## Spatial gradients of introgressed ancestry reveal cryptic connectivity patterns in a high gene flow marine fish

Tony Robinet  $(1)^*$ , Valérie Roussel (2), Karine Cheze (1) and Pierre-Alexandre Gagnaire (3)

------

**Figure S1**. Map of the 21 sampling localities of the 'main dataset', named by their locality code and represented with ICES fisheries statistical areas.

**Figure S2**. Map of the 31 sampling localities of the 'refined dataset', containing 761 individuals with a minimal number of 8 individuals per ICES rectangle.

**Figure S3. a:** Observed sample heterozygosity (Hobs) as a function of latitude for the 21 localities of the main dataset. **b:** Relationship between SNP polymorphism rate per locality and the number of specimens (*N*) in each locality, fitted by a logistic model.

**Figure S4.** Pairwise genetic differentiation represented as  $F_{sT}/(1-F_{st})$  as a function of the distance between the localities (main dataset of 21 localities).

**Figure S5. a:** Geographic pattern of mean MED ancestry proportions along the northeast ATL coastlines, from south to north, using the 'refined dataset' of 31 localities. **b:** Geographic pattern of mean MAF along the northeast ATL coastlines displayed by between-lineages outlier loci.

**Figure S6.** Outlier loci detection by LOSITAN within the Atlantic (**a**), and between ATL and MED lineages (**b**). Outlier loci detection by BAYESCAN within the Atlantic (**c**), and between ATL and MED lineages (**d**).

**Figure S7.** Geographic pattern of mean MAF for different subsets of loci, defined according to their allele frequency differences between ATL and MED lineages (ATL-MED  $\Delta$ -*p*), as a function of the distance to SINE (km). The analysis is based on the main dataset.

Table S1. Regions, localities and codes, ICES rectangles, dates of specimen collection.

**Table S2**. Summary of genetic variation at the 1012 retained loci, as grouped in the main dataset (21 localities from 7 regions).

Table S3. Pairwise FST for the 1012 loci (upper triangle) between pairs of 21 ATL localities.

**Table S4.** Pairwise FST calculated between pairs of 21 ATL localities from 980 non-outlier SNPs within the ATL (lower triangle) and from 32 within-ATL outlier SNPs (upper triangle).

Table S5. Fst p-values corresponding to the differentiation indices provided in Table S5.

**Table S6.** Pairwise Fst values at the 1012 retained loci (upper triangle), for each pair among the 7 Atlantic regions.

**Table S7.** Mantel test between matrices of geographic distances (nautic milles) and genetic distances [Fst/(1-Fst)] between pairs of localities within 3 groups of sampling locations that differ in their average ancestry proportions (see Figure 3).

<sup>1</sup> Biologie des Organismes et Ecosystèmes Aquatiques (BOREA) ; Muséum National d'Histoire Naturelle, CNRS, IRD, SU, UCN, UA ; Station marine de Concarneau, Quai de la Croix, 29900 Concarneau, France

<sup>2</sup> Université de Bretagne Occidentale, Institut GéoArchi EA7462, 6 avenue Le Gorgeu, 29100 Brest, France

<sup>3</sup> ISEM, Univ Montpellier, CNRS, EPHE, IRD, Montpellier, France.

<sup>\*</sup> corresponding author : <u>robinet@mnhn.fr</u>



**Figure S1**. Map of the 21 sampling localities of the 'main dataset', named by their locality code and represented with ICES fisheries statistical areas. See detailed information corresponding to each locality code in Table S1.



**Figure S2**. Map of the 31 sampling localities of the 'refined dataset', containing 761 individuals with a minimal number of 8 individuals per ICES rectangle. Empty rectangles represent samples from the main dataset that we excluded for the spatially refined analysis due to limited sample size.



**Figure S3. a:** Observed sample heterozygosity (Hobs) as a function of latitude for the 21 localities of the main dataset. The blue line represents the fitted linear model (slope p<0.001,  $r^2$ =0.66, grey shade 0.95 confidence interval). **b:** Relationship between SNP polymorphism rate per locality and the number of specimens (*N*) in each locality, fitted by a logistic model (orange line: Polymorphism = 0.05644 ln(*N*) + 0.74216, *p*<0.001 for each coefficient).



**Figure S4.** Pairwise genetic differentiation represented as  $F_{ST}/(1-F_{st})$  as a function of the distance between the localities (main dataset of 21 localities). **a**: Positive relationships between pairwise genetic and geographic distances illustrated for non-outlier SNPs (red line, slope p<0.001, r<sup>2</sup>=0.074) and outlier SNPs that were detected at the within-Atlantic scale with Lositan (green line, slope p<0.001, r<sup>2</sup>=0.089). **b**: Relationships between pairwise genetic and geographic distances illustrated for non-outlier SNPs separately for localities south to GONB (gold line, not significant) and north to GONB (purple line, slope p<0.001, r<sup>2</sup>=0.236). **c**: Same as b but for outlier SNPs that were detected at the within-Atlantic scale with Lositan. South to GONB (gold line, slope p<0.001, r<sup>2</sup>=0.083), north to GONB (purple line, slope p<0.001, r<sup>2</sup>=0.245).



**Figure S5. a:** Geographic pattern of mean MED ancestry proportions along the northeast ATL coastlines, from south (the locality SINE was not represented) to north (NEUK), using the 'refined dataset' of 31 localities. **b:** Geographic pattern of mean MAF along the northeast ATL coastlines displayed by between-lineages outlier loci. Blue line: localities from southern Portugal to PBRE; green line: localities from CRNW to SGEO in the Irish Sea (Celtic Sea-Irish Sea branch); red line: localities from GONB to NEUK in the North Sea (Channel-North Sea branch).



**Figure S6.** Outlier loci detection by LOSITAN within the Atlantic (**a**), and between ATL and MED lineages (**b**). Loci below and above the red line were classified as neutral and outlier loci, respectively. Outlier loci detection by BAYESCAN within the Atlantic (**c**), and between ATL and MED lineages (**d**). Loci left and right to the *q*-value threshold were classified as neutral and outlier loci, respectively.



**Figure S7.** Geographic pattern of mean MAF for different subsets of loci, defined according to their allele frequency differences between ATL and MED lineages (ATL-MED  $\Delta$ -*p*), as a function of the distance to SINE (km). The analysis is based on the main dataset.

Regions	Localities	Codes	ICES rectangles	Dates	Ng	Na
North Sea	North-East UK	NEUK	37E9, 38E9, 37E8	Win-Sum 2015	11	11
	East Anglia	ANGL	33F1	Sum 2014	30	29
	Dover-Boulogne	DOVE	30F1, 31F1,31F2	Win-Sum 2014, Win 2015	61	61
	Seine Bay	BSEI	28E8, 28E9	Win 2014	40	40
Channel	Normand-breton Gulf	GONB	26E7	Sum 2014	25	25
	Isle of Wight	ISWI	30E8, 30E9	Sum 2014	30	30
	English Cornwall	CRNW	28E3, 28E5, 29E4, 29E5	Win-Sum 2014	72	71
1.1.0	St George's Channel	SGEO	35E5	Sum 2013	22	21
Irish Sea	Wales	WALE	31E5, 32E5,	Sum 2013,	42	41
			32E6	Win-Sum 2014		
Calific Car	South-Ireland	IRIS	28E1	Win 2014	25	24
Celuc Sea	Britain's cap	PBRE	25E4, 25E5, 26E5	Win-Sum 2014	56	50
North Biscay	Loire Estuary	LOIR	22E6, 22E7, 23E6, 23E7,24E6	Sum 2012, Sum 2013, Win-Sum 2014, Win 2015	111	108
	Charentes	CHAR	20E7, 20E8, 21E7, 21E8	Sum 2013, Sum 2014, Win 2015	34	34
South Biscay	Aquitaine	AQUI	16E8, 17E8, 18E8	Sum 2013, Win-Sum 2014	81	79
	Cantabria	CANT	16E6	Sum 2014	21	21
	Asturias	ASTU	16E3, 16E4	Win 2014, Win 2015	39	37
	A Coruña	CORU	15E1	Win 2014	30	29
Portu gal	Vigo-Pontevedra	VIGO	13E1	Win 2014	30	30
Fortugai	Porto	PORT	10E1	Win 2015	30	30
	Peniche	PENI	7E0	Win 2015	29	29
	Sine	SINE	4E1	Win 2015	27	27
Total	21 localities		44 rectangles		846	827

**Table S1**. Regions, localities and codes, ICES rectangles (statistical squares for international management of fisheries), dates (season and year) of specimen collection. Ng: no. of specimens genotyped; Na: no. of specimens kept after quality filtering. Win: winter season (from Dec 1<sup>st</sup> of previous year to May 31<sup>st</sup> of the year); Sum: summer (from Jun 1st to Nov 30<sup>th</sup> of the year).

Region	Pool	Ν	H <sub>obs</sub>	Hexp	F <sub>IS</sub>	DAF	Polym.
North sea	ANGL	29	0.2464	0.2868	0.1409	0.8265	0.9259
	NEUK	11	0.2472	0.2952	0.1626	0.8200	0.8310
Channel	BSEI	40	0.2488	0.2881	0.1365	0.8255	0.9219
	CRNW	71	0.2483	0.2850	0.1287	0.8279	0.9615
	DOVE	61	0.2473	0.2835	0.1276	0.8290	0.9555
	GONB	25	0.2534	0.2932	0.1356	0.8216	0.9200
	ISWI	30	0.2483	0.2866	0.1337	0.8267	0.9259
Irish sea	SGEO	21	0.2472	0.2895	0.1464	0.8244	0.8874
	WALE	41	0.2531	0.2905	0.1286	0.8237	0.9536
Celtic sea	IRIS	24	0.2380	0.2785	0.1454	0.8328	0.8874
	PBRE	50	0.2547	0.2929	0.1307	0.8218	0.9625
North Biscay	CHAR	34	0.2535	0.2904	0.1270	0.8238	0.9526
2	LOIR	108	0.2604	0.2958	0.1196	0.8195	0.9951
South Biscay	AQUI	79	0.2581	0.2929	0.1190	0.8218	0.9872
2	ASTU	37	0.2562	0.2936	0.1274	0.8212	0.9536
	CANT	21	0.2562	0.2945	0.1301	0.8205	0.9249
Portugal	CORU	29	0.2648	0.3026	0.1251	0.8141	0.9654
0	VIGO	30	0.2569	0.2944	0.1271	0.8206	0.9565
	PENI	29	0.2639	0.3036	0.1309	0.8134	0.9634
	PORT	30	0.2632	0.3033	0.1322	0.8136	0.9664
	SINE	27	0.2850	0.3175	0.1024	0.8021	0.9852

**Table S2**. Summary of genetic variation at the 1012 retained loci, as grouped in the main dataset (21 localities from 7 regions). N: Number of genotyped individuals kept after filtering; Hobs: mean observed heterozygosity; Hexp: mean expected heterozygosity; FIS index: (Hexp-Hobs)/Hexp; DAF: mean frequency of major alleles; Polym.: proportion of polymorphic SNPs in each locality.

## Following pages:

**Table S3.** Pairwise FST for the 1012 loci (upper triangle) between pairs of 21 ATL localities. Significant FST appear in bold black (p<0.05, p-values are given in lower triangle), pale grey values are not significant. Locality codes are provided in Table S1.

**Table S4.** Pairwise FST calculated between pairs of 21 ATL localities from 980 non-outlier SNPs within the ATL (lower triangle) and from 32 within-ATL outlier SNPs (upper triangle). Bold black numbers indicate significant FST values (p<0.05, p-values are given in Table S7), pale grey values indicate non-significant Fst values. Locality codes are provided in Table S1.

**Table S5.** Fst p-values corresponding to the differentiation indices provided in Table S5 between pairs of 21 ATL localities, obtained using 5000 permutations for the 980 non-outlier SNPs within the ATL (lower triangle) and the 32 within-ATL outlier SNPs (upper triangle). Bold black numbers indicate significant Fst values (p<0.05), and pale grey non-significant Fst values. Locality codes are provided in Table S1.

	NEUK	ANGL	DOVE	SGEO	WALE	IRIS	IWSI	BSEI	CRNW	GONB	PBRE	LOIR	CHAR	AQUI	CANT	ASTU	CORU	VIGO	Ŭ	PORT	PORT PENI
NEUK		0.0112	0,4853	0.0172	0.0100	0.0094	0.0232	0.0600	0.5995	0.0070	0.0084	0.9328	0.0328	0.5273	0.0010	0.0440	0.0148		0.0126	0.0126 0.0128	0.0126 0.0128 0.013
ANGL	0.0132		0.0062	0.0110	0.0081	0.0098	0.0090	0.0083	0.0056	0.0102	0.0069	0.0042	0.0095	0.0053	0.0109	0.0077	0.0094	0	.0088	.0088 0.0091	.0088 0.0091 0.009
DOVE	0.0074	0.4259		0.0060	0.0049	0.0060	0.0061	0.6563	0.8816	0,0056	0.0048	0.0029	0.5753	0.8142	0.4891	0.7103	0.5773	0	0050	0050 0.0056	0050 0.0056 0.005
SGEO	0.0012	0.0548	0.7305		0.0079	0.0758	0.2016	0,4025	0.7802	0.0698	0,7285	0.9788	0.2078	0.7750	0.0256	0.3853	0.1136	0	9600	0096 0.3087	0096 0.3087 0.211
WALE	0.3903	0.4545	0.9156	0.8388		0.5727	0.6985	0.7313	0.9250	0.6153	0.9300	0.9968	0.6555	0.8838	0.5155	0.6939	0.6571	0.9	809	608 0.8694	608 0.8694 0.836
IRIS	0.0146	0.0862	0.6119	0.0113	0.0085		0.0097	0.2683	0.6647	0.0880	0.0069	0.0040	0.1106	0.6127	0.0358	0.2370	0.1816	0.0	092	092 0.0100	092 0.0100 0.009
IWSI	0.0129	0.1546	0.6119	0.0099	0.0077	0.1448		0.3173	0.6909	0.0546	0.0066	0.0041	0.2010	0.6423	0.1254	0,3399	0.1834	0.00	85	0.0086	0.0086 0.009
BSEI	0.0105	0.1700	0.0050	0.0080	0.0060	0.0082	0.0079		0.0045	0.0079	0.0060	0.0035	0.0070	0.7087	0.0087	0.4167	0.0077	0.00	772	0.0071	0.0071 0.007
CRNW	0.0064	0.4889	0.0038	0.0054	0.0045	0.0055	0.0053	0.7415		0.0055	0.0041	0.0027	0.7415	0.9180	0.5767	0.8000	0.6311	0.00	047	0.0047	0.0047 0.005
GONB	0.0148	0.0540	0.6393	0.0114	0.0083	0.0102	0.0109	0.2977	0.6511		0.0069	0.0039	0.1974	0.6307	0.0240	0.2468	0.1358	0.00	91	91 0.0099	91 0.0099 0.010
PBRE	0.4107	0.4365	0.8568	0.0072	0.0056	0.6145	0,6681	0.5995	0.9038	0.5999		0.9950	0.5629	0.9030	0,4683	0.7023	0.5983	0.00	62	62 0.0064	62 0.0064 0.006
LOIR	0.0044	0.8042	0.9942	0.0038	0.0035	0.9322	0.9252	0.9480	0.9950	0.9256	0.0032		0.9476	0.9996	0.9118	0.9712	0.9226	0.00	33	0,0036	33 0.0036 0.003
CHAR	0.0116	0.0754	0.0056	0.0094	0.0073	0.0096	0.0089	0.3519	0.0046	0.0087	0.0065	0,0035		0.6555	0.0982	0.3391	0.0086	0.00	83	0.0084	0.0084 0.008
AQUI	0.0061	0.4331	0.0039	0.0049	0.0043	0.0052	0.0051	0.0042	0.0034	0.0050	0.0037	0.0025	0.0047		0.0051	0.0040	0.0047	0.00	042	0.0044	042 0.0044 0.004
CANT	0.0158	0.0234	0.0062	0.0127	0.0081	0.0110	0.0096	0.1560	0.0054	0.0115	0,0072	0.0037	0.0094	0.5379		0.2024	0.0094	0.00	83	93 0.0095	93 0.0095 0.009
ASTU	0.0108	0.2038	0.0047	0.0081	0.0061	0.0084	0.0076	0,0060	0.0040	0,0082	0.0051	0.0033	0.0070	0.7231	0.0082		0.0071	0,00	70	70 0.0073	70 0.0073 0.007
CORU	0.0132	0.0892	0.0057	0.0103	0.0072	0.0091	0,0091	0.2855	0.0053	0.0093	0.0064	0.0038	0.1838	0.6493	0.1028	0.3405		0,00	85	85 0.0087	85 0.0087 0.009
VIGO	0.0734	0.2825	0.8838	0.4385	0.0067	0.3619	0.5147	0.5685	0.8988	0.3663	0.8748	0.9976	0.4405	0.8764	0.2713	0.5639	0.4171			0.6529	0.6529 0.593
PORT	0.0372	0.1892	0.7682	0.0097	0.0073	0.1672	0.3989	0.5219	0.8748	0.1720	0.7948	0.9830	0.3401	0.8264	0.1706	0.4655	0.3083	0.008	4	34	0.474
PENI	0.0218	0.1678	0.7123	0.0102	0.0074	0.2274	0.2426	0.4653	0,7998	0.0928	0.7802	0.9770	0.2989	0.7768	0.1278	0.4047	0.1682	0.008	ő	36 0,0086	36 0,0086
SINE	0.0312	0.1916	0.7323	0.1736	0.0078	0.1468	0.3187	0.3787	0.7924	0.1408	0.7902	0.9648	0.2280	0.7524	0.0960	0.4551	0.2330	0.00	2	91 0.3259	91 0.3259 0.301

Table S3.

Fst	
_827ATL	
32outliers	
980neutrals	
_21sreg	

0.0078 0.0042 0.0085 0.0047 0.0111 0.0082 0.0092 0.0175 0.0115 0.0079 0.0041 0.0083 0.0047 0.0115 0.0086 0.0092 <b>0.0228</b> 0.0095 C	0.0078 0.0042 0.0085 0.0047 0.0111 0.0082 0.0092 0.0175 0.0079 0.0041 0.0083 0.0047 0.0115 0.0086 0.0092 <b>0.0228</b>	0.0078 0.0042 0.0085 0.0047 0.0111 0.0082 0.0092 0.0079 0.0041 0.0083 0.0047 0.0115 0.0086 0.0092	0.0078 0.0042 0.0085 0.0047 0.0111 0.0082 0.0079 0.0041 0.0083 0.0047 0.0115 0.0086	0.0078 0.0042 0.0085 0.0047 0.0111 0.0079 0.0041 0.0083 0.0047 0.0115	0.0078 0.0042 0.0085 0.0047 0.0079 0.0041 0.0083 0.0047	0.0078 0.0042 0.0085 0.0079 0.0041 0.0083	0.0078 0.0042 0.0079 0.0041	0.0078		0.0096	0.0052	0.0085	0.0092	0.0093	0.0115	0.0202	0.0059	0.0095	0.0141	PENI
0.0092 (	0.0092	~	0.0092	0.0081	0.0105	0.0051	0.0083	0.0040	0.0076	0.0091	0.0052	0.0084	0.0093	0.0106	0.0112	0.0120	0.0064	0.0090	0.0142	PORT
0.0091 (	01	Ch	0.008	0.0084	0.0105	0.0048	0.0087	0.0040	0.0084	0.0093	0.0054	0.0085	0.0088	0.0096	0.0112	0.0100	0.0062	0.0090	0.0140	VIGO
0.0087 0.0174 (	0.0087			0.0082	0.0108	0.0048	0.0090	0.0046	0.0239	0.0121	0.0028	0.0081	0.0115	0.0048	0.0093	0.0284	0.0071	0.0082	0.0378	CORU
83 0.0120 0.0131 (	83 0.0120	8	0.00		0.0047	0.0050	0.0047	0.0037	0.0115	0.0049	0.0052	0.0136	0.0098	0.0098	0.0078	0.0236	0.0039	0.0081	0.0164	ASTU
29 0.0115 0.0079 (	29 0.0115	29	0.01	0.0095		0.0055	0.0048	0.0014	0.0093	0.0071	0.0037	0.0086	0.0094	0.0149	0.0058	0.0228	0.0031	0.0110	0.0231	CANT
73 0.0066 0.0076 (	73 0.0066	73	0.00	0.0077	0.0041		0.0056	0.0027	0.0213	0.0072	0.0036	0.0067	0.0047	0.0077	0.0043	0.0153	0.0052	0.0044	0.0168	AQUI
5 0.0121 0.0114 (	5 0.0121	(71	0.008	0.0076	0.0094	0.0048		0.0031	0.0132	0.0109	0.0045	0.0074	0.0081	0.0117	0.0108	0.0256	0.0046	0.0086	0.0193	CHAR
17 0.0050 0.0040 (	0.0050	7	0.003	0.0036	0.0040	0.0027	0.0036		0.0154	0.0035	0.0030	0.0036	0.0039	0.0039	0.0033	0.0093	0.0030	0.0041	0.0111	LOIR
4 0.0237 0.0183 (	4 0.0237	E	0.008	0.0069	0.0079	0.0053	0.0071	0.0043		0.0075	0.0055	0.0065	0.0080	0.0084	0.0175	0.0230	0.0057	0.0085	0.0088	PBRE
0.0079 0.0118 <b>(</b>	0.0079	X	0.009	0.0090	0.0113	0.0046	0.0084	0.0031	0.0170		0.0048	0.0085	0.0116	0.0139	0.0066	0.0281	0.0059	0.0088	0.0333	GONB
1 0.0088 0.0081 (	1 0.0088	jus.	0.005	0.0052	0.0059	0.0033	0.0049	0.0026	0.0151	0.0062		0.0052	0.0052	0.0033	0.0047	0.0124	0.0041	0.0052	0.0129	CRNW
3 0.0153 0.0165 (	3 0.0153	ຜ	0.015	0.0078	0.0098	0.0050	0.0143	0.0067	0.0170	0.0140	0.0080		0.0101	0.0136	0.0060	0.0194	0.0127	0.0086	0.0223	BSEI
2 0.0113 0.0185 (	0.0113	2	0.009	0.0080	0.0106	0.0046	0.0085	0.0049	0.0201	0.0096	0.0052	0.0078		0.0097	0.0090	0.0354	0.0060	0.0088	0.0301	ISMI
6 0.0141 0.0168 (	6 0.0141	ő	0.009	0.0084	0.0118	0.0047	0.0097	0.0048	0.0199	0.0105	0.0056	0.0091	0.0136		0.0101	0.0228	0.0063	0.0094	0.0410	IRIS
72 0.0072 0.0071 (	0.0072	2	0.007	0.0070	0.0082	0.0041	0.0067	0.0034	0.0061	0.0069	0.0039	0.0066	0.0071	0.0084		0.0085	0.0047	0.0075	0.0108	WALE
8 0.0150 0.0097 (	0.0150	8	0.009	0.0096	0.0134	0.0050	0.0101	0.0042	0.0088	0.0103	0.0057	0.0094	0.0104	0.0110	0.0227		0.0066	0.0100	0.0184	SGEO
58 0.0103 0.0081 (	58 0.0103	8	0.00	0.0056	0.0064	0.0036	0.0056	0.0018	0.0162	0.0059	0.0037	0.0052	0.0099	0.0075	0.0066	0.0166		0.0056	0.0153	DOVE
32 0.0069 0.0186 (	32 0.0069	32	0.00	0.0118	0.0137	0.0073	0.0119	0.0050	0.0270	0.0128	0.0044	0.0133	0.0084	0.0068	0.0085	0.0302	0.0096		0.0418	ANGL
19 0.0410 0.0347 (	0.0410	19	0.014	0.0121	0.0160	0.0066	0.0125	0.0048	0.0081	0.0162	0.0067	0.0104	0.0147	0.0175	0.0230	0.0594	0.0079	0.0160		NEUK
U VIGO PORT	U VIGO	C	COR	ASTU	CANT	AQUI	CHAR	LOIR	PBRE	GONB	CRNW	BSEI	IWSI	IRIS	WALE	SGEO	DOVE	ANGL	NEUK	Fst

Table S4.

## Fst\_827ATL\_32outliers\_980neutrals\_21sreg

CINE 0.0046 0.24	PENI 0.0166 0.13	PORT 0.0168 0.2	VIGO 0.0334 0.2	CORU 0.0004 0.2	ASTU 0.0310 0.10	CANT 0.0044 0.0	AQUI 0.0226 0.6	CHAR 0.0154 0.1	LOIR 0.1982 0.8;	PBRE 0.3413 0.2	GONB 0.0006 0.1	CRNW 0.1028 0.54	BSEI 0.0054 0.1	ISWI 0.0010 0.1	IRIS 0.0004 0.10	WALE 0.2885 0.53	SGEO 0.0002 0.1	DOVE 0.0518 0.5(	ANGL 0.0002	NEUK 0.0	p-value NEUK AN
402 0.7538	390 0.7117	114 0.6821	689 0.7594	042 0.3955	690 0.7584	230 0.8798	579 0.5589	518 0.6813	206 0.9908	370 0.7247	616 0.6029	475 0.7694	386 0.0778	768 0.6279	070 0.5667	319 0.9324	134 0.6795	045	0.1774	024 0.3917	IGL DOVE
0.2250	0.0206	0.2138 (	0.3457	0.0022	0.0040	0.0082	0.0508	0.0040	0.3629 (	0.0102	0.0022	0.1344 (	0.0180	0.0004	0.0084	0.7626	-	0.0436	0.0016	0.0002	SGEO
0.0896 (	0.3423 (	0.3771 (	0.4499 (	0.3895 (	0.4715 (	0.6925	0.8190 (	0.2819 (	0.9668 (	0.0866 (	0.6853 (	0.8342 (	0.6585 (	0.4823 (	0.3885		0.0322	0.6691 (	0.3931 (	0.0222 (	WALE
0.1374 0	0.2330 0	0.0984 0	0.2877 0	0.6793 0	0.1864 0	0.0466	0.3241 0	0.1210 0	0.9296 0	0.3803 0	0.0774 0	0.8868 0	0.0632 0	0.1360		0.5839 0	0.0872 0	0.3977 0	0.3969 0	0.0002 0	IRIS
0.3949 0	0.2625 (	0.2651 (	0.4325 (	0.1368 (	0.1884 (	0.2202 (	1.6483 (	0.3147 (	0.9424 (	0.4627 (	0.1416 (	0.6503 (	0.1792		0.0802 (	0.7672 (	0.1372 (	0.2150 (	0.2665	0.0068	IWSI
0.3121	0.2767	0.3069	0.3591	0.2216	0.0460	0.1482	0.3385	0.2929	0.9204	0.5245	0.2126	0.5809		0.3203	0.1318	0.6455	0.1870	0.5961	0.0438	0.0558	BSEI
0.8422	0.7754	0.7984	0.8254	0.9118	0.5587	0.7932	0.7860	0.6847	0.9914	0.7195	0.7700		0.2825	0.7125	0.6379	0.9664	0.7435	0.9110	0.6603	0.5615	CRNW
0.2597	0.1856	0.2739	0.3401	0.1052	0.6197	0.3857	0.3477	0.1386	0.9708	0.5199		0.5153	0.0538	0.1518	0.0668	0.7512	0.1390	0.5559	0.0724	0.0014	GONB
0.6479	0.6001	0.6483	0.6011	0.0042	0.1272	0.2527	0.0060	0.0876	0.0574		0.0362	0.0554	0.0260	0.0152	0.0132	0.9108	0.4519	0.0434	0.0020	0.4475	PBRE
0.9752	0.9574	0.9688	0.9826	0.7469	0.8226	0.9972	0.9252	0.9072		0.9452	0.9170	0.9540	0.4587	0.7343	0.7634	0.9986	0.9692	0.9918	0.6201	0.8798	LOIR
0.4095	0.3073	0.3635	0.3681	0.1490	0.5987	0.6303	0.4785		0.9388	0.5069	0.2408	0.6941	0.0448	0.2587	0.1092	0.6885	0.1372	0.5785	0.0762	0.0168	CHAR
0.8198	0.7754	0.7277	0.8140	0.6451	0.5445	0.4891		0.6295	0.9926	0.6605	0.7363	0.9418	0.5509	0.7516	0.7233	0.9092	0.7554	0.8912	0.2595	0.4909	AQUI
0.0582	0.0508	0.0826	0.1354	0.0320	0.6055		0.6881	0.0984	0.8746	0.3687	0.0314	0.5161	0.1596	0.0440	0.0208	0.5051	0.0162	0.4625	0.0400	0.0022	CANT
0.3175	0.3137	0.3479	0.3821	0.2144		0.0842	0.2330	0.2669	0.9108	0.4791	0.1602	0.5819	0.2160	0.2891	0.2234	0.5981	0.1670	0.5277	0.0670	0.0226	ASTU
0.2779	0.2144	0.2412	0.4243		0.2452	0.0684	0.3011	0.2370	0.9352	0.3249	0.1378	0.6701	0.0344	0.1738	0.1214	0.6733	0.1754	0.5781	0.8154	0.0062	CORU
0.0228	0.0688	0.4703		0.3815	0.1492	0.1970	0.5369	0.1682	0.8386	0.0162	0.4995	0.3999	0.0748	0.2531	0.1228	0.9396	0.1312	0.2889	0.5015	0.0002	VIGO
0.3857	0.2278		0.5049	0.0306	0.0880	0.3865	0.3807	0.1628	0.8880	0.0338	0.1710	0.4007	0.0370	0.0252	0.0442	0.8796	0.2963	0.4083	0.0178	0.0010	PORT
0.4051		0.3857	0.5295	0.0484	0.0932	0.0658	0.4387	0.0848	0.6363	0.0340	0.0326	0.4213	0.0218	0.0264	0.2288	0.8470	0.1416	0.4591	0.0282	0.0010	PENI
	0.0220	0.1100	0.5627	0.0004	0.0036	0.0376	0.0848	0.0024	0.3333	0.0122	0.0006	0.0610	0.0886	0.0010	0.0008	0.8780	0.2324	0.0486	0.0004	0.0002	SINE

Table S5.

p-val / Fst	North Sea	Channel	Irish Sea	Celtic Sea	North Biscay	South Biscay	Portugal
North Sea		0.0021	0.0057	0.0046	0.0031	0.0032	0.0029
Channel	0.6877		0.0017	0.0017	0.0014	0.0014	0.0013
Irish Sea	0.0004	0.7758		0.0040	0.0026	0.0026	0.0024
Celtic Sea	0.0412	0.6915	0.0434		0.0025	0.0023	0.0022
North Biscay	0.0982	0.8728	0.1930	0.1442		0.0017	0.0019
South Biscay	0.1122	0.8856	0.2084	0.2691	0.5497		0.0015
Portugal	0.5577	0.9976	0.7341	0.7107	0.7696	0.9486	

**Table S6.** Pairwise Fst values at the 1012 retained loci (upper triangle), for each pair among the 7 Atlantic regions. Bold black numbers indicate significantly Fst values (p<0.05). Significance of Fst values (empirical *p*-values) are given in lower triangle. Pale grey values indicate not significant values.

Region		Distance	es between localities	Mantel's	p-value
name	no. localities	mean	min-max	neutrals	outliers
North of GONB <sup>(1)</sup>	9	546	117-1285	0.01510*	0.01110*
South of GONB to Galicia <sup>(2)</sup>	7	508	169-1111	0.17548	0.52045
Galicia to South Portugal <sup>(3)</sup>	5	372	146-730	0.96840	0.17838

**Table S7.** Mantel test between matrices of geographic distances (nautic milles) and genetic distances [Fst/(1-Fst)] between pairs of localities within 3 groups of sampling locations that differ in their average ancestry proportions (see Figure 3). The test was performed using 10.000 permutations, p-values<0.05 are indicated in bold and labeled by an asterisk. <sup>(1)</sup> ANGL, BSEI, CRNW, DOVE, IRIS, ISWI, NEUK, SGEO, WALE;

<sup>(2)</sup> AQUI, ASTU, CANT, CHAR, GONB, LOIR, PBRE;

<sup>(3)</sup> CORU, PENI, PORT, SINE, VIGO.