Ancestry, Language and Culture

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Online Appendix – Supplementary Empirical Results

<u>Part A – Alternative choices of sample and questions.</u> <u>Metric of cultural distance: Euclidian distance.</u>

In Part A of this appendix we consider alternative samples and sets of questions. In the baseline included in the paper, we sought to maximize the country coverage. The requirement that the same number of questions be included in the analysis for all country pairs led to the loss of many WVS questions. In the tables that follow we replicate Tables 7, 8 and 9 in the paper considering alternative tradeoffs between country coverage and question coverage, resulting in the use of more questions but fewer countries.

1. <u>Maximizing the number of questions, but including China.</u>

	CD, all questions	CD, cat. A	CD, cat. B	CD, cat. C	CD, cat. D	CD, cat. E	CD, cat. F	CD, cat. G	CD, binary questions	CD, Non- binary questions
CD, category A	0.911*	1								
CD, category B	0.098*	0.0318	1							
CD, category C	0.739*	0.587*	0.027	1						
CD, category D	0.611*	0.511*	0.024	0.469*	1					
CD, category E	0.851*	0.689*	0.064*	0.515*	0.431*	1				
CD, category F	0.673*	0.467*	0.097*	0.458*	0.403*	0.478*	1			
CD, category G	0.345*	0.299*	0.131*	0.117*	0.277*	0.251*	0.257*	1		
CD, binary questions	0.923*	0.948*	0.051	0.740*	0.535*	0.667*	0.538*	0.242*	1	
CD, non-binary questions	0.953*	0.783*	0.125*	0.658*	0.606*	0.905*	0.707*	0.389*	0.763*	1
Weighted F _{ST} gen. dist.	0.172*	0.191*	0.080*	0.078*	0.057*	0.127*	0.121*	0.078*	0.175*	0.151*

Table 7A1 – Correlations between Genetic Distance and Cultural Distance measures (Maximizing # of questions with China, Euclidian distance)

(Correlations based on 1,275 observations; * denotes 5% significance). We are using a matrix of 51 countries and 164 questions. We are trying to maximize the number of questions in our analysis. 51 country combinations gives us a total of 1275 observations. In this combination China is included.

	(1)	(2)	(3)	(4)	(5)	(6)
Variable	Total	Total	Binary	Binary	Non-binary	Non-binary
F _{ST} genetic distance,	187.893	117.739	89.831	79.399	98.061	38.341
weighted	(6.22)***	(3.42)***	(6.33)***	(4.82)***	(5.45)***	(1.88)*
Geodesic Distance,		-3.427		-2.060		-1.366
1000s of km		(2.93)***		(3.68)***		(1.97)**
Absolute difference in		0.322		0.187		0.134
longitudes		(4.27)***		(5.19)***		(3.00)***
Absolute difference in		0.174		0.071		0.103
latitudes		(1.43)		(1.21)		(1.43)
1 for contiguity		-55.985		-24.837		-31.147
		(9.37)***		(8.68)***		(8.80)***
Number of landlocked		-8.496		-5.486		-3.009
countries in the pair		(3.25)***		(4.38)***		(1.94)*
Number of island		19.074		1.974		17.100
countries in the pair		(5.21)***		(1.13)		(7.88)***
1 if pair shares at least		-21.332		-6.286		-15.046
one sea or ocean		(4.35)***		(2.68)***		(5.18)***
Constant	-11.184	-4.644	-5.164	-1.480	-6.020	-3.163
	(5.05)***	(1.48)	(4.95)***	(0.99)	(4.56)***	(1.70)*
Adjusted R ²	0.03	0.17	0.03	0.14	0.02	0.17
# of observations	1,275	1,250	1,275	1,250	1,275	1,250
Standardized beta	0.172	0.108	0.175	0.154	0.151	0.059

Table 8A1 – Cultural Distance and Genetic Distance (Maximizing # of questions with China, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 51 countries and 164 questions. We are trying to maximize the number of questions in our analysis. 51 country combinations gives us a total of 1,275 observations. In this combination China is included.

	(1)	(2)	(3)	(4)	(5)	(6)	(7)
Variable	Category A	Category B	Category C	Category D	Category E	Category F	Category G
			Univ	ariate Specificat	tion		
F _{ST} genetic distance,	93.152	2.845	16.405	4.579	43.713	24.133	3.067
weighted	(6.95)***	(2.87)***	(2.80)***	(2.04)**	(4.57)***	(4.34)***	(2.79)***
Adjusted R ²	0.04	0.01	0.01	0.00	0.02	0.01	0.01
Standardized beta	0.191	0.080	0.078	0.057	0.127	0.121	0.078
			Multi	variate Specifica	ation		
F _{ST} genetic distance,	67.806	5.129	22.249	-0.984	14.652	11.205	-2.318
weighted	(4.41)***	(4.27)***	(3.25)***	(0.40)	(1.30)	(1.69)*	(1.77)*
Adjusted R ²	0.16	0.05	0.11	0.21	0.11	0.09	0.08
Standardized beta	0.139	0.144	0.106	-0.012	0.043	0.056	-0.059

Table 9A1 - Cultural Distance and Genetic Distance, by question category (Maximizing # of questions with China, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 51 countries and 164 questions. We are trying to maximize the number of questions in our analysis. 51 country combinations gives us a total of 1,275 observations. In this combination China is included.

The univariate specification is based on 1,275 observations (country pairs). The multivariate specification is based on 1,250 observations. All specifications include an intercept.

The multivariate specification includes the following geographic controls: Geodesic distance, absolute difference in longitudes, absolute difference in latitudes, dummy for contiguity, number of landlocked countries in the pair, number of island countries in the pair, dummy=1 if pair shares at least one sea or ocean.

2. <u>Maximizing the number of questions, but excluding China (version 1).</u>

	CD, all questions	CD, cat. A	CD, cat. B	CD, cat. C	CD, cat. D	CD, cat. E	CD, cat. F	CD, cat. G	CD, binary questions	CD, Non- binary questions
CD, category A	0.863*	1								•
CD, category B	0.107*	0.006	1							
CD, category C	0.776*	0.615*	0.112*	1						
CD, category D	0.770*	0.547*	0.085*	0.612*	1					
CD, category E	0.825*	0.670*	0.071*	0.559*	0.583*	1				
CD, category F	0.686*	0.372*	0.100*	0.444*	0.548*	0.386*	1			
CD, category G	0.315*	0.230*	0.108*	0.146*	0.300*	0.219*	0.255*	1		
CD, binary questions	0.929*	0.913*	0.068*	0.750*	0.649*	0.635*	0.631*	0.233*	1	
CD, non-binary questions	0.957*	0.739*	0.127*	0.719*	0.789*	0.892*	0.661*	0.347*	0.781*	1
Weighted F _{ST} gen. dist.	0.131*	0.061	0.027	0.166*	0.267*	0.057	0.098*	0.035	0.091*	0.149*

 Table 7A2 – Correlations between Genetic Distance and Cultural Distance measures

 (Maximizing # of questions without China, version 1, Euclidian distance)

(Correlations based on 990 observations; * denotes 5% significance). In this version we are using a matrix of 45 countries and 226 questions. We are trying to maximize the number of questions in our analysis. 45 country combinations gives us a total of 990 observations. In this combination China is excluded.

	(1)	(2)	(3)	(4)	(5)	(6)
Variable	Total	Total	Binary	Binary	Non-	Non-
					binary	binary
F _{ST} genetic distance,	267.752	178.678	87.038	51.891	180.714	126.786
weighted	(4.15)***	(2.62)***	(2.88)***	(1.61)	(4.73)***	(3.16)***
Geodesic Distance, 1000s		-5.570		-2.781		-2.789
of km		(2.62)***		(2.77)***		(2.23)**
Absolute difference in		0.413		0.245		0.167
longitudes		(3.18)***		(4.00)***		(2.19)**
Absolute difference in		0.380		0.117		0.262
latitudes		(1.71)*		(1.12)		(2.01)**
1 for contiguity		-75.251		-33.302		-41.949
		(8.61)***		(8.06)***		(8.15)***
Number of landlocked		-7.310		-6.590		-0.720
countries in the pair		(1.81)*		(3.46)***		(0.30)
Number of island		22.254		1.572		20.682
countries in the pair		(3.51)***		(0.52)		(5.53)***
1 if pair shares at least		-29.907		-9.087		-20.820
one sea or ocean		(3.96)***		(2.54)**		(4.68)***
Constant	-11.198	-1.575	-3.640	1.873	-7.558	-3.449
	(3.18)***	(0.33)	(2.21)**	(0.82)	(3.63)***	(1.22)
Adjusted R ²	0.02	0.14	0.01	0.12	0.02	0.15
# observations	990	967	990	967	990	967
Standardized beta	0.131	0.087	0.091	0.054	0.149	0.105

Table 8A2 – Cultural Distance and Genetic Distance (Maximizing # of questions without China, version 1, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 45 countries and 226 questions. We are trying to maximize the number of questions in our analysis. 45 country combinations gives us a total of 990 observations. In this combination China is excluded (version 1).

	(1)	(2)	(3)	(4)	(5)	(6)	(7)
Variable	Category A	Category B	Category C	Category D	Category E	Category F	Category G
			Univ	variate Specificat	ion		
F _{ST} genetic distance,	49.263	1.311	58.608	68.532	32.067	55.986	1.984
weighted,	(1.93)*	(0.84)	(5.30)***	(8.69)***	(1.78)*	(3.10)***	(1.11)
Adjusted R ²	0.00	-0.00	0.03	0.07	0.00	0.01	0.00
Standardized beta	0.061	0.027	0.166	0.266	0.057	0.098	0.035
			Mult	ivariate Specifica	tion		
F _{ST} genetic distance,	6.524	4.062	64.066	66.716	12.201	28.538	-3.430
weighted	(0.24)	(2.36)**	(5.36)***	(7.96)***	(0.63)	(1.44)	(1.75)*
Adjusted R ²	0.14	0.04	0.10	0.18	0.10	0.07	0.06
Standardized beta	0.008	0.083	0.182	0.259	0.022	0.050	-0.061

Table 9A2 – Cultural Distance and Genetic Distance, by question category (Maximizing # of questions without China, version 1, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 45 countries and 226 questions. We are trying to maximize the number of questions in our analysis. 45 country combinations gives us a total of 990 observations. In this combination China is excluded.

The univariate specification is based on 990 observations (country pairs). The multivariate specification is based on 967 observations. All specifications include an intercept.

The multivariate specification includes the following geographic controls: Geodesic distance, absolute difference in longitudes, absolute difference in latitudes, dummy for contiguity, number of landlocked countries in the pair, number of island countries in the pair, dummy=1 if pair shares at least one sea or ocean.

3. Maximizing the number of questions, but excluding China (version 2).

	CD, all questions	CD, cat. A	CD, cat. B	CD, cat. C	CD, cat. D	CD, cat. E	CD, cat. F	CD, cat. G	CD, binary questions	CD, Non- binary questions
CD, category A	0.878*	1								
CD, category B	0.100*	0.006	1							
CD, category C	0.748*	0.603*	0.048	1						
CD, category D	0.617*	0.522*	0.024	0.472*	1					
CD, category E	0.828*	0.669*	0.062*	0.558*	0.444*	1				
CD, category F	0.694*	0.393*	0.119*	0.423*	0.382*	0.403*	1			
CD, category G	0.342*	0.284*	0.154*	0.118*	0.279*	0.229*	0.267*	1		
CD, binary questions	0.938*	0.914*	0.047	0.732*	0.555*	0.645*	0.655*	0.266*	1	
CD, non-binary questions	0.953*	0.758*	0.137*	0.685*	0.608*	0.906*	0.657*	0.374*	0.790*	1
Weighted F _{ST} gen. dist.	0.174*	0.127*	-0.008	0.177*	0.089*	0.117*	0.161*	0.089*	0.156*	0.172*

 Table 7A3 – Correlations between Genetic Distance and Cultural Distance measures

 (Maximizing # of questions without China, version 2, Euclidian distance)

(Correlations based on 1,081 observations; * denotes 5% significance). In this version we are using a matrix of 47 countries and 164 questions. We are trying to maximize the number of questions in our analysis. 47 country combinations gives us a total of 1,081 observations. In this combination China is excluded.

	(1)	(2)	(3)	(4)	(5)	(6)
Variable	Total	Total	Binary	Binary	Non-	Non-
					binary	binary
F _{ST} genetic distance,	278.916	145.647	123.624	70.541	155.291	75.106
weighted	(5.80)***	(2.72)***	(5.19)***	(2.62)***	(5.73)***	(2.49)**
Geodesic Distance, 1000s		-6.435		-3.516		-2.919
of km		(4.10)***		(4.44)***		(3.30)***
Absolute difference in		0.460		0.281		0.178
longitudes		(4.61)***		(5.60)***		(3.18)***
Absolute difference in		0.471		0.213		0.259
latitudes		(2.85)***		(2.55)**		(2.78)***
1 for contiguity		-63.734		-29.215		-34.519
		(8.97)***		(8.16)***		(8.63)***
Number of landlocked		-9.827		-6.826		-3.002
countries in the pair		(3.06)***		(4.22)***		(1.66)*
Number of island		29.562		8.912		20.650
countries in the pair		(6.39)***		(3.82)***		(7.93)***
1 if pair shares at least		-26.454		-9.043		-17.410
one sea or ocean		(4.41)***		(2.99)***		(5.15)***
Constant	-14.325	-4.927	-6.274	-1.751	-8.050	-3.176
	(4.95)***	(1.30)	(4.37)***	(0.91)	(4.94)***	(1.48)
Adjusted R ²	0.03	0.19	0.02	0.16	0.03	0.19
# observations	1,081	1,058	1,081	1,058	1,081	1,058
Standardized beta	0.174	0.091	0.156	0.089	0.172	0.083

Table 8A3 – Cultural Distance and Genetic Distance (Maximizing # of questions without China, version 2, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 47 countries and 188 questions. We are trying to maximize the number of questions in our analysis. 47 country combinations gives us a total of 1081 observations. In this combination China is excluded (version 2).

	(1)	(2)	(3)	(4)	(5)	(6)	(7)
Variable	Category A	Category B	Category C	Category D	Category E	Category F	Category G
			Univ	variate Specifica	tion		
F _{ST} genetic distance,	84.957	-0.359	47.729	9.016	56.265	76.791	4.518
weighted,	(4.20)***	(0.27)	(5.91)***	(2.93)***	(3.88)***	(5.36)***	(2.93)***
Adjusted R ²	0.02	-0.00	0.03	0.01	0.01	0.03	0.01
Standardized beta	0.127	-0.008	0.177	0.089	0.117	0.161	0.089
			Mult	ivariate Specific	ation		
F _{ST} genetic distance,	25.567	3.975	51.135	0.855	21.984	44.913	-2.782
weighted	(1.13)	(2.51)**	(5.49)***	(0.26)	(1.32)	(2.68)***	(1.54)
Adjusted R ²	0.16	0.05	0.13	0.23	0.12	0.11	0.08
Standardized beta	0.038	0.091	0.190	0.008	0.046	0.094	-0.055

Table 9A3 – Cultural Distance and Genetic Distance, by question category (Maximizing # of questions without China, version 2, Euclidian distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 47 countries and 188 questions. We are trying to maximize the number of questions in our analysis. 47 country combinations gives us a total of 1,081 observations. In this combination China is excluded.

The univariate specification is based on 1,081 observations (country pairs). The multivariate specification is based on 1,058 observations. All specifications include an intercept.

The multivariate specification includes the following geographic controls: Geodesic distance, absolute difference in longitudes, absolute difference in latitudes, dummy for contiguity, number of landlocked countries in the pair, number of island countries in the pair, dummy=1 if pair shares at least one sea or ocean.

<u>Part B – Alternative metrics of cultural distance: Manhattan Distance.</u> <u>Sample choice: maximizing the number of countries available.</u>

The three tables that follow redo Tables 7, 8 and 9 in the main paper, using Manhattan distance as a measure of cultural distance from the WVS.

	CD, all questions	CD, cat. A	CD, cat. C	CD, cat. D	CD, cat. E	CD, cat. F	CD, cat. G	CD, binary questions	CD, Non- binary questions
CD, category A	0.879*	1							
CD, category C	0.644*	0.481*	1						
CD, category D	0.620*	0.530*	0.375*	1					
CD, category E	0.802*	0.598*	0.284*	0.392*	1				
CD, category F	0.729*	0.544*	0.394*	0.399*	0.476*	1			
CD, category G	0.443*	0.327*	0.311*	0.217*	0.308*	0.287*	1		
CD, binary questions	0.870*	0.837*	0.816*	0.584*	0.514*	0.544*	0.315*	1	
CD, non-binary questions	0.949*	0.787*	0.448*	0.560*	0.878*	0.749*	0.466*	0.670*	1
Weighted F _{ST} gen. dist.	0.269*	0.245*	0.075*	0.026	0.298*	0.216*	0.129*	0.147*	0.311*

Table 7B – Correlations between Genetic Distance and Cultural Distance measures (Maximizing # of countries, Manhattan distance)

(Correlations based on 2,628 observations; * denotes 5% significance). We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations.

	(1)	(2)	(3)	(4)	(5)	(6)
Variable	Total	Total	Binary	Binary	Non-	Non-
					binary	binary
F _{ST} genetic distance,	143.659	137.954	33.291	29.873	110.368	108.081
weighted	(14.30)***	(12.18)***	(7.60)***	(6.02)***	(16.75)***	(14.53)***
Geodesic Distance,		-1.071		-0.334		-0.737
1000s of km		(2.14)**		(1.52)		(2.24)**
Absolute difference in		0.071		0.019		0.052
longitudes		(1.96)**		(1.21)		(2.18)**
Absolute difference in		0.107		0.037		0.071
latitudes		(2.14)**		(1.67)*		(2.14)**
1 for contiguity		-34.388		-13.900		-20.488
		(9.92)***		(9.16)***		(9.00)***
Number of landlocked		-8.808		-4.027		-4.781
countries in the pair		(7.32)***		(7.65)***		(6.05)***
Number of island		-3.252		-2.704		-0.548
countries in the pair		(2.07)**		(3.93)***		(0.53)
1 if pair shares at least		-12.103		-2.999		-9.103
one sea or ocean		(5.25)***		(2.98)***		(6.02)***
Constant	-10.549	-3.685	-2.368	0.702	-8.182	-4.386
	(10.84)***	(2.50)**	(5.58)***	(1.09)	(12.83)***	(4.53)***
Adjusted R ²	0.07	0.14	0.02	0.08	0.10	0.16
# observations	2,628	2,513	2,628	2,513	2,628	2,513
Standardized beta	0.269	0.254	0.147	0.130	0.311	0.300

Table 8B – Cultural Distance and Genetic Distance (Maximizing # of countries, Manhattan distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations.

	(1)	(3)	(4)	(5)	(6)	(7)
Variable	Category A	Category C	Category D	Category E	Category F	Category G
			Univariate	Specification		
F _{ST} genetic distance,	47.646	9.019	1.455	57.263	24.171	4.105
weighted,	(12.93)***	(3.85)***	(1.34)	(15.99)***	(11.33)***	(6.67)***
Adjusted R ²	0.06	0.01	0.00	0.09	0.05	0.02
Standardized beta	0.245	0.075	0.026	0.298	0.216	0.129
			Multivariate	Specification		
F _{ST} genetic distance,	47.659	7.000	1.406	54.871	24.842	2.175
weighted	(11.56)***	(2.62)***	(1.15)	(13.50)***	(10.09)***	(3.11)***
Adjusted R ²	0.14	0.05	0.08	0.14	0.08	0.04
Standardized beta	0.241	0.058	0.025	0.283	0.218	0.069

Table 9B – Cultural Distance and Genetic Distance, by question category (Maximizing # of countries, Manhattan distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations

The univariate specification is based on 2,628 observations (country pairs). The multivariate specification is based on 2,513 observations. All specifications include an intercept.

The multivariate specification includes the following geographic controls: Geodesic distance, absolute difference in longitudes, absolute difference in latitudes, dummy for contiguity, number of landlocked countries in the pair, number of island countries in the pair, dummy=1 if pair shares at least one sea or ocean.

<u>Part C – Alternative metrics of cultural distance: F_{ST} Cultural Distance.</u> <u>Sample choice: maximizing the number of countries available.</u>

The three tables that follow redo Tables 7, 8 and 9 in the main paper, using the F_{ST} functional form to compute cultural distance from the WVS.

	CD, all questions	CD, cat. A	CD, cat. C	CD, cat. D	CD, cat. E	CD, cat. F	CD, cat. G	CD, binary questions	CD, Non- binary questions
CD, category A	0.903*	1							
CD, category C	0.750*	0.586*	1						
CD, category D	0.675*	0.622*	0.440*	1					
CD, category E	0.736*	0.550*	0.368*	0.410*	1				
CD, category F	0.775*	0.566*	0.509*	0.395*	0.556*	1			
CD, category G	0.411*	0.296*	0.277*	0.159*	0.366*	0.387*	1		
CD, binary questions	0.926*	0.930*	0.805*	0.6538*	0.515*	0.588*	0.274*	1	
CD, non-binary questions	0.916*	0.727*	0.571*	0.588*	0.850*	0.847*	0.490*	0.697*	1
Weighted F _{ST} gen. dist.	0.132*	0.070*	0.001	-0.047*	0.235*	0.222*	0.103*	0.026	0.222*

Table 7C – Correlations between Genetic Distance and Cultural Distance measures (Maximizing # of countries, F_{ST} distance)

(Correlations based on 2,628 observations; * denotes 5% significance). We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations.

	(1)	(2)	(3)	(4)	(5)	(6)
Variable	Total	Total	Binary	Binary	Non-	Non-
					binary	binary
F _{ST} genetic distance,	0.041	0.038	0.013	0.015	0.057	0.050
weighted	(6.80)***	(5.50)***	(1.34)	(1.41)	(11.66)***	(9.18)***
Geodesic Distance, 1000s		0.000		-0.001		0.001
of km		(1.30)		(1.08)		(3.75)***
Absolute difference in		-0.000		0.000		-0.000
longitudes		(1.49)		(0.21)		(3.13)***
Absolute difference in		-0.000		0.000		-0.000
latitudes		(1.86)*		(0.32)		(3.99)***
1 for contiguity		-0.016		-0.025		-0.011
		(7.62)***		(7.36)***		(6.69)***
Number of landlocked		-0.008		-0.009		-0.007
countries in the pair		(10.47)***		(7.92)***		(11.64)***
Number of island countries		-0.001		-0.005		0.001
in the pair		(1.15)		(3.08)***		(1.18)
1 if pair shares at least one		-0.006		-0.008		-0.005
sea or ocean		(4.09)***		(3.46)***		(4.14)***
Constant	0.036	0.042	0.048	0.057	0.030	0.034
	(61.73)***	(47.11)***	(51.85)***	(39.74)***	(62.79)***	(47.81)***
Adjusted R ²	0.02	0.09	0.00	0.05	0.05	0.13
# observations	2,628	2,513	2,628	2,513	2,628	2,513
Standardized beta	0.132	0.119	0.026	0.031	0.222	0.193

Table 8C – Cultural Distance and Genetic Distance (Maximizing # of countries, F_{ST} distance)

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations.

	(1)	(3)	(4)	(5)	(6)	(7)	
Variable	Category A	Category C	Category D	Category E	Category F	Category G	
	Univariate Specification						
F _{ST} genetic distance,	0.029	0.001	-0.024	0.055	0.129	0.029	
weighted,	(3.60)***	(0.07)	(2.42)**	(12.39)***	(11.66)***	(5.30)***	
Adjusted R2	0.00	-0.00	0.00	0.05	0.05	0.01	
Standardized beta	0.070	0.001	-0.047	0.235	0.222	0.103	
	Multivariate Specification						
F _{ST} genetic distance,	0.029	0.003	-0.022	0.050	0.112	0.013	
weighted	(3.19)***	(0.22)	(1.98)**	(10.12)***	(9.00)***	(2.22)**	
Adjusted R2	0.06	0.04	0.08	0.12	0.12	0.04	
Standardized beta	0.069	0.005	-0.043	0.214	0.190	0.049	

$\label{eq:constraint} \begin{array}{l} \mbox{Table 9C} - \mbox{Cultural Distance and Genetic Distance, by question category} \\ (Maximizing \# of countries, F_{ST} distance) \end{array}$

(t-statistics in parentheses; * significant at 10%; ** significant at 5%; ** significant at 1%)

We are using a matrix of 73 countries and 98 questions. We are trying to maximize the number of countries in our analysis. 73 country combinations gives us a total of 2,628 observations

The univariate specification is based on 2,628 observations (country pairs). The multivariate specification is based on 2513 observations.

All specifications include an intercept.

The multivariate specification includes the following geographic controls: Geodesic distance, absolute difference in longitudes, absolute difference in latitudes, dummy for contiguity, number of landlocked countries in the pair, number of island countries in the pair, dummy=1 if pair shares at least one sea or ocean.

Part D – Simple Correlations between Various Metrics of Cultural Distance, by question type and category

This section explores the simple correlations between Euclidian, Manhattan and FST measures of cultural distance for the baseline sample of countries and set of questions used in the main paper. Note that for binary questions Manhattan and Euclidian distances are the same.

Table D1. Correlation matrices between Euclidian, Manhattan and F_{ST} distances for various question types

Panel A – Total Index of Cultural Distance (all questions together)

	Euclidian	Manhattan
Manhattan	0.9999	1
F _{ST}	0.8734	0.8731

Panel B – Breakdown by question type

	Euclidian	Manhattan	Euclidian	Manhattan	
	Binary	Distance	Non-binary Distance		
Manhattan	1	1	0.9998	1	
F _{ST}	0.9172	0.9172	0.8173	0.8166	

Panel C – Breakdown by Question Category

	Euclidian	Manhattan	Euclidian	Manhattan		
	A Categor	y Distance	C Category Distance			
Manhattan	1	1	0.9998	1		
F _{ST}	0.8611	0.8614	0.9122	0.9118		
	D Categor	D Category Distance		E Category Distance		
Manhattan	0.9997	1	0.9997	1		
F _{ST}	0.8371	0.8360	0.8416	0.8400		
	F Categor	F Category Distance		G Category Distance		
Manhattan	1	1	0.9889	1		
F _{ST}	0.7050	0.7051	0.7701	0.7602		

Sample: 73 countries, i.e. 2,628 observations