

National-scale changes in crop diversity through the Anthropocene

Rachel O Mariani, Marc W Cadotte, Marney E Isaac, Denis Vile, Cyrille Violle, Adam R Martin

To cite this version:

Rachel O Mariani, Marc W Cadotte, Marney E Isaac, Denis Vile, Cyrille Violle, et al.. National-scale changes in crop diversity through the Anthropocene. Scientific Reports, 2021 , 11, $10.1038/s41598-$ 021-99728-5 . hal-03382443

HAL Id: hal-03382443 <https://hal.inrae.fr/hal-03382443v1>

Submitted on 18 Oct 2021

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

[Distributed under a Creative Commons Attribution 4.0 International License](http://creativecommons.org/licenses/by/4.0/)

scientific reports

Check for updates

National‑scale changes OPEN in crop diversity through theAnthropocene

Rachel O. Mariani1 , Marc W. Cadotte2 , Marney E. Isaac1 , DenisVile3 , CyrilleViolle4 & Adam R. Martin1*

Expansion of crops beyond their centres of domestication is a defning feature of the Anthropocene Epoch. This process has fundamentally altered the diversity of croplands, with likely consequences for the ecological functioning and socio-economic stability of agriculture under environmental change. While changes in crop diversity through the Anthropocene have been quantifed at large spatial scales, the patterns, drivers, and consequences of change in crop diversity and biogeography at nationalscales remains less explored. We use production data on 339 crops, grown in over 150 countries from 1961 to 2017, to quantify changes in country-level crop richness and evenness. Virtually all countries globally have experienced signifcant increases in crop richness since 1961, with the early 1980s marking a clear onset of a ~ 9-year period of increase in crop richness in countries worldwide. While these changes have increased the similarity of diversity of croplands among countries, only half of countries experienced increases in crop evenness through time. Ubiquitous increases in crop richness within nearly all countries between 1980 and 2000 are a unique biogeographical feature of the Anthropocene. At the same time, we detected opposing changes in crop evenness, and only modest signatures of increased homogenization of croplands among countries. Therefore context-dependent and, at least, national-scale assessments are needed to understand and predict how changes in crop diversity infuence agricultural resistance and resilience to environmental change.

The Anthropocene Epoch defines the current epoch of geologic time, where human activity is the dominant force shaping Earth's abiotic and biotic environmental processes, systems, and cycles¹⁻³. This novel geological epoch is defined in part by human-caused changes to Earth's biogeography e.g.⁴, including the introduction of non-native or domesticated species e.g.⁵, and climate- and land-use change-induced shifts in species distributions e.g.⁶. However, arguably the most pronounced human-mediated changes in species biogeography defining the Anthropocene, albeit still largely understudied⁷, is the deliberate spread of crops outside of their centres of domestication into other parts of the world⁸⁻¹⁰. Large-scale anthropogenic influences on crop biogeography first emerged during the Columbian Exchange, which is a period in the early sixteenth century defned by massive exchanges of crop plants and animals between West Africa, Europe, and the Americas, during the time of New World colonization by Europeans¹⁰. During this time there was major movement of crops such as corn and beans from the Americas to Europe, while other crops such as wheat and barley were transported the opposite direction^{9,10}.

While research defining the scientific basis of the Anthropocene have focused on the Columbian Exchange², analyses of contemporary global food production trends have demonstrated that there have been more recent (i.e., post-1950) and widespread movement of new crops—i.e., those previously not present in large, industrial agricultural lands—into nearly all regions of the world⁸. Specifically since the early 1960s, the range of crops cultivated in agricultural lands across virtually all continents and regions, has changed in remarkably similar patterns: (1) in the 1960s continents experienced a period of little change in the number of new crops being cultivated in large (i.e., industrial) agricultural lands; this was followed by (2) a period of rapid increase in the number of crop groups being cultivated, beginning in the late 1970s and continuing through to the 1980s; and fnally, (3) there exists a period of little change or levelling of in the number of crops being cultivated beginning

¹Department of Physical and Environmental Sciences, University of Toronto Scarborough, Toronto, Canada. ²Department of Biological Sciences, University of Toronto Scarborough, Toronto, Canada. ³Laboratoire d'Ecophysiologie des Plantes sous Stress Environnementaux (LEPSE, UMR759), Institut National de la Recherche Agronomique (INRA), Université de Montpellier, Montpellier, France. ⁴CEFE, CNRS, EPHE, IRD, University Montpellier, University Paul Valéry, 34293 Montpellier, France.[⊠]email: adam.martin@utoronto.ca

in the 1990s and persisting through to present day⁸. These changes have drastically shaped not only the diversity of agricultural lands worldwide, but have greatly infuenced global food supplies and security, diets, and agricultural economics¹¹⁻¹⁷.

Coinciding with these human-mediated increases in the number of crops being cultivated in agricultural lands, this time period (particularly the 1980s) is also marked by an overall increase in the similarity of crop composition globally⁸. As an example of this homogenization of croplands at supra-national scales, wheat, maize, soy, and rice, now dominate over 50% of the world's agricultural lands¹⁸. These changes have contributed to major shifs in the world's agricultural landscapes and economies, including the homogenization of global food supplies and diets¹⁴, greater interdependency in agricultural trade between countries to maintain food security and potentially increases susceptibility of agricultural lands to pests, diseases, and climate stressors¹⁹.

Researchers have argued that the magnitude of these contemporary changes in crop biogeography, indicates that the 1980s are also a notable marker of the onset of the Anthropocene8 . However, this aspect of a crop biogeography-based line of evidence supporting the Anthropocene hypothesis remains limited to research on changes in crop diversity at very large supranational regional, continental, or global spatial scales 8 , or a small number of national-scale analyses (e.g., within the United States²⁰). This is despite reason to expect that the timing and rates of change in crop diversity is likely to difer widely among countries. For example previous work has shown that regional-scale agricultural policy initiatives including the Caribbean Basin Initiative and the North American Free Trade Agreement, were a primary catalyst for changes in crop diversity in the Caribbean and North American, respectively8,21. Yet such regional initiatives are not necessarily common in many agricultural economies. For instance, throughout regions of Africa, country-specifc structural readjustment policies and colonial histories, but not necessarily region-scale policy initiatives, have played a key role in determining crop composition on agricultural lands²². Similarly, in certain countries Western philanthropic organizations have played a signifcant role in transferring crop technologies including disease resistant crops and pressuring shifs to cash crops, which have had unintended consequences for national agrobiodiversity $22,23$. This and a multitude of similar examples (e.g., $21,24-26$) indicate that country-scale analyses are needed for a nuanced understanding of how crop diversity and composition has changed through the Anthropocene.

Supranational assessments also preclude nuanced analyses of the correlates or consequences of crop diversity change. Specifcally, much of the structural adjustment policies that emphasized enhanced balance of trade and exports were imposed on the agricultural economies of developing nations throughout the 1980s e.g.^{21,25}. Since these policies tended to focus on cultivation of new crops for international export markets (e.g., the introduction of cocoa in India and oil palm in Peru in the early and mid-1970s), one might expect that changes in agricultural diversity profles (i.e., the timing, duration, and rate of change in crops being cultivated) vary systematically with socio-economic development indices. More specifcally, one may hypothesize that a certain socio-economic group (e.g., developing nations with lower Human Development Index (HDI) values) to have broadly shown similar timing and rates of change in crop diversity, compared to countries with higher HDI.

Finally, one might expect that the number of crops cultivated within a given country also varies systemically with latitude, across a latitudinal gradient in crop diversity¹⁵. This pattern is likely to emerge as a function of multiple factors including (1) seasonality and limited growing conditions towards the poles²⁷, and (2) the centres of crop domestication being disproportionately situated in tropical and sub-tropical regions^{28,29}. Moreover, if a latitudinal gradient in the number of crops cultivated in agricultural lands does exist, these patterns may have been drastically altered by the incorporation of new crops into agricultural lands that began in the 1980s $^{\rm 8}.$

National-scale analyses of crop cultivation are needed to test these questions surrounding crop diversity change, and its role in defning the Anthropocene. To address this, we used crop production data collected by the Food and Agricultural Organization $(FAO)^{18}$ to execute national-scale analyses of changes in crop diversity. Tis analysis evaluated production data for 339 crop species groups, grown in 201 United Nations-recognized countries, over the past 56 years (from 1961 to 2017), resulting in a dataset of over~2.36 million data points. Our analysis was designed to address the following: (1) How do patterns of change in the diversity of crops grown on agricultural lands, vary across countries in recent decades? (2) Have changes in country-scale crop diversity resulted in a detectable signal of "homogenization" across the world's agricultural lands? (3) Are patterns of change in crop diversity across countries, systematically related to country-scale socio-economic indicators? (4) Does the number of crops cultivated across countries follow a latitudinal diversity gradient? And if so, (5) has the introduction of new crops into diferent countries altered this gradient?

Results

Changes in crop richness within countries through time. Our results are based on trends and three diferent "Indicator" values inferred from a piecewise modelling framework, which is described in detail in the Methods and presented conceptually in Fig. 1. Across all countries we detected signifcant changes, largely increases, in crop commodity group richness over time $(r^2 = 0.356 - 0.998$ across 165 countries, $p < 0.01$ in all cases; Fig. 2, Table S1). Across 165 countries for which piecewise models converged, the average initial onset of crop group richness changes (Indicator 1) occurred in 1983 (±9.2 years s.d.; Fig. 2A,B). Te onset of changes in crop group richness occurred at the earliest in 1962 (in India), with the most recent onset of crop group richness change beginning in 2007 (in Serbia; Fig. 2A,B). Richness began to change at or afer 1980 in 113 countries, of which 71% (or 80 countries) show changes in crop richness beginning from 1980 to 1989 (Fig. 2A; Table S1).

The period of crop richness change (Indicator 2) lasted approximately 9 years on average $(\pm 8.9$ years m.a.d.; Fig. 2C,D). Nineteen countries had a period of richness change lasting only one year, while nine countries experienced a more prolonged period of crop richness change lasting≥30 years (Fig. 2C,D). National changes in richness began to saturate (i.e., ψ 2 in Eq. (2), or breakpoint 2 in Fig. 1) on average in 1995 (\pm 8.3 years s.d.), with richness increases levelling off in 117 countries prior to the turn of the century at or before 1999 (Fig. 2C,D).

2

Figure 1. Schematic representation of three indicators of change in crop commodity group diversity, derived from piecewise models predicting crop commodity group richness as a function of year. Detailed explanations of Indicator 1–3 are presented in the Methods section. Data shown here as the example is from Canada, with black dots representing the number of commodities reported by the Food and Agricultural Organization, for a given year. Black trendline represents the piecewise model ft, gray bands represent the 95% confdence limits surrounding the model, and red lines represent model parameters and indicators derived from the model. *Note*: the fgure presented here demonstrates changes in crop commodity group richness (*S*), though this framework was also employed for assessing change in crop group evenness (*J′*).

Over the period of change (i.e., between ψ 1 and ψ 2, Fig. 1), crop richness increased in 151 of 165 countries evaluated. Across all countries, richness increased (Indicator 3) on average by 0.8 species per year $(\pm 0.9 \text{ m.a.d.};$ Fig. 2E,F). Only 13 countries reported decreases in richness per year, with these countries showing declines of 0.4±0.6 crop groups per year on average (Fig. 2E,F, Table S1). In all of our datasets, Indicators 1–3 associated with changes in *S* were unrelated to the total area of a country (linear regression $r^2 = 0.0-0.004$, $p \ge 0.888$), or the area of a country under crop cultivation (linear regression r^2 = 0.0–0.01, $p \ge 0.208$).

Across the 164 countries for which data was available in both 1961 and 2017, crop richness varied signifcantly as a function of latitude in similar patterns in both years (Table 1, Fig. 3). Latitude and a 2nd order polynomial term explained 10.5% and 12.9% of the variation in crop group richness in 1961 and 2017, respectively (model *p*≤0.001 in both cases), with the richness-latitude relationship being similar in both years (Table 1). Specifically, regression models indicated that crop group richness increased from equatorial regions towards mid-latitude countries, with modeled peak crop group richness occurring at~37° latitude in both 1961 and 2017; richness then declined at higher latitudes, denoted by statistically signifcant (*p*≤0.002) negative 2nd order polynomial terms (Fig. 3). While these trends were similar between years, this analysis did refect the increased crop group richness that occurred between 1961 and 2017, in 152 of 164 of the countries included in this analysis.

Changes in crop evenness and composition across countries through time. Piecewise models evaluating changes in crop group evenness (*J′*) through time converged for 185 countries; in these countries year explained on average 80.9% of the variation in *J'* (model $p < 0.01$ in all cases, r^2 range = 0.198–0.992; Table S2). In these countries changes in *J′* began on average in 1981 (±11.9 years m.a.d.), although compared to changes in group richness, periods of change in evenness were more prolonged lasting on average for 15 years (±13.3 years m.a.d.) (Fig. 4A–D). Unlike analyses of crop richness, changes in evenness were less systematic, such that through the period of change evenness declined in 97 countries, increased in 88 countries, and average

Indicator 1: Onset of change in crop commodity group richness

Indicator 2: Duration of change in crop commodity group richness

Figure 2. Maps and histograms of three indicators of crop commodity group richness (*S*) change across 165 countries. Values for all three indicators for each country were derived from piecewise linear models predicting *S* as a function of year (see Fig. 1 for example). Countries coloured gray in the maps were those where either data was not available or the piecewise models failed to converge (denoted in Table S1). Histograms and associated descriptive statistics for each indicator are also presented, with means $(\pm s.d.)$ or medians $(\pm m.a.d.)$ denoted visually by the points and error bars below the histograms. All piecewise model parameters for each country are presented in Table S1. Maps were generated using the mapCountryData function in the rworldmap R package³⁰.

changes in *J′* centred on zero (mean=− 0.004 year−1±0.04 s.d.; Fig. 4E,F). However, similar to patterns of change in richness, Indicators 1–3 associated with changes in evenness were also independent of total area of a country

4

Table 1. Parameters and diagnostics for linear regression models (including a 2nd-order polynomial term) predicting crop group richness as a function of latitude in 1961 and 2017. Only the 164 countries with data from both years were included in these analyses. Parameter estimate standard errors shown in parentheses. Also shown are the latitudes at which crop group richness peaked, according to these models, which are shown visually in Fig. 3.

(linear regression r^2 =0.001–0.004, $p \ge 0.404$), or the area of a country under crop cultivation (linear regression r^2 =0.0–0.013, $p \ge 0.12$).

Multivariate analysis detected a signifcant infuence of country, year, and a year-by-country interaction term on crop composition (Adonis *p* < 0.01 in all cases; Table 2). Of these variables, country diferences were most pronounced with country identity explaining 89.5% of the variability in crop composition. Year explained \sim 1% of the variability in crop composition, while a country-by-year interaction term explained an additional 6.3% of the variation in crop composition (Table 2). Based on the non-metric multidimensional scaling (NMDS) analysis, there was trend of increasing similarity in crop composition among countries over time. This is illustrated by increasingly smaller 95% confdence bands surrounding the data points along NMDS axes 1 and 2 from 1961 through to 1983 (i.e., approximately the average year in which changes in crop group richness and evenness commence); multivariate space encapsulated by the 95% confdence band is then further reduced through 2017 (Figure S1).

Indicator 1: Onset of change in crop commodity group evenness

Indicator 2: Duration of change in crop commodity group evenness

Figure 4. Maps and histograms of three indicators of crop commodity group evenness (Pielou's evenness index (*J*^{*′*})) across 185 countries. Values for all three indicators for each country were derived from piecewise linear models predicting *J'* as a function of year, where harvested area (in ha) was used to approximate group abundance. Countries coloured gray in the maps were those where either data was not available or the piecewise models failed to converge (denoted in Table S2). Histograms and associated descriptive statistics for each indicator are also presented, with means $(\pm s.d.)$ or medians $(\pm m.a.d.)$ denoted visually by the points and error bars below the histograms. All piecewise model parameters for each country are presented in Table S2. Maps were generated using the mapCountryData function in the rworldmap R package³⁰.

Socioeconomics correlates of changes in crop group diversity and evenness. Patterns of change in crop richness or evenness were not systematically related to country socio-economic status, with HDI values predicting only 1.3% of Indicators 1–3 for both *S* and *J'* on average (Table 3). The only exception to this is that

Table 2. Results from an Adonis test evaluating changes in crop commodity group composition at national scales from 1961 to 2017. The distance matrix employed in this analysis was based on non-metric multidimensional scaling, whereby cultivation area was used as an estimate for crop group abundance. Results are presented visually in Figure S1.

Table 3. Results and variance components predicting Indicators of crop group diversity (*S*) and evenness (*J′*) change, as a function of Human Development Index (HDI), continent identity, and region identity. In these models an intercept and slope related to HDI were included as fxed efects, while region within continent were included as nested random effects. Statistically significant (p <0.05) fixed effect model parameters are highlighted in bold, and data transformations (based on distribution ftting) precede the Indicator numbers (as per Figs. 2 and 4).

the rate of change in *S* was signifcantly negatively related to HDI: countries with lower HDI scores expressed greater increases in *S* (i.e., Indicator 3 values) during their period of crop diversity change vs. countries with higher HDI (Table 3). This analysis did point to stronger explanatory power of spatial location in determining patterns of change in crop *S* and *J′*: both continent and region identity explained an average of 9.5% and 2.0% of the variation in Indicators 1–3, respectively (Table 3). However, ultimately country-by-country variation in crop group diversity change was largely idiosyncratic, with 87.0% of the variation in Indicators 1–3 for both *S* and *J′* being unaccounted for by socio-economic and spatial factors included in our models.

Discussion

The vast majority of the world's countries have experienced significant increases in the number of crop groups being cultivated over recent decades; changes that contribute to a detectable increase in the similarity of crops being grown among countries, that has occurred since the 1960s. Our fndings align with those from previous studies⁸ that detected similar changes in crop taxonomic and phylogenetic diversity at supra-national regional, continental, and global scales: (1) a period of little change through the 1960s and 1970s, followed by (2) the onset of increases in crop commodity group richness commencing the 1980s and extending through the end of the 1990s, followed by (3) a levelling of of commodity group richness beyond the 2000s. While the concordance of these two studies should not be surprising, our fndings contribute a new and more nuanced understanding of the remarkably similar patterns in crop group richness that have occurred across virtually all nations in recent decades; the consistency of which supports the idea that changes in crop diversity and biogeography occurring since the Columbian Exchange, are a ubiquitous feature of the Anthropocene.

However, these near-universal patterns in crop richness increases over recent decades (observed in all but 13 countries evaluated here) have had unequal and more variable efects on the evenness of crops being cultivated. The average duration of change in evenness (15 years) was nearly twice as long vs. the average duration of changes in richness (9 years). Previous work has noted that the majority of introduced crops into national agricultural production portfolios, largely owes to cultivation of crops beyond their country or region of origin¹³. So while introduction of novel crop groups has been rather succinct—corresponding to periods of rapid change in structural adjustment through the 1980s and 1990s^{21,22}—expansion of these crops across more cultivated lands draws out over longer periods. Tis discrepancy likely refects the lag between new crop introductions, compared to longer-term process associated with agricultural economic adjustments for these crops including expansion of export markets or crop-specific subsidies¹⁵, and to a much lesser extent, expansion of domestic consumption markets for new crops 11 .

The specific model parameters for country-specific trends may be sensitive to data quality¹⁷. However, here we observed a clear signal of both decreasing and increasing trends in crop evenness across 52% and 48% of the

7

world's countries, respectively. Tis approximately even proportion of increase and decrease in evenness detected among all countries here, would explain why analyses of global scale crop production trends reported no change in evenness between 1961 and 2013¹¹. Yet similar to the growing number of analyses of production and consumption at multiple scales^{8,11,14,15}, our multivariate analysis does indicate that over recent decades, countries have expressed a statistically significant increase in the similarity of crops in their agricultural lands. The implications of these shifs remain speculative, and based on our analysis, are clearly scale-dependent.

Specifcally, increases in the similarity in crop composition across regions and continents could indicate growing susceptibility of agriculture to pest and pathogen outbreaks, and perhaps climate change efects like regional temperature increases or precipitation declines⁸. Alternatively, a wider geographic spread of crop groups could well represent a means of bufering production from localized disruptions including local climatic change or weather events, pest or pathogens, or civil unrest¹¹. While both are plausible, generalizing either hypothesis to predict the impacts of homogenization across all agricultural lands globally is inconsistent with our countryspecifc analyses here. Since 1961, the evenness of crops in agricultural lands is both increasing and decreasing in approximately the same proportion of countries and agricultural area globally. Therefore, while previous studies including our own have speculated on how changes in crop diversity will infuence global agricultural production and sustainability^{8,11}, our results suggest the largest spatial scale at which potential impacts of stressors on crop production—though not food consumption or food security per se—can be robustly predicted is on a per-country basis.

Knowledge of the high context-dependency of agricultural adaptation, management, and crop selection likely indicates even smaller spatial scales (i.e., communities, households) are needed in order to fully predict susceptibility of production in the future. Indeed, an important caveat for our analysis is that even national scales likely do not comprehensively indicate how agricultural functional diversity has changed in the past 60 years. Our work here focuses on large-scale industrial agriculture, with the contributions of small-scale agriculture to overall crop diversity, particularly in terms of locally adapted varieties or landraces, likely underestimated³¹. Spatially explicit records of crop genetic diversity including locally adapted or cultivated crop phenotypes and their wild relatives are becoming more widely available, yet such a comprehensive assessment of crop genetic diversity remains prohibitive; indeed, while instructive, even analyses of phylogenetic crop diversity^{8,32,33} are limited to the crop species or sub-species levels. Clearly more work is needed to better integrate the methodological frameworks and concepts used here and in related studies^{11,14}, with finer-scale datasets on crop genetic diversity.

Determination of potential driving factors behind diversification trends. The factors underpinning national-scale increases in crop richness or evenness include a nuanced mix of environmental, socio-economic, and cultural factors. Here we hypothesized that patterns of change in crop group richness would correlate with development status (quantifed as the Human Development Index). However, HDI was clearly a poor predictor of the rate, duration, and timing of changes in crop group richness and evenness. While national-scale agriculture portfolios did not systematically change as a function of HDI, spatial location (i.e., region and continent) did explain~10–12% of the variation in patterns of change. From a strictly socio-economics perspective, this fnding could point to regional-scale agricultural policy in driving similar changes in crop group richness, on a region-by-region basis^{21,25}. Alternatively, it could also reflect regional-scale similarities in non-governmental organization (NGO) interventions in the agricultural sector²⁶. Indeed, previous studies have indicated that governmental and NGO intervention has led to a \sim 17% increase in global crop diversity on average²⁶. Additionally, and perhaps surprisingly, this same study²⁶ found that while climate is an important driver of on-farm crop diversity change, its infuence is secondary compared to market conditions.

Agriculture and the reshufing of species through the Anthropocene. We detected a statistically signifcant and hump-shaped latitudinal gradient in crop group richness, with low- to mid-latitude regions (centred on \sim 37° latitude), expressing the largest number of cultivated crop groups in large-scale agricultural lands. Consistent with previous studies, this non-linear latitudinal trend most likely emerges due to: (1) countries at these latitudes having a range of climatic conditions (i.e., Köppen climate zones) that supports cultivation of a large diversity of crop functional types and year-round production; and (2) countries at these latitudes encapsulating many of the world's centres of crop domestication, and having received among the largest imports of crops during the Columbian Exchange^{13,29,33}

It is expected that the band of latitude supporting the highest number of crop groups could shif as a result of global environmental change drivers. Indeed, researchers have now long projected that crop diversity and richness within countries will change as climate change intensifes, with clusters of high crop diversity moving poleward through time34,35. However, similar to categorical comparisons of crop richness produced in tropical vs. temperature countries¹⁵, our analysis did not detect evidence of a disruption in the latitudinal trends of crop species richness in between 1961 and 2017. Instead, we fnd only a systematic increase in richness across all latitudes, indicating that the cultivation of new crop commodity groups in large-scale agricultural lands has not fundamentally altered the latitudes at which crop group richness is highest. Moreover, our analysis here (1) likely misses the role that high crop diversity of small holder (i.e. $<$ 2-ha in size) farms³⁶—which are disproportionately concentrated at lower latitudes³⁷—play in driving latitudinal crop diversity gradients; and (2) does not address functional- or phylogenetic crop diversity, which may show diferent latitudinal patterns. Including these factors in additional analyses is a key step for further resolving our understanding of how crop biogeography is changing during the Anthropocene.

Methods

Data acquisition. Our analysis was based on open access crop production data from the United Nation's Food and Agricultural Organization (FAO) spanning from 1961 to 201718. We extracted data on area harvested (in ha) for 339 FAO-defned crop groups being grown in all UN-recognized countries. Since our research centred on understanding, quantifying, and mapping changes in crop diversity in current agricultural lands, countries that cease to exist (e.g., Yugoslavia) were not included in our analysis, resulting in data for 201 countries (Table S1). Prior to analyses, we adjusted certain crop group listings following our previous analyses of global changes in crop diversity⁸. Specifically, "Cottonlint" and "Cottonseed" were duplicated in our dataset and were therefore compiled as "Seedcotton", while "Palmkernels" were renamed as "Oilpalmfruit." Additionally, "Fruitpomenes", "Fruitstonenes", and "Grainmixed" were removed from analysis since these crop groupings are not associated with any specific crop species in the FAO database¹⁸. Finally, "Mushroomsandtruffles" were removed since it relates to non-plant species, and "Coir" was removed because it is a plant by-product.

Changes in crop richness over time. All statistical analyses were performed using R version 3.3.3 statistical software (R Foundation for Statistical Computing, Vienna, Austria). The initial step in our analysis was to calculate both crop richness and evenness for each country, at each individual year, using the vegan R package³⁸. Based on these datasets, we then used the analytical framework developed by⁸ to evaluate how crop species richness and evenness have changed in each individual country across its entire data range.

Specifically, in their analysis Martin et al.⁸ found that piecewise linear regression models provided the strongest descriptions of crop species richness change over time, across 21 of 22 FAO-defned regions globally. We therefore followed this approach by ftting a piecewise linear regression model for each country individually, that predicts changes in species richness over time. Piecewise model ftting was a two-step process, whereby for each country we frst ft a linear regression model of the form:

$$
S = a + (b \times \text{year}) \tag{1}
$$

where *a* is the intercept and *b* represents the rate of change in crop group richness (S) through time. This linear model (Eq. 1) was then used as the basis of a piecewise linear regression model, which was ftted in order to estimate breakpoints in the relationship between *S* and year. Specifcally, piecewise models were ft using the segmented function in the segmented R package³⁹, and were of the form:

$$
S = a + b(\text{year}) + ((c(\text{year} - \psi_1) \times I(\text{year} > \psi_1)) + (d(\text{year} - \psi_2) \times I(\text{year} > \psi_2))
$$
 (2)

where *a* is as in Eq. (1), and *b* represents the slope of the *S*-year relationship prior to the frst breakpoint (ψ1). Here, *c* represents the difference in the slope of the *S*-year relationship between the first and second piecewise model segments; the *c* parameter therefore applies only when the frst conditional indicator function (denoted by "*I*") is true. Similarly, *d* represents the diference in slopes for the *S*-year relationship between the frst, second, and third segments, which only applies when the second conditional indicator function is true. In sum, the slope of the relationship between *S* and year is equal to *b* prior to the ψ 1, is equal to *b* + *c* between ψ 1 and ψ 2, and is equal to $b+c+d$ after ψ 2. Piecewise models were fit with initial starting parameters of 1975 and 2000 for ψ 1 and ψ 2, respectively. The ψ 1 and ψ 2 parameters were tuned manually for 29 countries with a shortened data range, following visual inspection of data (see Tables S1 and S2).

Based on this piecewise regression model procedure, we then used parameters from Eq. (2) to determine three key indicator points of crop diversity change through time for each country (displayed visually in Fig. 1). Indicator 1 refects the onset of diversifcation in each country, and was calculated as Breakpoint 1 (ψ1) in Eq. (2); this indicator therefore corresponds to the year in which notable changes in species richness began. Indicator 2 refects the duration of the crop diversifcation period in each country, and was calculated as the diference between breakpoints 2 and 1 (i.e., ψ2-ψ1 from Eq. 2); this indicator therefore represents the duration of the period when crop prominent changes in crop diversity occurred. Finally, Indicator 3 refects the rate at which crop diversity changed throughout the diversifcation period in each country; this indicator was calculated as the rate of crop diversity change (between ψ 1 and ψ 2), which in our models corresponded to the sum of the slopes (1) prior to the first breakpoint, and (2) between the first and second breakpoints (i.e., corresponding to $b + c$ in Eq. 2). For each indicator we then calculated summary statistics as either mean \pm standard deviations or median \pm median absolute deviations (m.a.d.), where data was normally or log-normally distributed, respectively. Country values for each indicator were mapped using the mapCountryData function in the rworldmap R package⁴⁰.

Changes in crop evenness over time. Evaluations of temporal changes in crop evenness at national scales followed this same analytical approach as above. First, for each country-by-year combination we calculated Pielou's evenness index (*J′*)—which ranges from 0 to 1, with values closer to 0 indicating less evenness or greater abundance of a few dominant crop groups, and values closer to 1 representing more equitable abundances of crop groups—as:

$$
J' = \frac{H'}{\ln(S)}\tag{3}
$$

where *S* is again crop richness, and *H′* is the Shannon–Weiner diversity index calculated as:

$$
H' = -\sum_{i=1}^{S} p_i l n p_i \tag{4}
$$

where p_i represents the relative proportion of the *i*th crop group for a given country-by-year combination. In these evenness calculations, all values of p_i were estimated as the relative proportion of agricultural area (measured in ha) occupied by a given crop commodity group, within a country at a given year; this analytical approach was employed by Martin et al.⁸ when assessing crop group composition at supra-national scales. We then evaluated how *J′* values changed in each country through time by replicating our stepwise modelling analyses above, substituting *J′* for *S* in Eqs. (1) and (2), and extracting the same model indicators (Fig. 1). Finally, we calculated summary statistics and mapped each of these indicators, as described above.

Changes in crop composition across countries and over time. We used multivariate analyses to evaluate how temporal changes in *S* and *J′* infuenced crop composition across countries and over time. To do so, we created a community composition matrix whereby national-level crop assemblages were estimated for each of the country-by-year combinations. In this matrix, area harvested was taken as an approximation of the abundance of each crop group within each country-by-year combination (again following Martin et al.⁸). Since these abundances (or area harvested) across country-by-year combinations varied over orders of magnitude, we used non-metric multidimensional scaling (NMDS) to analyze and visualize spatial (country) and temporal (year) differences in crop diversity. Specifically, we used the vegan R package³⁸ to calculate all 58,899,231 Bray– Curtis dissimilarities among all 10,854 data points (i.e., crop group composition in every country-by-year data point), as:

$$
BC_{jk} = \frac{\sum i |x_{ij} - x_{ik}|}{\sum i (x_{ij} + x_{ik})}
$$
\n⁽⁵⁾

where BC_{ik} represents the dissimilarity between the *j*th and *k*th community, x_{ii} represents the abundance (i.e., area harvested) of crop group *i* in sample *j*, and *xik* represents the abundance of crop group *i* in sample *k*. We then used a multivariate analysis of variance (i.e., an Adonis test), to test for signifcant diferences in Bray–Curtis distances as a function of country, year, and a country-by-year interaction. Signifcance was assessed using a permutation test, with 99 permutations used.

Latitudinal gradients in crop richness. To test our hypotheses surrounding the presence of, and temporal changes in, latitudinal gradients in crop group diversity, we focused on 164 countries for which crop group diversity was available in both 1961 and 2017. For each of these two datasets, we ft a separate linear regression model that predicts crop group richness as a function of latitude (expressed as an absolute value) and a 2ndorder polynomial term for the 'latitude²' variable. From both of these models, we extracted and compared latitude value at which crop group richness was estimated/ modelled to peak.

Predictors of change in crop diversity and composition. We tested if Human Development Index (HDI) was correlated with patterns of change in crop diversity and composition. Briefy, the HDI is a composite index of four metrics related to socio-economic status, including life expectancy at birth, expected years of schooling for children at a school-centring age, mean years of schooling for adults≥25 years of age, and logtransformed gross national income per capita. These values are then aggregated on a per country basis, into an HDI index that ranges from 0–1 with higher scores denoting higher performance in these indicators. We employed 2017 HDI values in our analysis here, in order to include the most countries possible in each analysis (since earlier HDI scores are less readily available) 41 .

We then used linear mixed efects models to test if patterns of change in crop diversity and evenness varied systematically with HDI values. Tis entailed ftting six linear mixed models, where each of our six indicators (i.e., Indicators 1–3 for both *S* and *J′*) were predicted as a function of HDI; these models also accounted for potential spatial autocorrelation in Indicator values by including the FAO-defned continent identity and FAOdefned region identity of each country, as a nested random variable. Models were ft using the lme function in the nlme R package⁴¹. We then estimated the proportion of variation in each indicator that is explained by HDI, continent identity, and region identity, using the varcomp function in the ape R package⁴²—which partitioned explained variation across continents and regions—as well as the sem.model.fts function in the piecewiseSEM R package43—which partitioned explained variation across the fxed (i.e., model intercept and HDI) vs. random (i.e., continent and region) efects. Due to diferences in HDI data availability and in the number of piecewise models that converged, $n=152$ countries for all models of *S* indicators and $n=139$ countries for all models of *J'* indicators. Log-transformed values of Indicators were used in these analyses where they better approximated a log-normal distribution, as determined using the fitdistrplus function in the fitdistrplus R package⁴⁴.

Received: 1 April 2021; Accepted: 29 September 2021

References

1. Crutzen, P. J. *Earth System Science in the Anthropocene* 13–18 (Springer, 2006).

2. Lewis, S. L. & Maslin, M. A. Defning the Anthropocene. *Nature* **519**, 171–180 (2015).

- 3. Malhi, Y. The concept of the Anthropocene. Annu. Rev. Environ. Resour. 42, 77-104 (2017).
- 4. Young, K. R. Biogeography of the Anthropocene: Novel species assemblages. *Prog. Phys. Geogr.* **38**, 664–673 (2014).
	- 5. Capinha, C., Essl, F., Seebens, H., Moser, D. & Pereira, H. M. Te dispersal of alien species redefnes biogeography in the Anthropocene. *Science* **348**, 1248–1251 (2015).
	- 6. Parmesan, C. & Yohe, G. A globally coherent fngerprint of climate change impacts across natural systems. *Nature* **421**, 37–42 (2003).
	- 7. Zimmerer, K. S. et al. The biodiversity of food and agriculture (Agrobiodiversity) in the anthropocene: Research advances and conceptual framework. *Anthropocene* **25**, 100192 (2019).
	- 8. Martin, A. R. *et al.* Regional and global shifs in crop diversity through the Anthropocene. *PLoS ONE* **14**, e0209788 (2019).
	- 9. Crosby, A. W. *Te Columbian Exchange: Biological and Cultural Consequences of 1492* Vol. 2 (Greenwood Publishing Group, 2003).
	- 10. Mann, C. C. *1493: How the ecological collision of Europe and the Americas gave rise to the modern world* (2011).
	- 11. Aguiar, S., Texeira, M., Garibaldi, L. A. & Jobbágy, E. G. Global changes in crop diversity: Trade rather than production enriches supply. *Glob. Food Secur.* **26**, 100385 (2020).
	- 12. Campi, M., Dueñas, M. & Fagiolo, G. Specialization in food production afects global food security and food systems sustainability. *World Dev.* **141**, 105411 (2021).
	- 13. Khoury, C. K. *et al.* Origins of food crops connect countries worldwide. *Proc. R. Soc. B Biol. Sci.* **283**, 20160792 (2016).
	- 14. Khoury, C. K. *et al.* Increasing homogeneity in global food supplies and the implications for food security. *Proc. Natl. Acad. Sci. U.S.A.* **111**, 4001–4006 (2014).
	- 15. Nelson, E. J. *et al.* Commercial plant production and consumption still follow the latitudinal gradient in species diversity despite economic globalization. *PLoS ONE* **11**, e0163002 (2016).
	- 16. Renard, D. & Tilman, D. National food production stabilized by crop diversity. *Nature* **571**, 257–260 (2019).
	- 17. Mahaut, L., Violle, C. & Renard, D. Complementary mechanisms stabilize national food production. *Sci. Rep.* **11**, 1–7 (2021).
	- 18. Nations, F. A. O. U. <http://www.fao.org/faostat/en/>(2020).
	- 19. Hessenauer, P. *et al.* Evolution and adaptation of forest and crop pathogens in the Anthropocene. *Phytopathology* **111**, 49–67 (2021).
	- 20. Aguilar, J. *et al.* Crop species diversity changes in the United States: 1978–2012. *PLoS ONE* **10**, e0136580 (2015).
	- 21. Isakson, S. R. Maize diversity and the political economy of agrarian restructuring in Guatemala. *J. Agrar. Chang.* **14**, 347–379 (2014)
	- 22. Bonneuil, C. Development as experiment: Science and state building in late colonial and postcolonial Africa, 1930–1970. *Osiris* 15, 251-281 (2000)
	- 23. Holt-Gimenez, E., Altieri, M. A. & Rosset, P. Food First Policy Brief No. 12: Ten Reasons Why the Rockefeller and the Bill and Melinda Gates Foundations' Alliance for Another Green Revolution Will Not Solve the Problems of Poverty and Hunger in Sub-Saharan Africa. (Institute of Food and Development Policy, Oakland, USA, 2006).
	- 24. Arnold, D. Europe, technology, and colonialism in the 20th century. *Hist. Technol.* **21**, 85–106 (2005).
	- 25. Meertens, B. Agricultural performance in Tanzania under structural adjustment programs: Is it really so positive?. *Agric. Hum. Values* **17**, 333–346 (2000).
	- 26. Chen, M. *et al.* Diversifcation and intensifcation of agricultural adaptation from global to local scales. *PLoS ONE* **13**, e0196392 (2018).
	- 27. Pontarp, M. *et al.* Te latitudinal diversity gradient: Novel understanding through mechanistic eco-evolutionary models. *Trends Ecol. Evol.* **34**, 211–223 (2019).
	- 28. Harlan, J. R. Agricultural origins: Centers and noncenters. *Science* **174**, 468–474 (1971).
	- 29. Meyer, R. S., DuVal, A. E. & Jensen, H. R. Patterns and processes in crop domestication: An historical review and quantitative analysis of 203 global food crops. *New Phytol.* **196**, 29–48 (2012).
	- 30. South, A. rworldmap: A new R package for mapping global data. *R J.* **3**, 35–43 (2011).
	- 31. Ricciardi, V., Mehrabi, Z., Wittman, H., James, D. & Ramankutty, N. Higher yields and more biodiversity on smaller farms. *Nat. Sustain.* 1–7, (2021).
	- 32. Milla, R. Crop origins and phylo food: A database and a phylogenetic tree to stimulate comparative analyses on the origins of food crops. *Glob. Ecol. Biogeogr.* **29**, 606–614 (2020).
	- 33. Milla, R. *et al.* Phylogenetic patterns and phenotypic profles of the species of plants and mammals farmed for food. *Nat. Ecol. Evol.* **2**, 1808–1817 (2018).
	- 34. Leemans, R. & Solomon, A. M. Modeling the potential change in yield and distribution of the earth's crops under a warmed climate. *Clim. Res.* 79–96 (1993).
	- 35. Zabel, F., Putzenlechner, B. & Mauser, W. Global agricultural land resources–a high resolution suitability evaluation and its perspectives until 2100 under climate change conditions. *PLoS ONE* **9**, e107522 (2014).
	- 36. Jarvis, D. I. *et al.* A global perspective of the richness and evenness of traditional crop-variety diversity maintained by farming communities. *Proc. Natl. Acad. Sci. U.S.A.* **105**, 5326–5331 (2008).
	- 37. Lowder, S. K., Skoet, J. & Raney, T. Te number, size, and distribution of farms, smallholder farms, and family farms worldwide. *World Dev.* **87**, 16–29 (2016).
	- 38. Oksanen, J. *et al.* vegan: Community Ecology Package. R package version 2.4–5. (2017).
	- 39. Vito, M. & Muggeo, R. Segmented: An R package to ft regression models with broken-line relationships. cran.r-project.org/doc/ Rnews/. *R News* **8**, 20–25 (2008).
	- 40. South, A. rworldmap: A new R PACKAGE FOR MAPPING GLOBAL DATa. *Te R Journal* **3**, 35–43 (2011).
	- 41. nlme: Linear and Nonlinear Mixed Efects Models. R package version 3.1–131 [\(https://CRAN.R-project.org/package=nlme](https://CRAN.R-project.org/package=nlme), 2017).
	- 42. Paradis, E., Claude, J. & Strimmer, K. APE: Analyses of phylogenetics and evolution in R language. *Bioinformatics* **20**, 289–290 (2004)
	- 43. Lefcheck, J. S. piecewiseSEM: Piecewise structural equation modelling in R for ecology, evolution, and systematics. *Methods Ecol. Evol.* **7**, 573–579 (2016).
	- 44. Delignette-Muller, M. L. & Dutang, C. ftdistrplus: An R package for ftting distributions. *J. Stat. Sofw.* **64**, 1–34 (2015).

Acknowledgements

Tis research was undertaken in part based on funding from Natural Sciences and Engineering Research Council of Canada (NSERC) Discovery Grants to Adam Martin, Marney Isaac, and Marc Cadotte. MEI is also supported by supported by NSERC's Canada Research Chairs program, while MWC is supported by the TD Professor of Urban Forest Conservation and Biology Chair program, the Canada Foundation for Innovation, and the Ontario Research Fund. Denis Vile is supported by the European Union and the Region Languedoc-Roussillon "Chercheur d'Avenir" (FEDER FSE IEJ 2014–2020; Grant Project "APSEVIR"). Cyrille Violle was supported by the European Research Council (ERC) Starting Grant Project StG-2014-639706-CONSTRAINTS. Funding from the University of Toronto Scarborough's (UTSC) Vice Principal Research Office in support of UTSC's Centre for Agroecosystems Research also supported this research.

Author contributions

R.O.M. lead data consolidation, analysis, and visualization, and wrote the manuscript. A.R.M. conceived the study, contributed to data analysis and visualization, and co-wrote the manuscript. M.W.C, M.E.I., R.M., D.V., and C.V. contributed to manuscript editing and data analysis.

Competing interests

The authors declare no competing interests.

Additional information

Supplementary Information The online version contains supplementary material available at [https://doi.org/](https://doi.org/10.1038/s41598-021-99728-5) [10.1038/s41598-021-99728-5](https://doi.org/10.1038/s41598-021-99728-5).

Correspondence and requests for materials should be addressed to A.R.M.

Reprints and permissions information is available at [www.nature.com/reprints.](www.nature.com/reprints)

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International \odot $\left[{\rm (cc)}\right]$ License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit<http://creativecommons.org/licenses/by/4.0/>.

 $© The Author(s) 2021$