARO-LISB	0.096	392	[23]
Mollusca	<i>Sepia officinalis</i>	COI	ADR	VENI-MALT	0.510	1419	FARO-LISB	0.096	392	[23]
Mollusca	<i>Sepia officinalis</i>	COI	AEG	AGNI-GUSI	0.474	753	FARO-LISB	0.096	392	[23]
Mollusca	<i>Solen marginatus</i>	Allozymes	SC	Biz-EIM	0.024	362	Biz-Rad	0.012	70	[24]
Crustacea	<i>Aristeomorpha foliacea</i>	ISSR	AEG	ION-AEG	0.033	650	PPA-ION	0.037	530	[25]
Crustacea	<i>Carcinus aestuarii</i>	Microsatellites	SC	CA-TA	0.028	1141	RA-FG	0.002	399	[26]
Crustacea	<i>Carcinus aestuarii</i>	Microsatellites	ADR	FG-TA	0.005	577	RA-FG	0.002	399	[26]
Crustacea	<i>Liocarcinus depurator</i>	COI	GS	CADIZ- MALAGA	0.061	218	VALENCIA- TARRAGONA	0.008	239	[27]
Crustacea	<i>Liocarcinus depurator</i>	COI	AOF	MALAGA- ALICANTE	0.015	499	VALENCIA- TARRAGONA	0.008	239	[27]
Crustacea	<i>Liocarcinus depurator</i>	COI	IC	ALICANTE- VALENCIA	0.124	191	VALENCIA- TARRAGONA	0.008	239	[27]
Crustacea	<i>Macropipus tuberculatus</i>	COI	GS	CADIZ- MALAGA	0.059	218	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Macropipus tuberculatus</i>	COI	AOF	MALAGA- ALICANTE	0.035	499	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Macropipus tuberculatus</i>	COI	IC	ALICANTE- VALENCIA	0.018	191	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Munida intermedia</i>	COI	GS	CADIZ- MALAGA	0.044	218	VALENCIA- TARRAGONA	0.024	239	[27]
Crustacea	<i>Munida intermedia</i>	COI	AOF	MALAGA- ALICANTE	0.014	499	VALENCIA- TARRAGONA	0.024	239	[27]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Crustacea	<i>Munida intermedia</i>	COI	IC	ALICANTE- VALENCIA	0.012	191	VALENCIA- TARRAGONA	0.024	239	[27]
Crustacea	<i>Pachygrapsus marmoratus</i>	COI	GS	MC-PS	0.003	705	VL-TA	0.080	211	[28]
Crustacea	<i>Pachygrapsus marmoratus</i>	COI	SC	BZ-SH	-0.036	250	VL-TA	0.080	211	[28]
Crustacea	<i>Pachygrapsus marmoratus</i>	COI	ADR	TG-PC	-0.102	200	VL-TA	0.080	211	[28]
Crustacea	<i>Pachygrapsus marmoratus</i>	COI	AEG	CH-OH	-0.030	590	VL-TA	0.080	211	[28]
Crustacea	<i>Pagurus alatus</i>	COI	GS	CADIZ- MALAGA	0.020	218	VALENCIA- TARRAGONA	0.000	239	[27]
Crustacea	<i>Pagurus alatus</i>	COI	AOF	MALAGA- ALICANTE	0.023	499	VALENCIA- TARRAGONA	0.000	239	[27]
Crustacea	<i>Pagurus alatus</i>	COI	IC	ALICANTE- VALENCIA	0.008	191	VALENCIA- TARRAGONA	0.000	239	[27]
Crustacea	<i>Pagurus excavatus</i>	COI	GS	CADIZ- MALAGA	0.030	218	VALENCIA- TARRAGONA	0.022	239	[27]
Crustacea	<i>Pagurus excavatus</i>	COI	AOF	MALAGA- ALICANTE	0.032	499	VALENCIA- TARRAGONA	0.022	239	[27]
Crustacea	<i>Pagurus excavatus</i>	COI	IC	ALICANTE- VALENCIA	0.022	191	VALENCIA- TARRAGONA	0.022	239	[27]
Crustacea	<i>Palaemon elegans</i>	16S-COI	GS	Gra-Cad	0.052	318	Gir-Ebro	0.021	252	[29]
Crustacea	<i>Palaemon elegans</i>	16S-COI	SC	Fus-Cal	-0.056	329	Gir-Ebro	0.021	252	[29]
Crustacea	<i>Palinurus elephas</i>	Microsatellites	BF	Columbretes- Mallorca	0.019	208	Mallorc- Menorca	0.001	223	[30]
Crustacea	<i>Parapenaeus longirostris</i>	COI	GS	CADIZ- MALAGA	0.023	218	VALENCIA- TARRAGONA	0.031	239	[27]
Crustacea	<i>Parapenaeus longirostris</i>	COI	AOF	MALAGA- ALICANTE	0.025	499	VALENCIA- TARRAGONA	0.031	239	[27]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Crustacea	<i>Parapenaeus longirostris</i>	COI	IC	ALICANTE- VALENCIA	0.051	191	VALENCIA- TARRAGONA	0.031	239	[27]
Crustacea	<i>Parapenaeus longirostris</i>	D-Loop	SC	TYR1-SS1	0.011	352	SS1-SS2	0.018	311	[31]
Crustacea	<i>Parapenaeus longirostris</i>	D-Loop	ADR	AD1-SS2	-0.019	749	SS1-SS2	0.018	311	[31]
Crustacea	<i>Parapenaeus longirostris</i>	D-Loop	AEG	AEG2-SS2	0.252	1221	SS1-SS2	0.018	311	[31]
Crustacea	<i>Penaeus kerathurus</i>	COI	SC	MON-KAL	0.452	275	MON-GAB	-0.021	144	[32]
Crustacea	<i>Plesionika heterocarpus</i>	COI	GS	CADIZ- MALAGA	0.016	218	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Plesionika heterocarpus</i>	COI	AOF	MALAGA- ALICANTE	0.036	499	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Plesionika heterocarpus</i>	COI	IC	ALICANTE- VALENCIA	0.018	191	VALENCIA- TARRAGONA	0.015	239	[27]
Crustacea	<i>Stenosoma nadejda</i>	COI	GS	2 y 4	0.872	236	4 y 5	0.603	49	[33]
Crustacea	<i>Stenosoma nadejda</i>	COI	AOF	5 y 7	0.793	129	4 y 5	0.603	49	[33]
Echinodermata	<i>Arbacia lixula</i>	COI	AOF	HER-CAR	0.002	135	CAR-PAL	-0.014	150	[34]
Echinodermata	<i>Arbacia lixula</i>	COI	IC	BEN-CLM	0.015	171	CAR-PAL	-0.014	150	[34]
Echinodermata	<i>Arbacia lixula</i>	COI	BF	CLM-FOR	-0.026	142	CAR-PAL	-0.014	150	[34]
Echinodermata	<i>Astropecten aranciacus</i>	Microsatellites	GS	LAH-FAR	-0.001	464	GAE-MUR	-0.004	389	[35]
Echinodermata	<i>Echinaster sepisotus</i>	Microsatellites	IC	Pal-Bla	-0.031	508	Car-Pal	-0.025	157	[36]
Echinodermata	<i>Echinaster sepisotus</i>	Microsatellites	BF	Cab-Bla	0.030	262	Car-Pal	-0.025	157	[36]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Echinodermata	<i>Echinaster sepiotus</i>	Microsatellites	SC	Tab-Mon	0.183	390	Car-Pal	-0.025	157	[36]
Echinodermata	<i>Echinaster sepiotus</i>	Microsatellites	ADR	Cro-Tao	0.115	874	Car-Pal	-0.025	157	[36]
Echinodermata	<i>Holothuria mammata</i>	COI	IC	GE-CP	-0.011	720	CP-ML	-0.010	415	[37]
Echinodermata	<i>Holothuria polii</i>	COI	IC	TH-GI	-0.020	510	MA-IS	-0.010	290	[39]
Echinodermata	<i>Holothuria polii</i>	COI	BF	GI-ML	0.004	255	MA-IS	-0.010	290	[39]
Echinodermata	<i>Holothuria polii</i>	Allozymes	SC	TAB-MON	0.050	390	MON-ZAR	0.049	280	[38]
Echinodermata	<i>Paracentrotus lividus</i>	COI	GS	CDr08-TFr08	-0.002	121	TS08-CL08	-0.002	200	[41]
Echinodermata	<i>Paracentrotus lividus</i>	COI	AOF	CBr08-TFr08	0.020	323	TS08-CL08	-0.002	200	[41]
Echinodermata	<i>Paracentrotus lividus</i>	COI	IC	CBr08-JVr08	0.028	355	TS08-CL08	-0.002	200	[41]
Echinodermata	<i>Paracentrotus lividus</i>	16S	BF	Cab-Tos	0.046	346	Col-Tos	0.003	281	[40]
Echinodermata	<i>Parastichopus regalis</i>	COI	AOF	AB-CG	0.047	205	CG-AL	-0.017	210	[42]
Echinodermata	<i>Parastichopus regalis</i>	COI	IC	AL-VAL	0.023	155	CG-AL	-0.017	210	[42]
Tunicata	<i>Pseudodistoma crucigaster</i>	COI	BF	Blanes-Cap Catalunya	0.929	191	Cadaqués- Blanes	0.416	61	[43]
Pisces	<i>Anguilla anguilla</i>	Microsatellites	GS	Oued Sebou- Moulonya	-0.001	544	Canet-Tour du Valat	0.001	184	[44]
Pisces	<i>Apogon imberbis</i>	Microsatellites	AOF	HE-CG	0.011	170	CG-MA	0.052	506	[45]
Pisces	<i>Apogon imberbis</i>	Microsatellites	IC	CG-BL	0.005	807	CG-MA	0.052	506	[45]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Pisces	<i>Apogon imberbis</i>	Microsatellites	BF	MA-BL	0.060	294	CG-MA	0.052	506	[45]
Pisces	<i>Atherina boyeri</i>	Microsatellites	ADR	Ku-Bt	0.030	330	Ca-Ob	0.050	214	[46]
Pisces	<i>Chromis chromis</i>	D-Loop	SC	Pro/Nap-Goz	0.090	638	Orb-Nap	0.000	384	[47]
Pisces	<i>Chromis chromis</i>	D-Loop	AEG	Don-Goz	0.740	1122	Orb-Nap	0.000	384	[47]
Pisces	<i>Coris julis</i>	tRNA thr	BF	TM-ML	0.003	298	SP-CI	0.003	264	[48]
Pisces	<i>Coris julis</i>	tRNA thr	SC	NA-AU	0.012	799	SP-CI	0.003	264	[48]
Pisces	<i>Coris julis</i>	tRNA thr	ADR	LE-SP	0.011	524	SP-CI	0.003	264	[48]
Pisces	<i>Coris julis</i>	tRNA thr	AEG	GA-AU	0.029	925	SP-CI	0.003	264	[48]
Pisces	<i>Coryphoblennius galerita</i>	D-Loop	IC	CG-Ba	-0.026	795	Cad-Luz	0.169	291	[49]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	GS	MKS1-MRBT	0.017	280	EGLV-FSET	0.017	680	[51]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	AOF	MRBT-EMUR	0.056	590	EGLV-FSET	0.017	680	[51]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	IC	EMUR-EGLV	0.017	320	EGLV-FSET	0.017	680	[51]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	SC	TISK-TMAH	0.010	431	EGLV-FSET	-0.003	680	[50]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	ADR	GMSL-ILDV	0.012	1305	EGLV-FSET	-0.003	680	[50]
Pisces	<i>Dicentrarchus labrax</i>	Microsatellites	AEG	YEGL-GTSK	0.050	1505	EGLV-FSET	-0.003	680	[50]
Pisces	<i>Diplodus sargus</i>	Allozymes	SC	BIZ-GA	0.026	460	MAH-GAB	0.035	170	[52]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Pisces	<i>Diplodus vulgaris</i>	Microsatellites	AOF	HE-CG	0.060	170	CG-MA	0.001	506	[45]
Pisces	<i>Diplodus vulgaris</i>	Microsatellites	IC	CG-BL	0.008	807	CG-MA	0.001	506	[45]
Pisces	<i>Diplodus vulgaris</i>	Microsatellites	BF	MA-BL	0.004	294	CG-MA	0.001	506	[45]
Pisces	<i>Engraulis encrasicolus</i>	RFLPs	GS	Mal-Olh	0.017	401	Lio-Liv	0.005	736	[53]
Pisces	<i>Engraulis encrasicolus</i>	RFLPs	SC	Liv-Pat	0.007	1651	Lio-Liv	0.005	736	[53]
Pisces	<i>Engraulis encrasicolus</i>	RFLPs	ADR	Chi-Pat	0.032	1210	Lio-Liv	0.005	736	[53]
Pisces	<i>Engraulis encrasicolus</i>	RFLPs	AEG	Sar-Pat	0.004	495	Lio-Liv	0.005	736	[53]
Pisces	<i>Epinephelus marginatus</i>	Microsatellites	IC	CGA-BCN	0.002	641	BCN-MED	0.004	130	[54]
Pisces	<i>Epinephelus marginatus</i>	Microsatellites	BF	MALL-BCN	-0.003	208	BCN-MED	0.004	130	[54]
Pisces	<i>Epinephelus marginatus</i>	Microsatellites	SC	ALG-TUN	0.013	717	BCN-MED	0.004	130	[54]
Pisces	<i>Epinephelus marginatus</i>	Microsatellites	AEG	GRE-TUN	0.004	1237	BCN-MED	0.004	130	[54]
Pisces	<i>Merluccius merluccius</i>	SNPs	GS	NPT-MAL	0.006	230	NAD-SAD	-0.007	377	[55]
Pisces	<i>Merluccius merluccius</i>	SNPs	SC	STY-MAB	0.013	751	NAD-SAD	-0.007	377	[55]
Pisces	<i>Merluccius merluccius</i>	SNPs	ADR	SAD-NWJ	0.007	596	NAD-SAD	-0.007	377	[55]
Pisces	<i>Merluccius merluccius</i>	SNPs	AEG	AEG-CYP	0.001	1009	NAD-SAD	-0.007	377	[55]
Pisces	<i>Mugil cephalus</i>	Microsatellites	IC	WMA-WME	0.040	355	WME-WMT	0.003	494	[56]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Pisces	<i>Mugil cephalus</i>	Microsatellites	SC	WMO-EMK	0.000	609	WME-WMT	0.003	494	[56]
Pisces	<i>Mugil cephalus</i>	Microsatellites	AEG	ASH-EMH	0.013	1322	WME-WMT	0.003	494	[56]
Pisces	<i>Mullus barbatus</i>	Microsatellites	AOF	FU-GA	-0.001	260	FU-AL	0.002	230	[60]
Pisces	<i>Mullus barbatus</i>	Microsatellites	IC	CG-BL	0.022	862	BL-IT	0.019	729	[57]
Pisces	<i>Mullus barbatus</i>	Microsatellites	SC	TIR5-SS3	0.008	320	IO1-SS3	0.006	100	[59]
Pisces	<i>Mullus barbatus</i>	Allozymes	AEG	4 y 5	0.001	1235	4 y 3	0.002	132	[58]
Pisces	<i>Mullus surmuletus</i>	Microsatellites	AOF	HE-CG	0.021	170	CG-MA	0.127	506	[45]
Pisces	<i>Mullus surmuletus</i>	Microsatellites	IC	CG-BL	0.001	862	CG-MA	0.127	506	[45]
Pisces	<i>Mullus surmuletus</i>	Microsatellites	BF	MA-BL	0.134	294	CG-MA	0.127	506	[45]
Pisces	<i>Oblada melanura</i>	Microsatellites	AOF	HE-CG	0.001	170	CG-MA	0.155	506	[45]
Pisces	<i>Oblada melanura</i>	Microsatellites	IC	CG-BL	0.052	807	CG-MA	0.155	506	[45]
Pisces	<i>Oblada melanura</i>	Microsatellites	BF	MA-BL	0.170	294	CG-MA	0.155	506	[45]
Pisces	<i>Sardina pilchardus</i>	EPIC	GS	Larache-Nador	0.034	416	Essaouira-Larache	0.002	319	[61]
Pisces	<i>Sardina pilchardus</i>	Allozymes	AOF	1 y 2	0.051	217	6 y 8	0.048	353	[62]
Pisces	<i>Sardina pilchardus</i>	Allozymes	IC	2 y 5	0.060	305	6 y 8	0.048	353	[62]
Pisces	<i>Sardina pilchardus</i>	Microsatellites	ADR	TR-IO	0.002	805	CH-SB	-0.001	346	[63]

Group	Species	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Pisces	<i>Scorpaena porcus</i>	Microsatellites	ADR	PC-TG	0.007	200	TG-TRE	0.008	272	[64]
Pisces	<i>Scyliorhinus canicula</i>	Microsatellites	AEG	ION-CRE	0.025	585	CRE-MYR	0.004	254	[65]
Pisces	<i>Seriola dumerilli</i>	D-loop	SC	TU-SR	0.031	530	TU-SA	-0.041	400	[66]
Pisces	<i>Serranus cabrilla</i>	Microsatellites	AOF	HE-CG	0.013	174	BL-CC	-0.002	120	[67]
Pisces	<i>Serranus cabrilla</i>	Microsatellites	IC	CP-C	0.044	386	BL-CC	-0.002	120	[67]
Pisces	<i>Serranus cabrilla</i>	Microsatellites	BF	MA-BL	0.008	221	BL-CC	-0.002	120	[67]
Pisces	<i>Serranus cabrilla</i>	Microsatellites	SC	CER-OT	0.000	1224	BL-CC	-0.002	120	[67]
Pisces	<i>Serranus cabrilla</i>	Microsatellites	AEG	GR-OT	0.002	888	BL-CC	-0.002	120	[67]
Pisces	<i>Solea vulgaris</i>	D-Loop	SC	SAL-SIC	0.043	789	LIV-SAL	-0.007	624	[68]
Pisces	<i>Solea vulgaris</i>	D-Loop	ADR	NER-GAL	0.024	437	LIV-SAL	-0.007	624	[68]
Pisces	<i>Solea vulgaris</i>	Allozymes	AEG	GR-EG	0.013	1308	E1-M1	0.007	458	[69]
Pisces	<i>Squalus blainville</i>	COI	SC	SIC-MAL	0.029	300	MAL-ION	-0.065	590	[70]
Pisces	<i>Squalus blainville</i>	COI	AEG	CRE-CYC	-0.278	292	MAL-ION	-0.065	590	[70]
Pisces	<i>Symphodus tinca</i>	Microsatellites	AOF	HE-CG	0.033	170	CG-MA	0.001	506	[45]
Pisces	<i>Symphodus tinca</i>	Microsatellites	IC	CG-BL	0.001	807	CG-MA	0.001	506	[45]
Pisces	<i>Symphodus tinca</i>	Microsatellites	BF	MA-BL	0.002	294	CG-MA	0.001	506	[45]

Group	<i>Species</i>	Marker	Front	Localities	Genetic	KM	Localities	Genetic	KM	References
Pisces	<i>Thalassoma pavo</i>	D-Loop	SC	Ust-Lin	0.000	416	Gen-Poz	0.000	736	[71]
Pisces	<i>Thalassoma pavo</i>	D-Loop	AEG	Kea-lin	0.571	1168	Gen-Poz	0.000	736	[71]
Pisces	<i>Tripterygion delaisi</i>	Microsatellites	AOF	TA-GA	0.046	293	CO-BL	0.029	263	[72]
Pisces	<i>Tripterygion delaisi</i>	Microsatellites	IC	PA-CO	0.036	386	CO-BL	0.029	263	[72]
Pisces	<i>Tripterygion delaisi</i>	Microsatellites	BF	CO-FO	0.010	167	CO-BL	0.029	263	[72]

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