The non-independence of nations and why it matters

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### Author Note

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### Abstract

Cross-national analyses test hypotheses about the drivers of variation in national outcomes. However, since nations are connected in various ways, such as via spatial proximity and shared cultural ancestry, cross-national analyses often violate assumptions of non-independence, inflating false positive rates. Here, we show that, despite being recognised as an important statistical pitfall for over 200 years, cross-national research in economics and psychology still does not sufficiently account for non-independence. In a review of the 100 highest-cited cross-national studies of economic development and values, we find that controls for non-independence are rare. When studies do control for non-independence, our simulations suggest that most commonly used methods are insufficient for reducing false positives in non-independent data. In reanalyses of twelve previous cross-national correlations, half of the estimates are compatible with no association after controlling for non-independence using global proximity matrices. We urge social scientists to sufficiently control for non-independence in cross-national research.

Keywords: spatial non-independence; cultural non-independence; cross-national analyses; simulations; replications; reanalyses

Word count: 4892 words excl. Methods (Methods: 1877 words)

## The non-independence of nations and why it matters

Nations are perhaps the single most important population unit structuring modern
human life. The nation in which someone is born has a large effect on what they can
expect out of life, including their income level<sup>1</sup>, life expectancy<sup>2</sup>, mental health<sup>3</sup>, subjective
well-being<sup>4</sup>, and educational attainment<sup>5</sup>. Nations are also among the most important
determinants of human cultural variation, with greater cultural similarity within than
outside national borders<sup>6</sup>.

Given the importance of nations for structuring human behaviour, there is justifiably huge interest in statistical analyses that attempt to predict variation in national outcomes of all kinds. At the time of writing, a search in Web of Science for the term "cross-national" in titles or abstracts returned over 13,000 unique hits. The standard 10 practice for cross-national analyses is to conduct bivariate correlations or multiple 11 regressions with individual data points representing different nations. Such analyses widen 12 the scope of social science beyond Western populations<sup>7,8</sup> and have been used to study. 13 among other topics, the causes of variation in the economic wealth of nations<sup>9–12</sup>, global 14 patternings of cultural norms and values 13-16, and the universality and diversity of human 15 behaviour and psychology around the world <sup>17–20</sup>. 16

However, cross-national analyses are complicated by the fact that nations are not statistically independent data points. Unlike independent random samples from a population, nations are related to one another in a number of ways. First, nations that are closer to one another tend to be more similar than distant nations. This phenomenon is known as spatial non-independence<sup>21</sup>, and it occurs because nations in close spatial proximity share characteristics due to local cultural diffusion of ideas<sup>22</sup> and regional variation in climate and environment<sup>21</sup>. For example, the neighbouring African nations Zambia and Tanzania have similar levels of terrain ruggedness, which has been used to partially explain their similar levels of economic development<sup>23</sup>. This pattern conforms to

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Tobler's first law of geography: "everything is related to everything else, but near things are more related than distant things" (p. 236).

Second, nations with shared cultural ancestry tend to be more similar than culturally 28 unrelated nations. This is known as cultural phylogenetic non-independence<sup>25–27</sup>, and 29 occurs because related nations share cultural traits inherited via descent from a common ancestor. Shared cultural ancestry can result in a form of pseudoreplication, whereby multiple instances of the same trait across nations are merely duplicates of the ancestral original. For example, the related island nations Tonga and Tuvalu share similar languages and customs due to cultural inheritance from a common Polynesian population dating back more than 1,000 years. More recently, shared ancestry explains similarities in institutions, norms, technologies, and values between colonial settlements and their colonisers (e.g. English, French, Spanish, and Portuguese settlements of the Americas). Importantly, 37 these deep cultural connections between nations often span large geographic distances 38 around the world. Tonga and Tuvalu share cultural traits despite being separated by over 39 1,500 kilometres of ocean, and South American and European nations remain culturally similar today despite their locations on two separate continents. Shared cultural ancestry 41 must therefore be considered independently of spatial proximity in the study of nations. 42

Spatial and cultural phylogenetic non-independence between nations make

cross-national inference challenging. A fundamental assumption of regression analysis is

that model residuals should be independently and identically distributed<sup>28</sup>. But without

accounting for spatial or cultural non-independence between nations, model residuals can

show structure that remains unaccounted for, violating this assumption. Treating nations

as independent can thus inflate false positive rates<sup>29</sup>, producing spurious "direct"

relationships between variables that in fact only indirectly covary due to spatial or cultural

connections<sup>30</sup> (see Supplementary Figure S1 for an example causal model).

Non-independence is widely acknowledged in fields that routinely deal with spatially

or culturally structured data. In ecology and sociology, studies with regional-level data use a variety of autoregressive models designed to account for spatial patternings<sup>31,32</sup>. In 53 anthropology, researchers have recognised cultural non-independence as an important statistical pitfall for over 200 years, with issues of cultural pseudoreplication being identified in early comparative studies of marriage practices across societies<sup>25</sup>. In the twentieth century, anthropologists began to emphasise that human societies do not develop 57 independently, but rather exist in a globally interconnected "world system" linked by shared history and cultural ancestry<sup>33</sup>. In order to minimise the confounding effects of this non-independence in comparisons of human societies, researchers compiled the Standard Cross-Cultural Sample of 186 cultures which deliberately avoids sampling closely related cultures<sup>34</sup>, though it is difficult to completely remove spatial and cultural dependencies<sup>35,36</sup> 62 and the smaller sample of cultures reduces statistical power. Today, anthropologists borrow phylogenetic comparative methods from evolutionary biology, such as phylogenetic least squares regression<sup>37</sup>, when comparing societies, treating culturally related societies in the same way as biologists treat genetically related species (e.g. <sup>38,39</sup>). 66

At the national level, recent reanalyses have revealed that several cross-national relationships reported in economics and psychology do not hold when controlling for non-independence between nations. One working paper replicated 25 analyses of "persistence" in economics, in which modern national outcomes are regressed against historical characteristics of those nations, and found that over half of the relationships were attenuated when controlling for spatial non-independence<sup>40</sup>. Another replication study found that many of the widely publicised relationships between national-level pathogen prevalence and political institutions and attitudes fail to hold when controlling for various kinds of non-independence<sup>41</sup>. These reanalyses, and others<sup>42–44</sup>, raise the question: how widespread a concern is non-independence in studies of national-level outcomes?

To address this question, we consider national-level variables of general interest across
the social sciences: economic development and cultural values. These variables are

frequently included as both outcomes and predictors in cross-national studies in economics and psychology<sup>9-16</sup>. First, we demonstrate that economic development and cultural values are spatially and culturally non-independent across nations, emphasising the need to 81 control for non-independence. Second, we systematically review the 100 highest-cited 82 cross-national studies of economic development and cultural values and determine baseline rates of controlling for non-independence in the literature. Third, we run simulations to 84 determine whether common methods of dealing with non-independence in the literature 85 sufficiently reduce false positive rates. Fourth, we reanalyse twelve previous cross-national analyses of economic development and cultural values from our systematic review, 87 incorporating global geographic and linguistic proximity matrices as controls for spatial and cultural non-independence.

90 Results

National-level economic development and cultural values are spatially and culturally non-independent

In order to motivate our research question, it is important to first quantify the degree of spatial and cultural non-independence for economic development and cultural values around the world. If these variables are independent or only weakly non-independent, then the issue might be safe to ignore. However, if they are more strongly non-independent, then there is a possibility that non-independence could be confounding cross-national inferences.

To this end, we used Bayesian multilevel models to simultaneously estimate
geographic and cultural phylogenetic signal for a range of economic development and
cultural values variables. For economic development, we focused on the Human
Development Index<sup>45</sup>, gross domestic product per capita, annual gross domestic product
per capita growth, and the Gini index of income inequality. For cultural values, we focused
on two primary dimensions of cultural values from the World Values Survey, traditional

vs. secular values and survival vs. self-expression values<sup>16</sup>, as well as cultural tightness<sup>14</sup> and individualism<sup>15</sup>. These variables are not intended to be a comprehensive list of all national-level variables included in cross-national research, but rather an illustrative set of variables that are widely used in the literature.

For all of these variables, we found that a substantial proportion of national-level 108 variation was explained by spatial proximity and/or shared cultural ancestry between 109 nations (Figure 1; see Supplementary Table S1 for numerical results). Signal estimates 110 were often markedly strong, with spatial proximity and shared cultural ancestry frequently 11: explaining over half of the national-level variation. For spatial proximity, Bayes Factors 112 indicated strong evidence that the geographic signal estimates differed from zero for all economic development variables and traditional values. However, the evidence was only 114 equivocal for survival values and individualism, and strong evidence was found that the 115 geographic signal estimate for tightness was equal to zero. For shared cultural ancestry, 116 Bayes Factors indicated strong evidence that the cultural phylogenetic signal estimates 117 differed from zero for all economic development and cultural values variables except for 118 gross domestic product per capita growth, for which the evidence was equivocal. These 110 findings emphasise the need to account for spatial and cultural phylogenetic 120 non-independence in cross-national analyses of economic development and cultural values. 121

# Previous cross-national analyses have not sufficiently accounted for non-independence

Given that economic development and cultural values show evidence of geographic and cultural phylogenetic signal, have cross-national analyses sufficiently accounted for this non-independence? To assess this, we systematically searched the published literature for articles that combined the search terms "economic development" or "values" with the search terms "cross-national", "cross-cultural", or "cross-country". We removed articles that did not report original research, were not relevant to economic development or

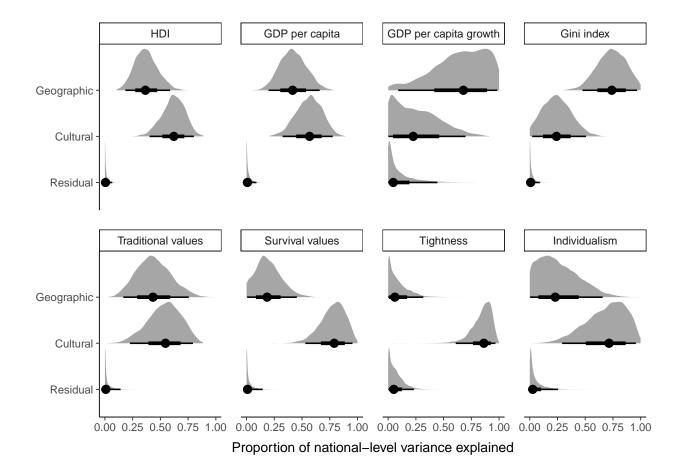


Figure 1. Posterior estimates of geographic and cultural phylogenetic signal for a range of economic development and cultural values variables. Geographic and cultural phylogenetic signal are operationalised as the proportion of national-level variance explained by geographic and linguistic proximity matrices. Grey ridges are full posterior distributions, points are posterior median values, and black lines are 50% and 95% credible intervals. HDI = Human Development Index; GDP = gross domestic product.

cultural values, or did not report at least one cross-national analysis. We then retained the
100 articles (50 for economic development, 50 for cultural values) with the highest annual
rate of citations (Supplementary Table S2). For each of these highly-cited articles, we
exhaustively recorded every cross-national analysis reported in the main text (n = 4,308),
identifying in each case whether or not the analysis attempted to control for spatial,
cultural, or any other form of non-independence between nations (see Methods for detailed
search criteria and coding decisions).

The results of our systematic review show that most published articles containing 137 cross-national analyses make no attempt to account for statistical non-independence. Figure 2a plots the proportion of articles that contain at least one cross-national analysis accounting for non-independence. We find that 42% of economic development articles 140 contain at least one attempt to control for non-independence (95% bootstrap confidence 141 interval [0.30 0.54]), while this proportion decreases to only 8% for cultural values articles 142 (95% bCI [0.02 0.16]). Both kinds of article are most likely to use regional fixed effects 143 (e.g. continent fixed effects) to account for non-independence, but some articles also 144 include controls for spatial distance (e.g. latitude) and shared cultural history (e.g. colony 145 status). These proportions are even lower when focusing on the full sample of 4,308 146 analyses: only 5% (95% credible interval [0.02 0.13]) of individual economic development 147 analyses and 1% (95% CI [0.00 0.02]) of individual cultural values analyses are estimated 148 to control for non-independence (Supplementary Figure S2). 149

While our review contains articles from journals with a range of impact factors, an anonymous reviewer suggested that our estimates could be biased downwards by analyses published in lower impact outlets with more relaxed standards for issues like non-independence. It is also possible that, since our systematic review goes back as far as 1993, our estimates are being biased downwards by earlier studies, and that controls for non-independence have increased over time with methodological advancements and greater awareness of the issue. To test these possible explanations for our low estimates, we fitted

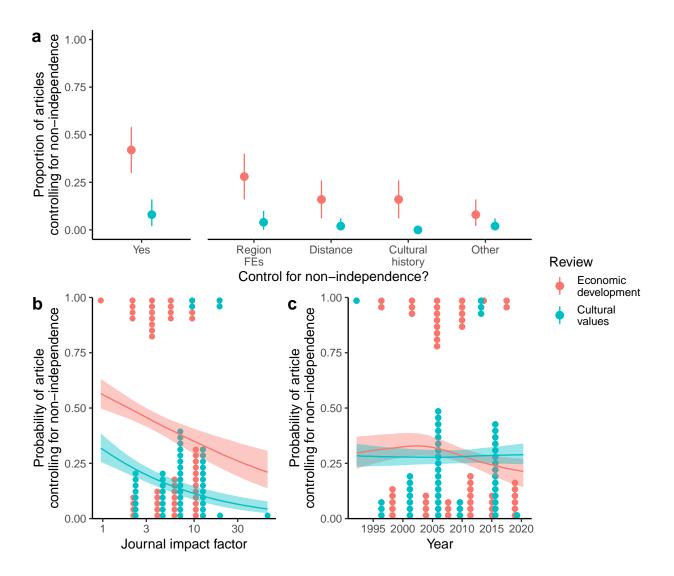


Figure 2. Results from systematic review of 100 highly-cited cross-national studies of economic development (red) and cultural values (blue). (a) Proportion of articles containing at least one analysis accounting for non-independence, overall and split by common methods of controlling for non-independence. (b) The association between journal impact factor and the probability that an article contains at least one analysis accounting for non-independence. (c) Estimated trend over time for the probability that an article contains at least one analysis accounting for non-independence. Point ranges represent proportions and 95% bootstrap confidence intervals. Lines and shaded areas are posterior median regression lines and 50% credible intervals from Bayesian logistic regression models. Dots represent raw counts of individual articles that did (top) or did not (bottom) account for non-independence. Region FEs = region fixed effects.

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logistic regression models to the data from the review, including log journal impact factor 157 and publication year as separate predictors. Interestingly, we found that, if anything, 158 studies from higher impact journals were less likely to include at least one control for 159 non-independence than studies from lower impact journals, both for studies of economic 160 development (b = -0.38, 95% CI [-0.87 0.09]) and for studies of cultural values (b = -0.56, 161 95% CI [-1.10 -0.05]; Figure 2b). Moreover, splines revealed no relationship between 162 publication year and the probability of including at least one control for non-independence, 163 both for studies of economic development (b = -0.13, 95% CI [-1.10 0.82]) and for studies 164 of cultural values (b = -0.02, 95% CI [ $-1.00 \ 0.96$ ]; Figure 2c). 165

# Common methods of controlling for non-independence are insufficient for 166 reducing false positive rates in non-independent data

Our systematic review revealed that most cross-national analyses in the literature do 168 not control for spatial or cultural phylogenetic non-independence. When they do, they tend 169 to include controls like latitude and regional fixed effects. Do these methods sufficiently 170 account for statistical non-independence?

To compare the efficacy of different methods in the literature, we conducted a 172 simulation study. We simulated national-level datasets (n = 236 nations) with varying 173 degrees of spatial or cultural phylogenetic autocorrelation (i.e. non-independence) for 174 outcome and predictor variables, but with no direct causal relationship between the 175 variables. We then fitted naive regressions without controls to these datasets, as well as 176 regression models with controls for latitude, longitude, and continent fixed effects. Despite not being identified in our systematic review, we also included other methods that are often 178 used in the literature to account for non-independence. Additional spatial controls included 179 the mean of the predictor variable within a surrounding 2000km radius (e.g. 46) and Conlev 180 standard errors<sup>47,48</sup> based on geographic distances between nations (e.g. <sup>19,46</sup>). Additional 181 cultural controls included fixed effects for the language families of the majority-spoken 182

languages in each nation (e.g. <sup>49</sup>) and Conley standard errors based on genetic distances
between nations (e.g. <sup>19,46</sup>). These fixed effect approaches attempt to account for
non-independence by holding geographic location constant (latitude, longitude), discarding
between-region variation and exploiting only local variation (continent fixed effects, mean
of surrounding 2000km), or correcting standard errors for autocorrelation post-hoc while
leaving model coefficients unchanged (Conley standard errors).

Beyond fixed effect approaches, we also fitted Bayesian random effects regressions 189 that explicitly model spatial and/or cultural phylogenetic non-independence by allowing 190 nations to covary according to geographic and/or linguistic proximity matrices. Geographic 191 proximity between nations is calculated from inverse distances between longitude and latitude coordinates. Linguistic proximity between nations is calculated from a global 193 phylogenetic tree that represents hierarchical relationships of genealogical descent for all 194 languages in the world. For each pair of nations, we calculated inverse phylogenetic 195 distances (i.e. number of branches separating two taxa) between all languages spoken in 196 that nation pair and produced an average "linguistic proximity" score weighted by the 197 percentages of speakers within those nations. To include the resulting geographic and 198 linguistic proximity matrices in our models, we included a Gaussian process<sup>50,51</sup> over 199 latitude and longitude values and/or assumed that nation random intercepts were 200 correlated in proportion to their linguistic proximity<sup>52</sup>. These random effects approaches 201 attempt to account for non-independence by modelling the covariance between nations that 202 is induced by their geographic or linguistic connections. 203

Figures 3 and 4 plot the estimated false positive rates from our simulation study, split
by different methods and different degrees of spatial or cultural phylogenetic
autocorrelation (see Supplementary Tables S3 and S4 for numerical results and
Supplementary Figures S3 and S4 for full distributions of effect sizes under strong
autocorrelation). Across all model types, false positive rates were measured as the
proportion of models that estimated a slope with a 95% confidence / credible interval

excluding zero, falsely inferring a relationship when none is present. For reference, "weak" autocorrelation in our simulation is comparable to the geographic signal for survival values in Figure 1 (i.e. 20% of the national-level variance is explained by non-independence), while "moderate" and "strong" levels of autocorrelation are comparable to the cultural phylogenetic signal for traditional and survival values, respectively (i.e. 50% and 80% of the national-level variance is explained by non-independence).

Our simulation study revealed that with at least moderate degrees of spatial or
cultural phylogenetic autocorrelation for both outcome and predictor variables, naive
regression models produce false positive rates above chance levels. This false positive rate
increases as the degree of autocorrelation increases. With strong spatial autocorrelation for
both outcomes and predictors, false positive rates reach as high as 77%. We find a slightly
lower false positive rate under strong cultural phylogenetic autocorrelation, though this
false positive rate is still greater than expected by chance (36%).

Most methods common in the literature do not reduce these high false positive rates. 223 With strong spatial autocorrelation for both outcome and predictor variables, false positive 224 rates remain above 50% when controlling for latitude, longitude, and language family fixed 225 effects (Figure 3). Similarly, Conley standard error corrections based on spatial and genetic 226 distances do not reduce false positive rates below 40% under strong spatial autocorrelation. 227 The most effective fixed effects methods are continent fixed effects, which continue to 228 produce a false positive rate of 35% under strong spatial autocorrelation, and controlling for the mean of the predictor variable within a 2000km radius, which eliminates false 230 positives under even strong spatial autocorrelation (6%). However, additional simulations 231 revealed that these reductions in false positive rates come at the cost of lower statistical 232 power (see Supplementary Figures S5 - S7). In additional simulations where the true 233 relationship between the predictor and outcome variable was known, power analyses 234 showed that both continent fixed effects and the 2000km radius control had less than 80% 235 power to detect moderate true correlations (r = 0.3) under strong spatial autocorrelation.

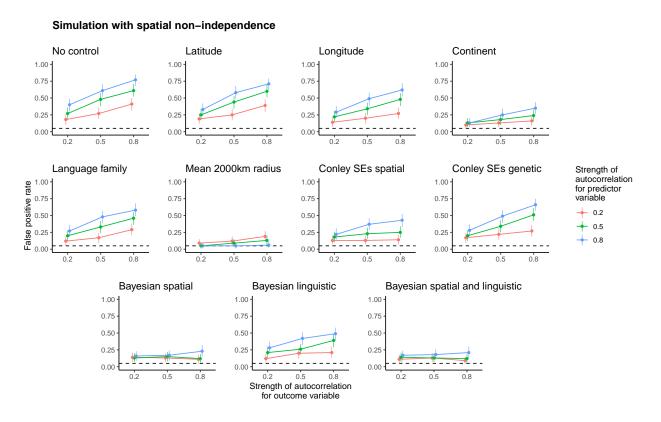


Figure 3. False positive rates for different methods of controlling for spatial non-independence in our simulation study. For simulated outcome and predictor variables, we systematically varied the strength of spatial autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination and fitted different models to each dataset. False positive rates were operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate the 5% false positive rate that is expected due to chance. SEs = standard errors.

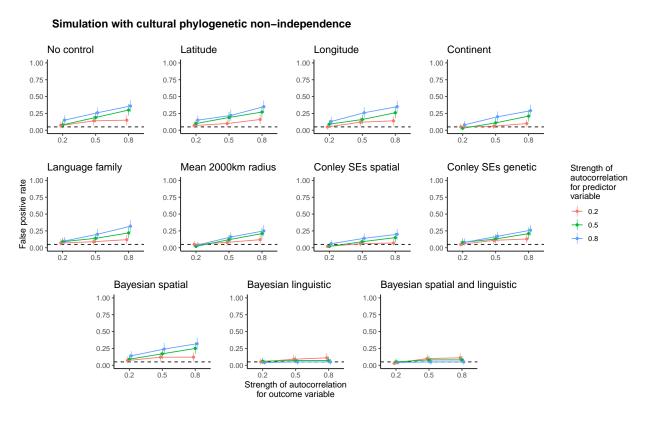


Figure 4. False positive rates for different methods of controlling for cultural phylogenetic non-independence in our simulation study. For simulated outcome and predictor variables, we systematically varied the strength of cultural phylogenetic autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination and fitted different models to each dataset. False positive rates were operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate the 5% false positive rate that is expected due to chance. SEs = standard errors.

By contrast, Bayesian spatial Gaussian process regression with longitude and latitude 237 strikes a balance between reducing false positives and retaining high statistical power to 238 detect true effects. This approach reduces false positives to 15% under moderate spatial 239 autocorrelation and 23% under strong spatial autocorrelation. Random effects models that 240 additionally account for linguistic proximity between nations perform equally well, though 241 models with only linguistic covariance continue to produce false positives. False positives 242 are not completely eliminated with these random effects models. Nevertheless, these 243 methods have at least 80% power to detect moderate (r = 0.3) and large (r = 0.5) true correlations between variables under all levels of spatial autocorrelation (see 245 Supplementary Figures S5 - S7). 246

In our simulation of cultural phylogenetic non-independence, we find that none of the 247 fixed effects methods reduce false positive rates (Figure 4). Controls for latitude, longitude, 248 continent fixed effects, the mean of the predictor variable in a 2000km radius, and Conley 240 standard error corrections based on spatial and genetic distances do little to change false 250 positive rates. Even language family fixed effects continue to produce a false positive rate 251 of 32% under strong cultural phylogenetic autocorrelation. By contrast, models with 252 random effects covarying according to linguistic proximity completely eliminate false 253 positives across all degrees of cultural phylogenetic autocorrelation. This approach is also 254 the only method that is able to detect large true correlations (r=0.5) with at least 80% 255 power (see Supplementary Figures S8 - S10). Random effects models that additionally 256 account for geographic proximity between nations perform equally well, though models 257 with only a spatial Gaussian process continue to produce false positives. 258

# Key findings in the literature are not robust to reanalysis with more rigorous methods

Our systematic review and simulation study have shown that controls for non-independence are rare in cross-national studies of economic development and cultural values, and when studies do attempt to control for non-independence, the methods
typically used are unable to sufficiently reduce false positive rates. This raises the worrying
possibility that the cross-national literature in economics and psychology is populated with
spurious relationships.

To determine how widespread this issue of spurious cross-national relationships might 267 be, we reanalysed a subset of twelve previous cross-national analyses from our systematic 268 review, controlling for spatial and cultural phylogenetic non-independence using global 269 geographic and linguistic proximity matrices. Out of the 100 papers included in our systematic review, primary or secondary data were publicly available for 47 papers. We 271 attempted to replicate key statistically significant cross-national correlations from these papers — mostly initial bivariate regression specifications without covariates (see Methods) 273 — and stopped when we had sampled a set of twelve analyses for which we were able to 274 replicate the original result. The final set included six analyses from our economic 275 development review<sup>53–58</sup> and six from our cultural values review<sup>13,14,16,59–61</sup> that had 276 available data and were able to be replicated. We pre-registered this set before running any 277 control models (https://osf.io/uywx8/). We controlled for non-independence by including 278 (1) a Gaussian process allowing nation random intercepts to covary according to a 279 geographic proximity matrix, and/or (2) nation random intercepts that covaried according 280 to a linguistic proximity matrix (see Supplementary Methods for full models). 281

Figure 5 visualises the results of our reanalysis (see Supplementary Table S5 for numerical results). Cross-national correlation effect sizes tended to reduce when controlling for statistical non-independence between nations, sometimes by as much as half of the original effect size. Overall, after controlling for non-independence, six out of twelve cross-national associations had 95% credible intervals that included zero. For the economic development analyses, four out of six cross-national relationships had 95% credible intervals including zero when controlling for spatial non-independence. For the cultural values analyses, two out of six cross-national relationships had 95% credible intervals including

zero when controlling for cultural phylogenetic non-independence. Supplementary Figure
 S11 shows these cross-national correlations plotted against the raw data.

292 Discussion

In a systematic literature review and simulation, we found that cross-national studies in economics and psychology rarely account for non-independence between nations, and, when they do, the methods they use are insufficient to reduce false positives in non-independent data. In a reanalysis of twelve cross-national correlations, we further showed that neglecting to account for non-independence has resulted in potentially spurious relationships in the published literature, with half of the correlations failing to replicate when controlling for spatial or cultural non-independence with more rigorous methods. These findings suggest that cross-national analyses in economics and psychology should be interpreted with caution until non-independence is sufficiently accounted for.

Our initial analyses add to and clarify existing evidence regarding the degree of 302 non-independence for national-level economic and cultural variables. One previous study 303 suggested that geographic proximity is more important than deep cultural ancestry in 304 explaining the distribution of human development across Eurasian nations, though the 305 authors noted that their small sample of 44 nations and regional focus limited their 306 statistical power<sup>62</sup>. By contrast, our global samples of over 160 nations revealed strong 307 cultural phylogenetic signal, as well as geographic signal, for the Human Development 308 Index, GDP per capita, and the Gini index of inequality. Another previous study found 309 that similarities in the cultural values of nations are predicted by linguistic, but not geographic, distances between those nations<sup>6</sup>. We find this same result for survival 311 vs. self-expression values, cultural tightness, and individualism, but for traditional 312 vs. secular values we find that both linguistic and geographic proximity are important 313 independent predictors of global variation. These findings emphasise the need to account 314 for both spatial and cultural phylogenetic non-independence in cross-national studies of 315

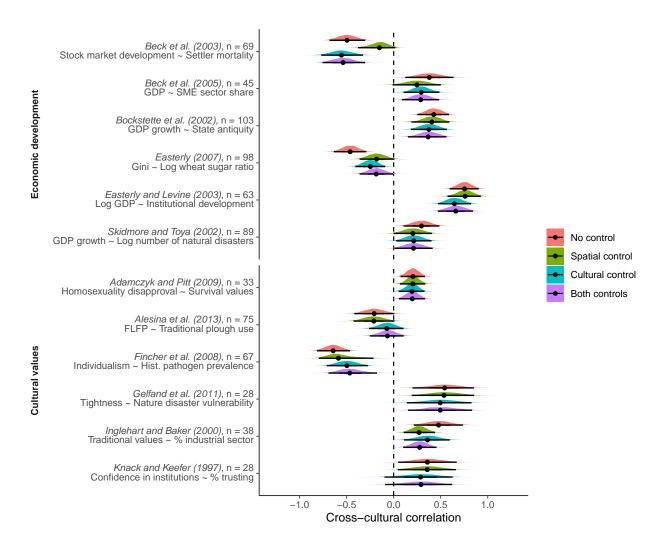


Figure 5. Posterior correlations from our reanalysis of twelve previous cross-national analyses. For each previous cross-national relationship, we plot the posterior slopes from a naive regression (red), a regression controlling for spatial non-independence (green), a regression controlling for cultural phylogenetic non-independence (blue), and a regression controlling for both spatial and cultural phylogenetic non-independence simultaneously (purple). All outcome and predictor variables are standardised. Most analyses are simple bivariate cross-national correlations, but Gelfand et al. (2011) is a partial correlation controlling for log gross national income and Adamczyk and Pitt (2009) is a multilevel model including several covariates. Coloured ridges are full posterior distributions, and points and black lines represent posterior medians and 95% credible intervals. GDP = gross domestic product. FLFP = female labour force participation.

economic development and cultural values.

Crucially, our systematic literature review and simulation study revealed that the 317 most commonly used controls for non-independence do not sufficiently deal with the issue. 318 In our simulations, controlling for either latitude or longitude did not reduce false positive 319 rates. This result calls into question the use of controls like distance to the equator to 320 account for non-independence in cross-national regression models, though these controls 321 may still be suitable to account for regional or latitudinal variation in climate, ecology, and 322 natural threats (e.g. pathogens), which we did not simulate. High false positive rates 323 persisted with Conley standard errors, despite recent claims that these standard error 324 corrections are sufficient to deal with spatial non-independence<sup>63</sup>. The simulation also confirmed the assertion that fixed effects for spatial or cultural groupings (e.g. continent or 326 language family fixed effects) are insufficient because non-independence still remains within 327 groupings<sup>41</sup>. This logic further applies to analyses that control for non-independence by 328 separately analysing different regions (e.g.<sup>64</sup>). Controlling for the mean of the predictor 329 variable within a 2000km radius<sup>46</sup> eliminated false positive rates in spatially autocorrelated 330 data, but had reduced statistical power to detect true associations. Across all model types 331 in our simulation, the only methods that markedly reduced false positive rates while 332 retaining high statistical power were the random effects models with covariance matrices. 333 The important advantage of these models is not that they are Bayesian per se (any of the 334 approaches used here could be implemented in a Bayesian framework), but rather that 335 they explicitly model covariance as a function of spatial or cultural distance. 336

There are other approaches to controlling for non-independence that we did not include in our simulation. For example, conditional autoregressive models<sup>31</sup> and generalised additive models<sup>65</sup> are approaches that can be applied in both frequentist and Bayesian frameworks. There are also alternative ways to operationalise cultural distances between nations beyond linguistic distances, including metrics like cultural fixation indices (e.g. cultural  $F_{ST}^{6,66}$ ), covariance based on genetic distances<sup>67</sup>, and phylogenetic distances

between religious traditions. We see merit in each of these approaches and the use of one over another will depend on the specific question at hand. We decided to focus on 344 linguistic distances in this study since language is a tangible socially-learned trait that has 345 previously been used to successfully track the effects of deep cultural ancestry on modern 346 national outcomes<sup>62,68,69</sup>. Future work should explore whether other approaches are 347 sufficient to reduce false positive rates in spatially and culturally non-independent data. In 348 addition to spatial proximity and shared cultural ancestry, we did not simulate other 349 sources of non-independence that potentially exist in real cross-national datasets, such as 350 modern connections between nations due to flows of people and information (e.g. flight 351 networks, social media networks) and shared histories of colonialism and capitalist modes 352 of production that have shaped today's global landscape<sup>70,71</sup>. Additional controls will be 353 required to ensure that these sources of non-independence do not confound cross-national inferences. 355

Ours is not the first review to show that studies are misapplying statistical methods 356 in ways that inflate false positive rates. For example, other literature reviews have shown 357 that studies in the social sciences tend to use small samples of participants<sup>72</sup>, treat ordinal 358 data as metric<sup>73</sup>, incorrectly handle missing values<sup>74</sup>, and ignore best practices in 359 meta-analyses<sup>75</sup>. Why do cross-national studies also rarely account for non-independence? 360 At the institutional level, one possibility is that such practices are incentivised because they 361 generate statistically significant relationships, which increase the probability that a study is 362 published<sup>72</sup>. Indeed, we found that controls for non-independence were less common among 363 articles published in high impact journals, suggesting that researchers are rewarded for 364 such practices. At the individual level, another possibility is that researchers outside of 365 anthropology and ecology are less aware of the problem, or believe that the problem does not apply to analyses of nations. Even if researchers appreciate the problem, they might not know of suitable controls or perceive the methods to be too complex.

These institutional- and individual-level barriers can be combatted. First,

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cross-national replication studies like ours and others<sup>40–44</sup>, combined with the 370 methodological reviews included in Registered Reports<sup>76</sup>, might change incentive structures 371 and encourage researchers to analyse the world's nations with more rigorous methods. 372 Second, more explicit descriptions of causal models could promote controls for 373 non-independence by clearly outlining the nature of confounding and the sources of 374 autocorrelation in cross-national data<sup>77</sup>. The causal model outlined in Supplementary 375 Figure S1 is a useful example, but individual studies must outline their own particular 376 causal assumptions, which may include further sources of non-independence and 377 confounding variables to control for (e.g. post-communist status, colony status). These 378 causal models can then be used to design tailored statistical estimation strategies. Indeed, 379 in our review, economists studying economic development dealt with national-level 380 non-independence more than psychologists studying cultural values, likely because economics studies tend to be lengthy statistical exercises that systematically incorporate or 382 exclude numerous variables in an attempt to infer causation. Third, the recent widespread accessibility of open source statistical software, such as the programming language Stan<sup>78</sup> 384 and the R package brms<sup>79</sup>, should promote the use of more rigorous methods to control for 385 non-independence. Using brms, for example, Bayesian Gaussian process regression is straightforward to conduct, requiring only longitude and latitude values for nations. We 387 have provided an online tutorial to help researchers apply these methods to their own 388 cross-cultural datasets (https://scottclaessens.github.io/blog/2022/crossnational/). 389

Until such changes are implemented and sufficient controls for non-independence are
the norm, existing cross-national correlations should be interpreted with caution. In our
reanalyses, we found that half of the cross-national correlations had 95% credible intervals
that included zero when controlling for spatial and/or cultural phylogenetic
non-independence. While these results are in line with previous reanalyses<sup>40</sup>, we note that
we are unable to outright reject the claims from these studies, since we only reanalysed the
first bivariate regression specifications presented in the papers<sup>63</sup>. More detailed sets of

reanalyses would be required to comprehensively challenge the claims from these specific 397 papers. Nevertheless, these reanalyses do show, more broadly, that the problem of 398 statistical non-independence applies to a wider range of national-level variables than those 399 identified by previous work, such as parasite stress and democratic outcomes<sup>41</sup>. Moreover, 400 given our finding that most studies in the current cross-national literature do not deal with 401 non-independence at all, our reanalyses raise the worrying possibility that this literature is 402 populated with spurious relationships. Future work should expand our set of reanalyses to 403 determine the extent of this problem in the literature. 404

We do not wish to dissuade researchers from conducting cross-national studies. On 405 the contrary, such work promises to deepen understanding of our world, including the causes and consequences of economic development and cultural values. Moreover, 407 cross-national studies allow social scientists to broaden their scope of study beyond 408 Western populations<sup>7</sup>, providing the representative samples necessary to test evolutionary 400 and socio-ecological theories of human behaviour<sup>8,80</sup>. But in order to minimise spurious 410 relationships in global datasets, we urge researchers to control for spatial and cultural 411 phylogenetic non-independence when reporting cross-national correlations. Nations are not 412 independent, and our statistical models must reflect this. 413

414 Methods

## 5 Geographic and cultural phylogenetic signal

To estimate the degree of spatial and cultural phylogenetic non-independence in
economic development and cultural values, we calculated geographic and cultural
phylogenetic signal for global measures of development and values. For economic
development variables, we retrieved longitudinal data on the Human Development Index
(1990 - 2019; n = 189 nations), gross domestic product per capita (1960 - 2021; n = 209
nations), annual percentage growth in gross domestic product per capita (1961 - 2021; n =

208 nations), and the Gini coefficient of income inequality (1967 - 2021; n = 167 nations). Human development data were retrieved from the United Nations Development 423 Programme (https://hdr.undp.org/en/content/download-data) and data for all other 424 economic development variables were retrieved from the World Bank 425 (https://data.worldbank.org/). For cultural values variables, we retrieved longitudinal data 426 on traditional vs. secular values and survival vs. self-expression values from the World 427 Values Survey<sup>16</sup> (1981 - 2019; n = 116 nations). We downloaded the full Integrated Values 428 Survey, which included all waves from the World Values Survey and the European Values 429 Survey, and computed the two dimensions of cultural values following procedures from 430 previous research<sup>16</sup>. Additionally, we retrieved cross-sectional data on cultural tightness (n 431 = 57 nations) and individualism (n = 97 nations) from previous work $^{13,81}$ . 432 To calculate geographic and cultural phylogenetic signal, we created two proximity 433 matrices for 269 of the world's nations: a geographic proximity matrix and a linguistic 434 proximity matrix. Geographic proximity was converted from logged geodesic distances 435 between nation capital cities. Linguistic proximity was calculated as the cultural proximity 436 between all languages spoken within nations, weighted by speaker percentages (see 437 Supplementary Methods). We included these matrices in Bayesian multilevel models, 438 allowing nation random intercepts to covary according to both geographic and linguistic 439 proximity simultaneously. These models were fitted with the R package brms<sup>79</sup> and 440 converged normally ( $\hat{R} < 1.1$ ). Estimates of geographic and cultural phylogenetic signal 441 were computed as the proportion of national-level variance in these models explained by geographic and linguistic proximity matrices.

## Systematic literature review

We exported two searches from Web of Science (https://www.webofknowledge.com/)
on 27<sup>th</sup> September 2021, restricting our searches to articles published between 1900 and
The first search was for the terms "economic development" AND ("cross-national"

OR "cross-cultural" OR "cross-country"), which returned 965 articles. The second search was for the terms "values" AND ("cross-national" OR "cross-cultural" OR "cross-cuntry"), which returned 6806 articles. Once exported, we ordered the articles by descending number of citations per year since initial publication, using citation counts reported by Web of Science.

We then systematically coded each article, in order, for inclusion in our review.

Articles were only included if: (1) they were judged to be relevant to economic

development or cultural values; (2) they were an original empirical research article; and (3)

they contained at least one analysis with national-level outcome or predictor variables. We

stopped when we had included 50 articles for the economic development review and 50

articles for the cultural values review.

Within each included article, we exhaustively coded every individual cross-national 459 analysis reported in the main text. We coded mainly correlation or regression analyses, and 460 explicitly excluded meta-analyses, factor analyses, measurement invariance analyses, 461 multidimensional scaling analyses, hierarchical clustering analyses, multiverse analyses, and 462 scale development / validation analyses. We also excluded analyses that compared only 463 two, three, four, five, or six nations. For each included analysis, we recorded the year, 464 impact factor of the journal (retrieved from https://jcr.clarivate.com/jcr/home), outcome 465 variable, all predictor variables, test statistic, p-value, number of nations, number of data points, model type, if the data were available, and whether and how the analysis attempted to control for non-independence.

We coded common attempts to control for non-independence between nations. These included: (1) any higher-level control variables for spatial regional groupings (e.g. continent fixed effects); (2) any geographic distance control variables (e.g. distance between capital cities, distance from equator, latitude); (3) any control variables capturing shared cultural history (e.g. former colony, legal origin fixed effects, linguistic history, cultural influence);

and (4) any other control variables, tests, or approaches that were deemed as attempts to
control for non-independence (e.g. eigenvector filtering<sup>82</sup>, controls for trade-weightings
between nations, cross-sectional dependence tests<sup>83</sup>, separate analyses for subsets of
nations). These were coded by the first author.

Once we had compiled our review database, we calculated the proportion of articles attempting to control for non-independence at least once. We also calculated the proportion of articles employing the different types of control listed above at least once: regional fixed effects, distance, shared cultural history, or other. For these proportions, we calculated 95% bootstrap confidence intervals with 1,000 bootstrap iterations. Additionally, we predicted the probability of an article attempting to control for non-independence at least once using Bayesian logistic regression, including in separate models log journal impact factor and year of publication as linear and spline predictors, respectively.

For individual analyses, we dealt with the nested nature of the data (analyses nested 486 within articles) by fitting Bayesian multilevel logistic regression models with review type 487 (economic development vs. cultural values) as the sole fixed effect and random intercepts 488 for articles. We fitted these models separately for overall attempts to control for 480 non-independence and split by method type. We report the adjusted proportions with 95% 490 credible intervals (see Results and Supplementary Figure S2). Additionally, we predicted 491 the probability of an analysis attempting to control for non-independence using Bayesian 492 multilevel logistic regression with random intercepts for articles. In separate models, we 493 included log journal impact factor and year of publication as linear and spline predictors, respectively. All Bayesian models were fitted with the brms R package<sup>79</sup>. Our priors were informed by prior predictive checks, and all models converged normally ( $\hat{R} < 1.1$ ).

### 497 Simulations

We simulated data for 236 nations i with varying degrees of spatial or cultural phylogenetic signal for outcome y and predictor x using the following generative model:

$$\begin{bmatrix} y_i \\ x_i \end{bmatrix} \sim \text{MVNormal} \begin{pmatrix} \alpha_y \\ \alpha_x \end{pmatrix}, \mathbf{S}$$

$$\alpha_y \sim \text{Normal}(0, \sqrt{\lambda} \cdot \Sigma)$$

$$\alpha_x \sim \text{Normal}(0, \sqrt{\rho} \cdot \Sigma)$$

$$\mathbf{S} = \begin{pmatrix} \sqrt{1-\lambda} & 0 \\ 0 & \sqrt{1-\rho} \end{pmatrix} \begin{pmatrix} 1 & r \\ r & 1 \end{pmatrix} \begin{pmatrix} \sqrt{1-\lambda} & 0 \\ 0 & \sqrt{1-\rho} \end{pmatrix}$$

where  $\Sigma$  is a correlation matrix proportional to either geographic or linguistic proximities between nations,  $\lambda$  and  $\rho$  are autocorrelation parameters that represent the expected spatial or cultural phylogenetic signal for outcome and predictor variables, respectively, and r is the true cross-national correlation between the variables after accounting for autocorrelation. Importantly, when r=0 in this simulation, we know that there is no direct causal relationship between y and x. Instead, any relationship between the two variables is merely the result of autocorrelation.

We set the autocorrelation parameters  $\lambda$  and  $\rho$  to either 0.2 (weak), 0.5 (moderate), or 0.8 (strong). We also initially set the true cross-national correlation to 0 in order to determine false positive rates, and then additionally set r to 0.1 (small effect), 0.3 (medium effect), and 0.5 (large effect) in order to determine statistical power to detect true effects. For each parameter combination, we simulated 100 datasets, resulting in 3600 datasets. Each dataset had 236 rows representing different nations, with the following associated data for each nation: latitude, longitude, continent (Africa, Asia, Europe, North America,
Oceania, or South America), language family of the nation's majority spoken language
(Afro-Asiatic, Atlantic-Congo, Austroasiatic, Austronesian, Eskimo-Aleut, Indo-European,
Japonic, Kartvelian, Koreanic, Mande, Mongolic-Khitan, Nilotic, Nuclear Trans New
Guinea, Sino-Tibetan, Tai-Kadai, Tupian, Turkic, or Uralic), the mean of the predictor
variable within a 2000km radius, and coordinates for genetic distances from a previous
study<sup>19</sup> (only available for 177 nations).

With the resulting simulated datasets, we standardised outcome and predictor 520 variables and fitted eleven different models: (1) naive regression without controls, (2) regression with latitude control, (3) regression with longitude control, (4) regression with 522 continent fixed effects, (5) regression with language family fixed effects, (6) regression 523 controlling for the mean of the predictor variable in a 2000km radius, (7) regression 524 employing Conley standard errors based on geographic distances, (8) regression employing 525 Conley standard errors based on genetic distances, (9) Bayesian regression including a 526 Gaussian process over latitudes and longitudes, (10) Bayesian regression including random 527 intercepts covarying according to linguistic proximity, and (11) Bayesian regression 528 including both a Gaussian process over latitudes and longitudes and random intercepts 529 covarying according to linguistic proximity. 530

Models employing Conley standard errors either required latitude and longitude 531 values or coordinates for genetic distances. To determine distance cutoffs, we employed an 532 approach recommended in previous work<sup>84</sup>: we fitted models with a range of feasible 533 distance cutoffs and retained the model with the largest standard error for the slope parameter. These models were fitted using the conleyeeq R package<sup>85</sup>. Bayesian models 535 were fitted using the brms R package<sup>79</sup>. Our choice of priors was based on prior predictive 536 simulation. All models converged normally ( $\hat{R} < 1.1$ ). Across all model types and 537 parameter combinations, we calculated the false positive rate as the proportion of models 538 that estimated slopes with a 95% confidence / credible interval excluding zero when r=0. We calculated statistical power as the proportion of models that estimated slopes with a 95% confidence / credible interval excluding zero when r > 0. We calculated 95% bootstrap confidence intervals for these false positive rates and statistical power estimates with 1,000 bootstrap iterations.

## Reanalyses

We searched the individual analyses from our systematic review for statistically significant cross-national correlations with available primary or secondary data. We restricted our search to one analysis per paper, and searched until we had a set of twelve analyses, six from economic development papers and six from cultural values papers, for which we were able to replicate the original result (i.e. find a cross-national correlation with the same sign and roughly the same effect size). We also ensured that at least one analysis was a multilevel model, with multiple observations per nation.

The twelve analyses that we settled on 13,14,16,53-61 were mostly bivariate cross-national 552 correlations, except for two. One analysis <sup>14</sup> additionally controlled for log gross national 553 income, and another analysis<sup>58</sup> is a multilevel model including random intercepts for 554 nations and several individual-level and national-level covariates (see Model 5 in original 555 paper). Before running any additional models, we pre-registered these twelve analyses on 556 the Open Science Framework on  $25^{\rm th}$  January 2022 (https://osf.io/uywx8/). We 557 endeavoured to keep the sample sizes of our reanalyses as close to the original analyses as possible, though there were some deviations (see Supplementary Table S6). Despite these 559 slight deviations from the original analyses, all models reported in Figure 5 are fitted to the same number of data points, meaning that any changes in effect sizes are solely due to 561 controlling for non-independence. 562

For each individual analysis, we ran four models: (1) a naive regression replicating
the original finding, (2) a regression including a Gaussian process allowing nation random

intercepts to covary according to a geographic proximity matrix from latitude and longitude values, (3) a regression including nation random intercepts that covaried according to a linguistic proximity matrix, and (4) a regression including both a geographic Gaussian process and nation random intercepts with linguistic covariance. See Supplementary Methods for full models.

We fitted these models using the *brms* R package<sup>79</sup>. Our choice of priors was based on prior predictive simulation. All models converged normally ( $\hat{R} < 1.1$ ), though for some models we resorted to using approximate Gaussian processes<sup>86</sup> to reach convergence.

# 573 Reproducibility

All data and code are accessible at our Open Science Framework repository

(https://osf.io/uywx8/). We used the *targets* R package<sup>87</sup> to create a reproducible data

analysis pipeline and the *papaja* R package<sup>88</sup> to reproducibly generate the manuscript.

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### **Author Contributions**

SC and QDA conceived of and designed the study. SC curated the data, produced all code for analysis and visualisation, and wrote the original draft of the manuscript. TK and QDA developed and compiled the geographic and linguistic distance matrices. QDA provided funding and input on manuscript preparation and revision. All authors reviewed and edited the final draft of the manuscript.

## Competing Interests

The authors declare no competing interests.

## Data Availability

All data can be found on the Open Science Framework: https://osf.io/uywx8/

# Code Availability

All code to reproduce the statistical analyses in this manuscript can be found on the Open Science Framework: https://osf.io/uywx8/

## Ethical Approval

Ethical approval was not required for the study as we analysed only secondary and simulated data.

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#### Supplementary Material

#### Supplementary Methods

#### Calculating global geographic and linguistic proximity matrices.

Geographic distance between two nations was calculated as the logged geodesic distance between nation capital cities (data from the R package *maps*; Brownrigg, 2018) using the R package *geosphere* (Hijmans, 2019). The geographic proximity matrix was computed as one minus the log geographic distance matrix scaled between 0 and 1.

Linguistic proximity between two nations was calculated as the cultural proximity between all languages spoken within those nations, weighted by speaker percentages. We acquired cultural proximity data by combining the language family trees provided by Glottolog v3.0 (Hammarström et al., 2017) into one global language tree (undated and unresolved). We calculated cultural proximity s between two languages j and k as the distance (in number of nodes traversed) of their most recent common ancestor i to the root of the tree, through the formula:

$$s_{jk} = \frac{n_r - n_i}{n_r}$$

where  $n_r$  is the maximum path length (in number of nodes traversed) leading to the pan-human root r, and  $n_i$  is the maximum path length leading to node i. We then combined these proximities with speaker data from Ethnologue 21 (Ethnologue, 2018) and compared every language spoken within those nations by at least 1 permille of the population, weighted by speaker percentages, through the formula:

$$w_{lm} = \Sigma \Sigma p_{lj} p_{mk} s_{jk}$$

where  $p_{lj}$  is the percentage of the population in nation l speaking language j,  $p_{mk}$  is the percentage of the population in nation m speaking language k, and  $s_{jk}$  is the proximity measure between languages j and k (Eff, 2008). The resulting linguistic proximity matrix was also scaled between 0 and 1 before analysis.

Bayesian models for reanalysis. We provide model formulae for our reanalyses of cross-national correlations, for a general bivariate case with standardised outcome Y and predictor X variables. In the naive regression model without controls for non-independence:

$$Y_i \sim \text{Normal}(\mu_i, \sigma)$$
  
 $\mu_i = \alpha + \beta X_i$   
 $\alpha \sim \text{Normal}(0, 0.4)$   
 $\beta \sim \text{Normal}(0, 0.4)$ 

The priors in this model were arrived at by prior predictive checks, with wider priors making predictions beyond the scale of standardised outcome variables and narrower priors being too informative.

 $\sigma \sim \text{Exponential}(5)$ 

To control for spatial non-independence, we add a Gaussian process to this model and feed it a scaled geographic distance matrix D based on Euclidean distances between latitude and longitude coordinates. This distance matrix is computed internally by the R package brms (Bürkner, 2017). The Gaussian process uses an exponentiated quadratic covariance kernel, the only covariance kernel currently supported by brms. The model formula is:

$$Y_{i} \sim \text{Normal}(\mu_{i}, \sigma)$$

$$\mu_{i} = \alpha + \kappa_{\text{NATION}[i]} + \beta X_{i}$$

$$\begin{pmatrix} \kappa_{1} \\ \kappa_{2} \\ \kappa_{3} \end{pmatrix} \sim \text{MVNormal} \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \end{pmatrix}, \mathbf{K}$$

$$\dots$$

$$\langle \kappa_{n} \rangle$$

$$\mathbf{K}_{ij} = sdgp^{2} \exp\left(-D_{ij}^{2}/(2lscale^{2})\right)$$

$$\alpha \sim \text{Normal}(0, 0.4)$$

$$\beta \sim \text{Normal}(0, 0.4)$$

$$\beta \sim \text{Normal}(0, 0.4)$$

$$\sigma \sim \text{Exponential}(5)$$

$$sdgp \sim \text{Exponential}(5)$$

$$lscale \sim \text{InverseGamma}(?,?)$$

where n is the number of nations, and  $D_{ij}^2$  reflects the squared Euclidean distances between latitude and longitude coordinates for the i-th and j-th nations. Notice that the inverse gamma prior on lscale is left undetermined. This is because the brms package intelligently tunes the prior for this parameter based on the covariates of the Gaussian process (see https://betanalpha.github.io/assets/case\_studies/gp\_part3/part3.html).

To control for cultural phylogenetic non-independence, we manually specify the covariance structure for nation random intercepts using a pre-computed linguistic proximity matrix L (see previous section). The covariance between two nations is assumed to be linearly proportional to the linguistic proximity between those nations. This assumption is justified if we assume that cultural traits evolve neutrally via Brownian motion along a language phylogeny. Although this is a conservative assumption, we follow this approach here rather than another Gaussian Process for several reasons: (1) to avoid model non-convergence when later including a spatial Gaussian Process in the same model, (2) to work around the lack of a coordinate system for linguistic distances, as opposed to latitude and longitude values for geographic distance, and (3) to showcase different ways that researchers can allow nations to covary to control for non-independence.

The non-centered parameterisation of this model is:

$$Y_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + z_{\text{NATION}[i]} \sigma_{\alpha} L + \beta X_i$$

 $\alpha \sim \text{Normal}(0, 0.4)$ 

 $\beta \sim \text{Normal}(0, 0.4)$ 

 $z_j \sim \text{Normal}(0, 1)$ 

 $\sigma_{\alpha} \sim \text{Exponential}(5)$ 

 $\sigma \sim \text{Exponential}(5)$ 

Finally, we can control for spatial and cultural phylogenetic non-independence simultaneously by including both a Gaussian process over latitude and longitude coordinates and nation random intercepts that covary according to linguistic proximity. The resulting model is as follows:

$$Y_{i} \sim \text{Normal}(\mu_{i}, \sigma)$$

$$\mu_{i} = \alpha + \kappa_{\text{NATION}[i]} + z_{\text{NATION}[i]} \sigma_{\alpha} L + \beta X_{i}$$

$$\begin{pmatrix} \kappa_{1} \\ \kappa_{2} \\ \kappa_{3} \\ \dots \\ \kappa_{n} \end{pmatrix} \sim \text{MVNormal} \begin{pmatrix} \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \\ \end{pmatrix}, \mathbf{K} \\ \dots \\ 0 \end{pmatrix}$$

$$\mathbf{K}_{ij} = sdgp^{2} \exp\left(-D_{ij}^{2}/(2lscale^{2})\right)$$

$$\alpha \sim \text{Normal}(0, 0.4)$$

$$\beta \sim \text{Normal}(0, 0.4)$$

$$z_{j} \sim \text{Normal}(0, 1)$$

$$\sigma_{\alpha} \sim \text{Exponential}(5)$$

$$\sigma \sim \text{Exponential}(5)$$

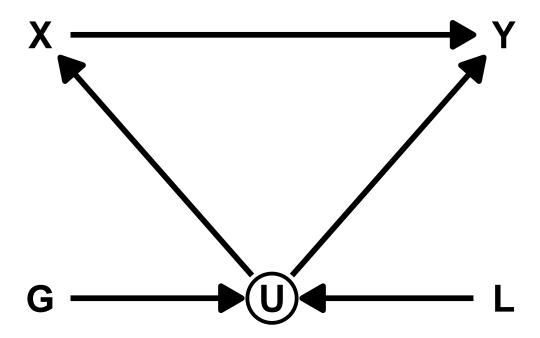
$$sdgp \sim \text{Exponential}(5)$$

$$lscale \sim \text{InverseGamma}(?,?)$$

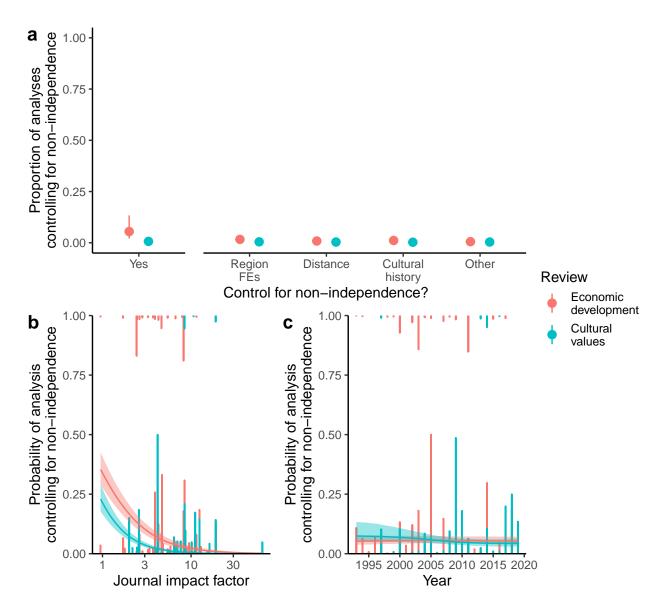
#### Supplementary Results

Further exploration of twelve reanalyses. To understand why some cross-national correlations were attenuated by controls for non-independence while others were robust, we further explored our fitted models for evidence of spatial and cultural autocorrelation. For each outcome variable, our Gaussian process models provided varying estimates of how quickly spatial autocorrelation declined with distance (Supplementary Figure S12). For example, in Skidmore and Toya<sup>58</sup> gross domestic product growth was only moderately spatially autocorrelated at 1,000 km distance (posterior median spatial autocorrelation at 1,000 km = 0.42, 95% CI  $[0.07 \ 0.90]$ ), whereas in Inglehart and Baker<sup>16</sup> traditional values were strongly spatially autocorrelated at the same distance (posterior median spatial autocorrelation at 1,000 km = 0.96, 95% CI [0.81 0.99]). We also found varying estimates of cultural phylogenetic signal (Supplementary Figure S13), with some outcome variables expressing low signal (e.g. confidence in institutions<sup>61</sup>; posterior median = 0.07, 95% CI [0.00 0.53]) and others expressing high signal (e.g. female labour force participation<sup>60</sup>; posterior median = 0.89, 95% CI [0.63 0.98]). Exploratory regressions provided suggestive evidence that stronger estimates of spatial autocorrelation or cultural phylogenetic signal resulted in a more pronounced reduction in the effect size when controlling for non-independence between nations (Supplementary Figure S14). However, these negative slopes for spatial autocorrelation (b = -0.19, 95% CI [-1.33 1.04]) and cultural phylogenetic signal (b = -0.28, 95\% CI [-1.26 0.70]) were both very uncertain, due to the small number of analyses and the posterior uncertainty in effect sizes and estimates of non-independence.

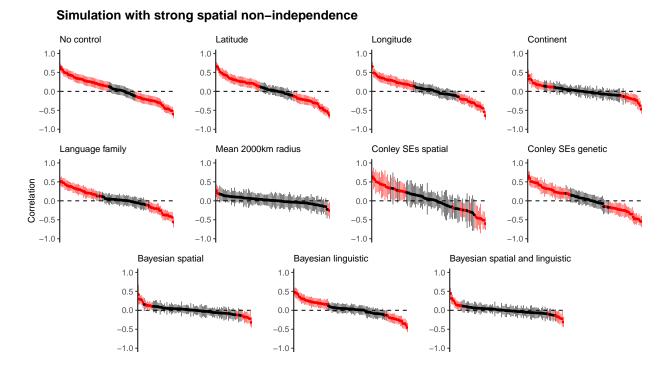
#### Supplementary Figures



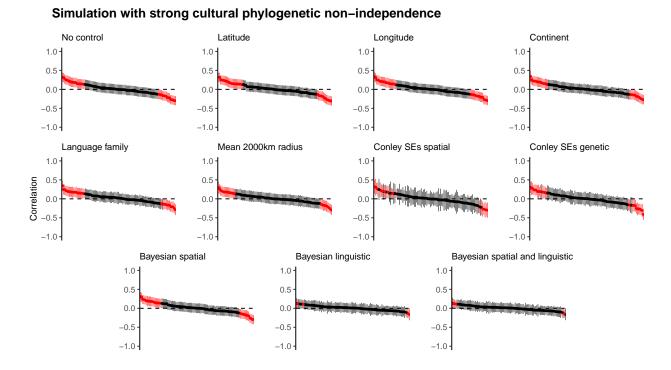
Supplementary Figure S1. A causal directed acyclic graph of spatial and cultural phylogenetic non-independence in cross-national studies. We are interested in estimating the direct effect of national-level exposure X on national-level outcome Y. But these variables are confounded by their common unobserved cause U. U is a stand-in for shared environmental, ecological, and geographic causes (e.g. climate, biodiversity, physical topography) and cultural and institutional causes (e.g. cultural norms, technologies, and institutions). In this causal model, we need to condition on U to estimate the direct path from X to Y, but we cannot since it is unobserved. However, geographic G and linguistic L relationships between societies influence U, since changing a nation's spatial distance to or shared cultural ancestry with other nations will change its environmental and cultural traits. We can thus use G and L to model the covariation between X and Y induced by U. Failing to do this and simply estimating the bivariate correlation between X and Y will produce spurious relationships and residuals that are spatially and culturally non-independent around the world.



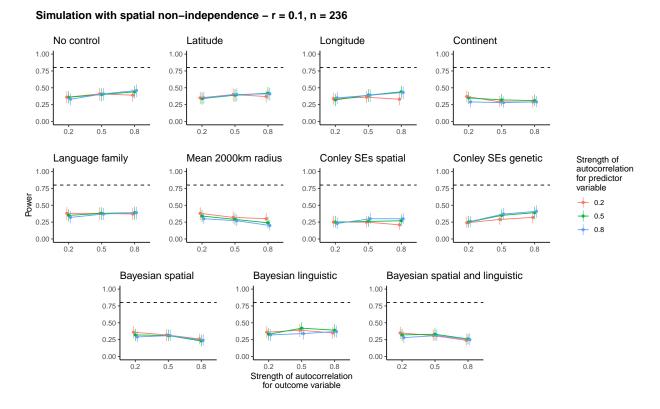
Supplementary Figure S2. Analysis-level results from systematic review of 100 highly-cited cross-national studies of economic development (red) and cultural values (blue). (a) Proportion of analyses accounting for non-independence, overall and split by common methods of controlling for non-independence. (b) The association between journal impact factor and the probability that an analysis accounts for non-independence. (c) Estimated trend over time for the probability that an analysis accounts for non-independence. Point ranges represent estimated proportions and 95% credible intervals. Lines and shaded areas are posterior median regression lines and 50% credible intervals from Bayesian multilevel models. Histograms represent relative counts for individual analyses that did (top) or did not (bottom) account for non-independence. Region FEs = region fixed effects.



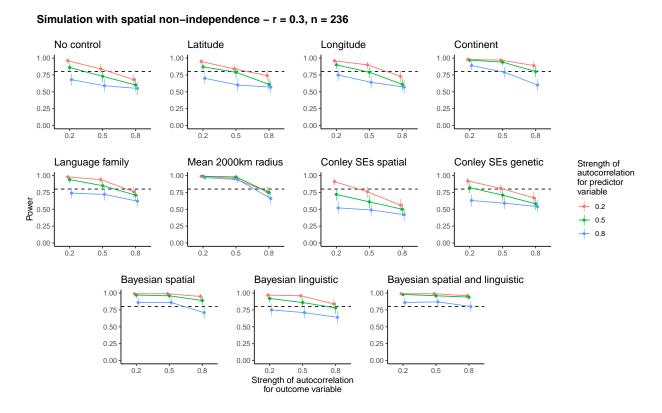
Supplementary Figure S3. Distribution of cross-national correlations from simulation study under strong spatial autocorrelation. In these simulations, the strength of spatial autocorrelation is set to 0.8 for both outcome and predictor variables, and the true correlation is set to 0. For frequentist regression models, point ranges represent correlation estimates and 95% confidence intervals. For Bayesian regression models, point ranges represent posterior means and 95% credible intervals. Correlations are ordered by effect size independently in each panel. Red point ranges indicate that the 95% confidence / credible interval excludes zero. Black point ranges indicate that the 95% confidence / credible interval includes zero.



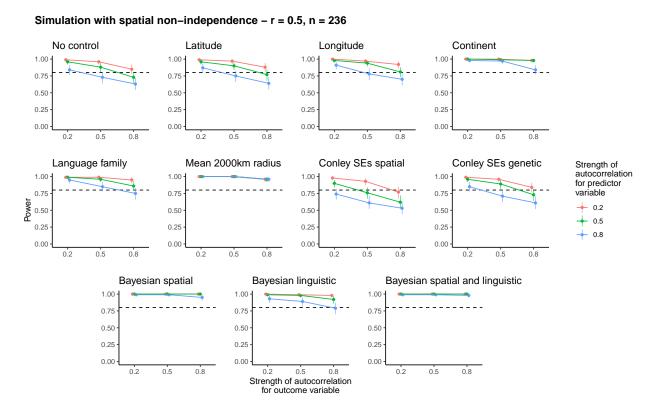
Supplementary Figure S4. Distribution of cross-national correlations from simulation study under strong cultural phylogenetic autocorrelation. In these simulations, the strength of cultural phylogenetic autocorrelation is set to 0.8 for both outcome and predictor variables, and the true correlation is set to 0. For frequentist regression models, point ranges represent correlation estimates and 95% confidence intervals. For Bayesian regression models, point ranges represent posterior means and 95% credible intervals. Correlations are ordered by effect size independently in each panel. Red point ranges indicate that the 95% confidence / credible interval excludes zero. Black point ranges indicate that the 95% confidence / credible interval includes zero.



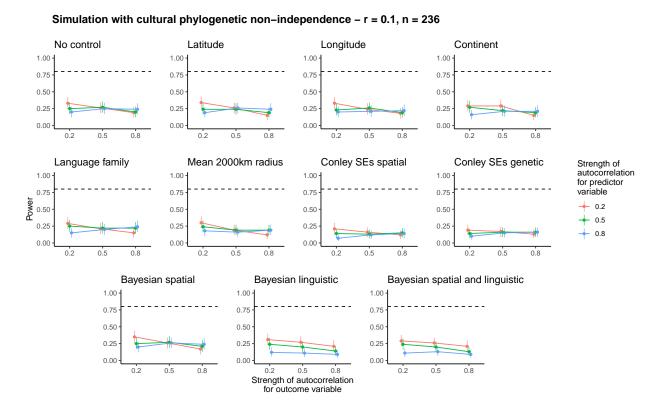
Supplementary Figure S5. Statistical power estimates for different methods of controlling for spatial non-independence in our simulation study, assuming a small true effect size (r = 0.1). For simulated outcome and predictor variables, we systematically varied the strength of spatial autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a small true correlation between variables (r = 0.1) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



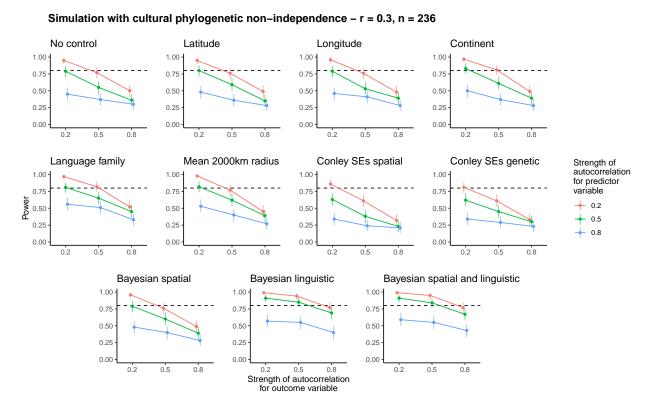
Supplementary Figure S6. Statistical power estimates for different methods of controlling for spatial non-independence in our simulation study, assuming a medium true effect size (r = 0.3). For simulated outcome and predictor variables, we systematically varied the strength of spatial autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a medium true correlation between variables (r = 0.3) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



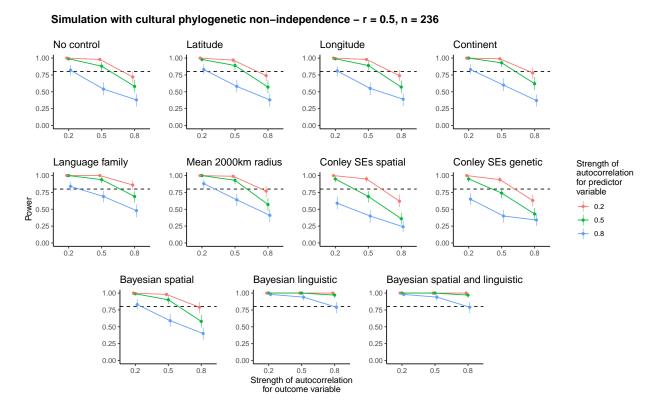
Supplementary Figure S7. Statistical power estimates for different methods of controlling for spatial non-independence in our simulation study, assuming a large true effect size (r = 0.5). For simulated outcome and predictor variables, we systematically varied the strength of spatial autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a large true correlation between variables (r = 0.5) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



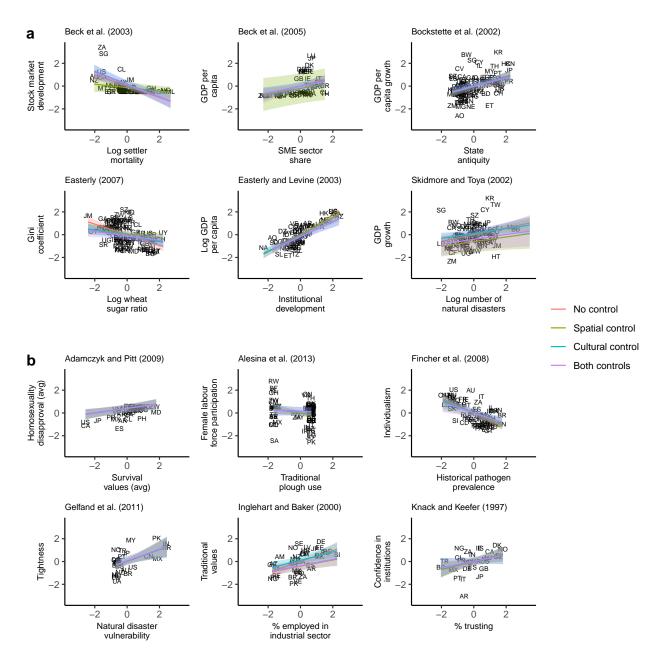
Supplementary Figure S8. Statistical power estimates for different methods of controlling for cultural phylogenetic non-independence in our simulation study, assuming a small true effect size (r=0.1). For simulated outcome and predictor variables, we systematically varied the strength of cultural phylogenetic autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a small true correlation between variables (r=0.1) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



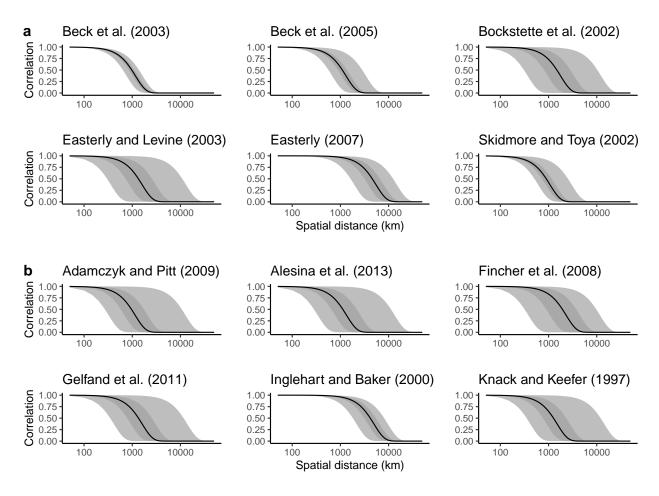
Supplementary Figure S9. Statistical power estimates for different methods of controlling for cultural phylogenetic non-independence in our simulation study, assuming a medium true effect size (r = 0.3). For simulated outcome and predictor variables, we systematically varied the strength of cultural phylogenetic autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a medium true correlation between variables (r = 0.3) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



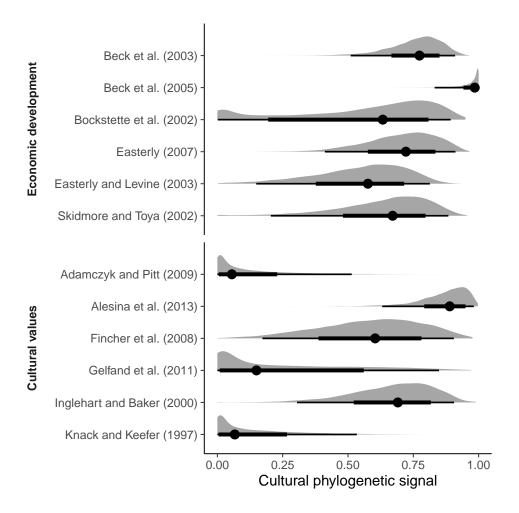
Supplementary Figure S10. Statistical power estimates for different methods of controlling for cultural phylogenetic non-independence in our simulation study, assuming a large true effect size (r=0.5). For simulated outcome and predictor variables, we systematically varied the strength of cultural phylogenetic autocorrelation, from weak (0.2) to moderate (0.5) to strong (0.8). We simulated 100 datasets per parameter combination assuming a large true correlation between variables (r=0.5) and fitted different models to each dataset. Statistical power was operationalised as the proportion of models that estimated a slope with a 95% confidence / credible interval excluding zero. Point ranges represent proportions and 95% bootstrap confidence intervals, and dashed lines indicate 80% power. SEs = standard errors.



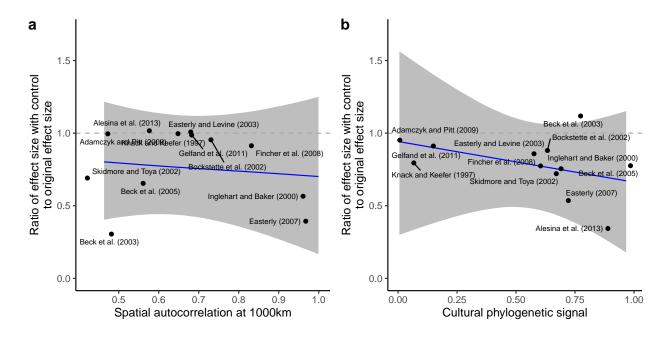
Supplementary Figure S11. Reanalysis models fitted to raw data, for economic development (a) and cultural values (b) studies. Data points are labelled using ISO 3166-1 alpha-2 letter country codes. In all reanalyses, outcome and predictor variables are standardised, making regression slopes comparable to Pearson's correlation coefficients. Lines and shaded areas represent posterior median regression lines and 95% credible intervals. For models with covariates (Adamzyck and Pitt 2009; Gelfand et al. 2011), marginal effects are presented holding all covariates at zero or their reference categories.



Supplementary Figure S12. Posterior estimates of Gaussian process functions mapping spatial autocorrelation onto geographic distance from our reanalyses of economic development (a) and cultural values (b) studies. The y-axis represents the amount of spatial autocorrelation between data points with increasing distance between those points on the x-axis (logged distance in kilometres). Lines and shaded areas represent median posterior spatial autocorrelation functions and 50% and 95% credible intervals.



Supplementary Figure S13. Posterior estimates of cultural phylogenetic signal from our reanalyses. Cultural phylogenetic signal is operationalised as the proportion of national-level variance explained by linguistic proximity between nations. Ridges are full posterior distributions, and points are posterior medians, and lines represent 50% and 95% credible intervals.



Supplementary Figure S14. The estimated degree of spatial and cultural phylogenetic non-independence predicts reductions in effect size in our reanalysis. (a) Higher estimated degrees of spatial autocorrelation at 1,000 km distance predict more pronounced reductions in effect sizes when controlling for non-independence. (b) Higher estimated levels of cultural phylogenetic signal predict more pronounced reductions in effect sizes when controlling for non-independence. In both panels, the y-axis represents the ratio of the effect size when controlling for spatial and cultural non-independence to the original effect size (from naive regression model), and the x-axis represents posterior median model estimates. Regression lines are plotted with 95% credible intervals.

### **Supplementary Tables**

### Supplementary Table S1

Geographic and cultural phylogenetic signal results for economic development and cultural values variables. Signal estimates reflect the proportion of national-level variance explained by geographic and linguistic covariance matrices. Bayes Factors (BF) reflect support for the hypothesis that the signal estimate differs from zero. HDI = Human Development Index; GDPpc = gross domestic product per capita.

Outcome	Geographic signal	Cultural phylogenetic signal
HDI CDPno	0.37, 95% CI [0.18, 0.59], BF > 100	0.62, 95% CI [0.40, 0.80], BF > 100 0.56, 95% CI [0.33, 0.78], BF > 100
GDPpc GDPpc growth	0.42, 95% CI [0.20, 0.66], BF > 100 0.65, 95% CI [0.09, 0.98], BF = 16.79	0.56, 95% CI [0.00, 0.70], BF = 1.16
Gini index	0.74, 95% CI [0.48, 0.97], BF > 100	0.25, 95% CI [0.02, 0.51], BF = 3.97
Traditional values Survival values	0.44, 95% CI [0.17, 0.76], BF > 100 0.20, 95% CI [0.01, 0.45], BF = 1.96	0.54, 95% CI [0.23, 0.79], BF > 100 0.78, 95% CI [0.53, 0.96], BF > 100
Tightness	0.09, 95%  CI  [0.00, 0.32], BF = 0.33	0.84, 95%  CI  [0.61, 0.97],  BF > 100
Individualism	0.26, 95%  CI  [0.00, 0.66], BF = 1.97	0.69, 95%  CI  [0.29, 0.96],  BF > 100

Supplementary Table S2

List of 100 papers included in literature review, sorted by annual rate of citations since publication.

Review	Reference	Citations per year
Cultural wal	Knack, S., & Keefer, P. (1997). Does social capital have an economic	
Cultural val-	payoff? A cross-country investigation. The Quarterly journal of eco-	121.50
ues	nomics, 112(4), 1251-1288.	
	Inglehart, R., & Baker, W. E. (2000). Modernization, cultural change,	
	and the persistence of traditional values. American sociological review,	119.33
	19-51.	
	Gelfand, M. J., Raver, J. L., Nishii, L., Leslie, L. M., Lun, J., Lim, B.	
	$\mathrm{C.,}\ \dots\ \&\ \mathrm{Yamaguchi,}\ \mathrm{S.}$ (2011). Differences between tight and loose	91.33
	cultures: A 33-nation study. science, $332(6033)$ , $1100-1104$ .	
	Alesina, A., Giuliano, P., & Nunn, N. (2013). On the origins of	
	gender roles: Women and the plough. The quarterly journal of eco-	57.12
	nomics, 128(2), 469-530.	
	Schmitt, D. P., Realo, A., Voracek, M., & Allik, J. (2008). Why can't a	
	man be more like a woman? Sex differences in Big Five personality traits	40.15
	across 55 cultures. Journal of personality and social psychology, 94(1),	49.15
	168.	
	Hofstede, G., & McCrae, R. R. (2004). Personality and culture revisited:	
	Linking traits and dimensions of culture. Cross-cultural research, $38(1)$ ,	35.53
	52-88.	
	Delhey, J., Newton, K., & Welzel, C. (2011). How general is trust in	
	"most people"? Solving the radius of trust problem. American Sociolog-	35.10
	ical Review, 76(5), 786-807.	
	Fincher, C. L., Thornhill, R., Murray, D. R., & Schaller, M. (2008).	
	Pathogen prevalence predicts human cross-cultural variability in indi-	20.22
	vidualism/collectivism. Proceedings of the Royal Society B: Biological	33.38
	Sciences, 275(1640), 1279-1285.	

Table S2 continued

Review	Reference	Citations per year
	Schneider, S. L. (2008). Anti-immigrant attitudes in Europe: Outgroup	
	size and perceived ethnic threat. European Sociological Review, 24(1),	30.54
	53-67.	
	Franzen, A., & Meyer, R. (2010). Environmental attitudes in cross-	
	national perspective: A multilevel analysis of the ISSP 1993 and	30.27
	2000. European sociological review, 26(2), 219-234.	
	Steenkamp, J. B. E., Ter Hofstede, F., & Wedel, M. (1999). A cross-	
	national investigation into the individual and national cultural an-	29.32
	tecedents of consumer innovativeness. Journal of marketing, $63(2)$ , $55-69$ .	
	Paxton, P. (2002). Social capital and democracy: An interdependent	20.21
	relationship. American sociological review, 254-277.	29.21
	Santos, H. C., Varnum, M. E., & Grossmann, I. (2017). Global increases	20.25
	in individualism. Psychological science, $28(9)$ , $1228-1239$ .	28.25
	Van Zanten, B. T., Van Berkel, D. B., Meentemeyer, R. K., Smith, J.	
	W., Tieskens, K. F., & Verburg, P. H. (2016). Continental-scale quan-	27.00
	tification of landscape values using social media data. Proceedings of the	21.00
	National Academy of Sciences, 113(46), 12974-12979.	
	Beugelsdijk, S., & Welzel, C. (2018). Dimensions and dynamics of na-	
	tional culture: Synthesizing Hofstede with Inglehart. Journal of cross-	24.00
	cultural psychology, 49(10), 1469-1505.	
	Stephan, U., & Uhlaner, L. M. (2010). Performance-based vs socially	
	supportive culture: A cross-national study of descriptive norms and en-	23.91
	trepreneurship. Journal of International Business Studies, $41(8),\ 1347$	20.01
	1364.	
	Dakhli, M., & De Clercq, D. (2004). Human capital, social capital, and	
	innovation: a multi-country study. Entrepreneurship & regional devel-	23.53
	opment, $16(2)$ , $107-128$ .	

Table S2 continued

Review	Reference	Citations per year
	Adamczyk, A., & Pitt, C. (2009). Shaping attitudes about homosex-	
	uality: The role of religion and cultural context. Social Science Re-	21.75
	search, 38(2), 338-351.	
	Mayda, A. M., & Rodrik, D. (2005). Why are some people (and	
	countries) more protectionist than others?. European Economic Re-	21.69
	view, 49(6), 1393-1430.	
	Johnson, T., Kulesa, P., Cho, Y. I., & Shavitt, S. (2005). The relation	
	between culture and response styles: Evidence from 19 countries. Journal	20.62
	of Cross-cultural psychology, 36(2), 264-277.	
	Tam, K. P., & Chan, H. W. (2018). Generalized trust narrows the gap	
	between environmental concern and pro-environmental behavior: Mul-	20.00
	tilevel evidence. Global Environmental Change, 48, 182-194.	
	Kuppens, P., Realo, A., & Diener, E. (2008). The role of positive and	
	negative emotions in life satisfaction judgment across nations. Journal	19.62
	of personality and social psychology, 95(1), 66.	
	Tam, K. P., & Chan, H. W. (2017). Environmental concern has a weaker	
	association with pro-environmental behavior in some societies than oth-	10.75
	ers: A cross-cultural psychology perspective. Journal of Environmental	18.75
	Psychology, 53, 213-223.	
	Miska, C., Szőcs, I., & Schiffinger, M. (2018). Culture's effects on corpo-	
	rate sustainability practices: A multi-domain and multi-level view. Jour-	17.67
	nal of World Business, 53(2), 263-279.	
	Oishi, S., Diener, E., Lucas, R. E., & Suh, E. M. (2009). Cross-cultural	
	variations in predictors of life satisfaction: Perspectives from needs and	16.00
	values. In Culture and well-being (pp. 109-127). Springer, Dordrecht.	
	Malka, A., Soto, C. J., Inzlicht, M., & Lelkes, Y. (2014). Do needs	
	for security and certainty predict cultural and economic conservatism?	16.00
	A cross-national analysis. Journal of personality and social psychol-	16.00
	ogy, 106(6), 1031.	

Table S2 continued

Review	Reference	Citations per year
	Eom, K., Kim, H. S., Sherman, D. K., & Ishii, K. (2016). Cultural	
	variability in the link between environmental concern and support for	15.80
	environmental action. Psychological Science, 27(10), 1331-1339.	
	Larsen, C. A. (2008). The institutional logic of welfare attitudes: How	
	welfare regimes influence public support. Comparative political stud-	15.62
	ies, 41(2), 145-168.	
	Sortheix, F. M., & Schwartz, S. H. (2017). Values that underlie and	
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Supplementary Table S3

False positive rates in simulation with varying levels of spatial autocorrelation for the predictor variable  $(\rho)$  and for the outcome variable  $(\lambda)$ , assuming that the true correlation is 0. Numbers represent the total number of analyses, out of 100, that estimated a slope with a 95% confidence / credible interval excluding zero, falsely inferring a relationship when none is present.

		$\rho = 0.2$			$\rho = 0.5$			$\rho = 0.8$	
Model	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$
No control	18	27	41	27	48	61	40	61	77
Latitude	19	25	39	25	44	09	33	58	71
Longitude	14	20	27	22	34	48	29	49	62
Continent	10	13	16	13	18	24	13	25	35
Language family	12	17	29	20	33	46	27	48	58
Mean 2000km radius	6	12	19	2	6	13	2	5	9
Conley SEs spatial	13	13	14	18	23	25	22	37	43
Conley SEs genetic	17	22	27	20	34	51	28	49	99
Bayesian spatial	14	13	11	13	15	12	16	17	23
Bayesian linguistic	12	20	21	21	26	39	28	42	49
Bayesian spatial & linguistic	11	13	6	14	13	12	17	18	21

Supplementary Table S4

total number of analyses, out of 100, that estimated a slope with a 95% confidence / credible interval excluding variable  $(\rho)$  and for the outcome variable  $(\lambda)$ , assuming that the true correlation is 0. Numbers represent the False positive rates in simulation with varying levels of cultural phylogenetic autocorrelation for the predictor zero, falsely inferring a relationship when none is present.

		$\rho = 0.2$			$\rho = 0.5$			$\rho = 0.8$	
Model	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$	$\lambda = 0.2$	$\lambda = 0.5$	$\lambda = 0.8$
No control	7	14	15	$\infty$	19	30	15	26	36
Latitude	2	10	16	10	19	27	15	22	35
Longitude	5	12	14	6	16	26	13	26	35
Continent	5	9	10	3	11	21	~	20	29
Language family	2	6	12	6	14	22	10	20	32
Mean 2000km radius	5	<b>∞</b>	12	2	12	21	4	16	25
Conley SEs spatial	2	9	2	2	6	15	9	14	20
Conley SEs genetic	5	11	13	∞	13	21	8	17	26
Bayesian spatial	2	12	12	6	17	25	14	24	32
Bayesian linguistic	5	6	11	9	2	2	4	2	5
Bayesian spatial & linguistic	က	10	11	ಬ	$\infty$	$\infty$	4	ಬ	ರ

Supplementary Table S5

Estimated cross-national correlations from our reanalysis of twelve previous cross-national analyses. Numbers are posterior median slopes (equivalent to Pearson's r correlation coefficients) with 95% credible intervals.

Analysis	No control	Spatial control	Cultural control	Both controls
Adamczyk and Pitt (2009)	[0.07, 0.07]	95% CI [ 0.07,	95% CI [ 0.05,	95% CI [ 0.06,
Alesma et al. $(2013)$ Beck et al. $(2003)$	-0.21, 93% C1 [-0.42, 0.01] -0.50, 95% CI [-0.68, -0.30]		-0.07, 95% CI [-0.20, 0.11] -0.56, 95% CI [-0.77, -0.33]	-0.01, 95% CI [-0.23, 0.11] -0.54, 95% CI [-0.76, -0.30]
Beck et al. (2005)		95% CI [-0.01,	95% CI [ 0.10,	95% CI [ 0.09,
Bockstette et al. (2002)		95% CI [ 0.19,	95% CI [ 0.18,	95% CI [ 0.15,
Easterly and Levine (2003)		95% CI [ 0.57,	95% CI [ 0.47,	95% CI [ 0.47,
Easterly $(2007)$	٠.	95% CI [-0.36,	95% CI [-0.41,	95% CI [-0.36,
Fincher et al. (2008)	٠.	95% CI [-0.80,	95% CI [-0.71,	95% CI [-0.69,
Gelfand et al. (2011)	_	95% CI [ 0.19,	95% CI [ 0.14,	95% CI [ 0.15,
Inglehart and Baker (2000)		95% CI [ 0.11,	95% CI [ 0.11,	95% CI [ 0.10,
Knack and Keefer (1997)		95% CI [ 0.04,	95% CI [-0.10,	95% CI [-0.09,
Skidmore and Toya (2002)	_	95% CI [ 0.00,	95% CI [ 0.03,	95% CI [ 0.01,

Supplementary Table S6
List of sample size deviations between original analyses and reanalyses.

Analysis	Original N	Reanalysis N	Reason for deviation
Adamczyk and Pitt (2009)	33	33	N/A
			Original analysis used imputed values for tra-
			ditional plough use which were unavailable to
			us. For our reanalysis, we collected data from
Alesina et al. (2013)	>177	75	Tables A4 and A9 and ran our regression on
			the 75 nations with available data for female
			labour force participation in 2000 and tradi-
			tional plough use.
T. 1. (2222)			We removed one nation (Zaire) as it was not
Beck et al. (2003)	70	69	present in our linguistic distance matrix.
Beck et al. (2005)	45	45	N/A
			After manually linking state history data from
Bockstette et al. (2002)	94	103	Appendix A with world development indica-
			tors, we had complete data for 103 nations.
			After manually linking log GDP per capita
			1995 data from Appendix Table A2 in Ace-
Easterly and Levine	<b>T</b> 0	40	moglu et al. (2001) with an institutions index
(2003)	72	63	self-constructed from world governance indi-
			cators, we only had complete data for 63 na-
			tions.
			After manually linking log wheat-sugar ratio
F . 1 (222 <b>7</b> )	440	0.0	data from Appendix A with longitudinal Gini
Easterly (2007)	118	98	data, we only had complete data for 98 na-
			tions.
			We combined England and Northern Ireland
Fincher et al. (2008)	68	67	into a single nation (United Kingdom) for the
			reanalysis.

Table S6 continued

Analysis	Original N	Reanalysis N	Reason for deviation
			We combined East and West Germany into a
C 16 1 4 1 (2011)	20	90	single nation (Germany) and removed a high
Gelfand et al. (2011)	30	28	leverage point with a substantially higher dis-
			aster rate than other nations (Venezuela).
			After manually linking traditional values data
T 11 / 1 D 1			from the World Values Survey with data on
Inglehart and Baker	er 65	38	percentage total employment in industry from
(2000)			World Development Indicators, we only had
			complete data for 38 nations.
Knack and Keefer (1997)	28	28	N/A
Skidmore and Toya (2002)	89	89	N/A

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