

The GGCMI Phase II experiment: global crop yield responses to changes in carbon dioxide, temperature, water, and nitrogen levels

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Abstract

Concerns about food security under climate change have motivated efforts to better understand the future changes in yields by using detailed process-based models in agronomic sciences. Process-based models differ on many details affecting yields and considerable uncertainty remains in future yield projections. Phase II of the Global Gridded Crop Model Intercomparison (GGCMI), an activity of the Agricultural Model Intercomparison and Improvement Project (AgMIP), consists of a large simulation set with perturbations in atmospheric CO₂ concentrations, temperature, precipitation, and applied nitrogen inputs and constitutes a data-rich basis of projected yield changes across twelve models and five crops (maize, soy, rice, spring wheat, and winter wheat) using global gridded simulations. In this paper we present the simulation output database from Phase II of the GGCMI effort, a targeted experiment aimed at understanding the sensitivity to and interaction between multiple climate variables (as well as management) on yields, and illustrate some initial summary results from the model intercomparison project. We also present the construction of a simple “emulator or statistical representation of the simulated 30-year mean climatological output in each location for each crop and model. The emulator captures the response of the process-based models in a lightweight, computationally tractable form that facilitates model comparison as well as potential applications in subsequent modeling efforts such as integrated assessment.

Keywords: climate change, food security, model emulation, AgMIP, crop model

1. Introduction

2 Projecting crop yield response to a changing climate is of
3 great importance, especially as the global food production sys-
4 tem will face pressure from increased demand over the next
5 century. Climate-related reductions in supply could therefore
6 have severe socioeconomic consequences. Multiple studies
7 with different crop or climate models predict sharp reduction in
8 yields on currently cultivated cropland under business-as-usual
9 climate scenarios, although their yield projections show consid-
10 erable spread (e.g. Porter et al. (IPCC), 2014, Rosenzweig et al.,
11 2014, Schauberger et al., 2017, and references therein). Model
12 differences are unsurprising because crop responses in models
13 can be complex, with crop growth a function of complex inter-
14 actions between climate inputs and management practices.

15 Computational Models have been used to project crop yields
16 since the 1950's, beginning with statistical models (Heady,
17 1957, Heady & Dillon, 1961) that attempt to capture the rela-
18 tionship between input factors and resultant yields. These sta-
19 tistical models were typically developed on a small scale for lo-
20 cations with extensive histories of yield data. The emergence of
21 computers allowed development of numerical models that sim-
22 ulate the process of photosynthesis and the biology and phe-
23 nology of individual crops (first proposed by de Wit (1957),
24 Duncan et al. (1967) and attempted by Duncan (1972)). For a
25 history of crop model development see the appendix of Rosen-
26 zweig et al. (2014). A half-century of improvement in both
27 models and computing resources means that researchers can
28 now run crop simulation models for many years at high spatial
29 resolution on the global scale.

30 Both types of models continue to be used, and compara-
31 tive studies have concluded that when done carefully, both ap-
32 proaches can provide similar yield estimates (e.g. Lobell &

33 Burke, 2010, Moore et al., 2017, Roberts et al., 2017, Zhao
34 et al., 2017). Models tend to agree broadly in major response
35 patterns, including a reasonable representation of the spatial
36 pattern in historical yields of major crops (e.g. Elliott et al.,
37 2015, Müller et al., 2017) and projections of decreases in yield
38 under future climate scenarios.

Process models do continue to struggle with some important details, including reproducing historical year-to-year variability (e.g. Müller et al., 2017), reproducing historical yields when driven by reanalysis weather (e.g. Glotter et al., 2014), and low sensitivity to extreme events (e.g. Glotter et al., 2015). These issues are driven in part by the diversity of new cultivars and genetic variants, which outstrips the ability of academic modeling groups to capture them (e.g. Jones et al., 2017). Models do not simulate many additional factors affecting production, including pests/diseases/weeds. For these reasons, individual studies must generally re-calibrate models to ensure that short-term predictions reflect current cultivar mixes, and long-term projections retain considerable uncertainty (Wolf & Oijen, 2002, Jagtap & Jones, 2002, Iizumi et al., 2010, Angulo et al., 2013, Asseng et al., 2013, 2015). Inter-model discrepancies can also be high in areas not yet cultivated (e.g. Challinor et al., 2014, White et al., 2011). Finally, process-based models present additional difficulties for high-resolution global studies because of their complexity and computational requirements. For economic impacts assessments, it is often impossible to integrate a set of process-based crop models directly into an integrated assessment model to estimate the potential cost of climate change to the agricultural sector.

Nevertheless, process-based models are necessary for understanding the global future yield impacts of climate change for many reasons. First, cultivation may shift to new areas, where no yield data are currently available and therefore statistical models cannot apply. Yield data are also often limited in the de-

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67 developing world, where future climate impacts may be the most
 68 critical. Second, only process-based models can capture the
 69 growth response to elevated CO₂, novel conditions that are not
 70 represented in historical data (e.g. Pugh et al., 2016, Roberts
 71 et al., 2017). Similarly process-based models can represent
 72 novel changes in management practices (e.g. fertilizer input)
 73 that may ameliorate climate-induced damages.

74 Statistical emulation of crop simulations has been used to
 75 combine advantageous features of both statistical and process-
 76 based models. The statistical representation of complicated nu-
 77 matical simulation (e.g. O'Hagan, 2006, Conti et al., 2009), in
 78 which simulation output acts as the training data for a statisti-
 79 cal model, has been of increasing interest with the growth of
 80 simulation complexity and volume of output. Such emulators
 81 or "surrogate models" have been used in a variety of fields in-
 82 cluding hydrology (e.g. Razavi et al., 2012), engineering (e.g.
 83 Storlie et al., 2009), environmental sciences (e.g. Ratto et al.,
 84 2012), and climate (e.g. Castruccio et al., 2014, Holden et al.,
 85 2014). For agricultural impacts studies, emulation of process-
 86 based models allows exploring crop yields in regions outside
 87 ranges of current cultivation and with input variables outside
 88 historical precedents, in a lightweight, flexible form that is com-
 89 patible with economic studies.

90 In the past decade, many studies have developed emulators of
 91 crop yields from process-based models. Early studies propos-
 92 ing or describing potential emulators include Howden & Crimp
 93 (2005), Räisänen & Ruokolainen (2006) and Lobell & Burke
 94 (2010). In an early application, Ferrise et al. (2011) used a Arti-
 95 ficial Neural Net trained on simulation outputs to predict wheat₁₀₁
 96 yields in the Mediterranean. Studies developing single-model₁₀₂
 97 emulators include Holzkämper et al. (2012) for the CropSyst₁₀₃
 98 model, Ruane et al. (2013) for the CERES wheat model, Oye-₁₀₄
 99 bamiji et al. (2015) for the LPJmL model (for multiple crops,₁₀₅
 100 using multiple scenarios as a training set). In recent years, emu-₁₀₆

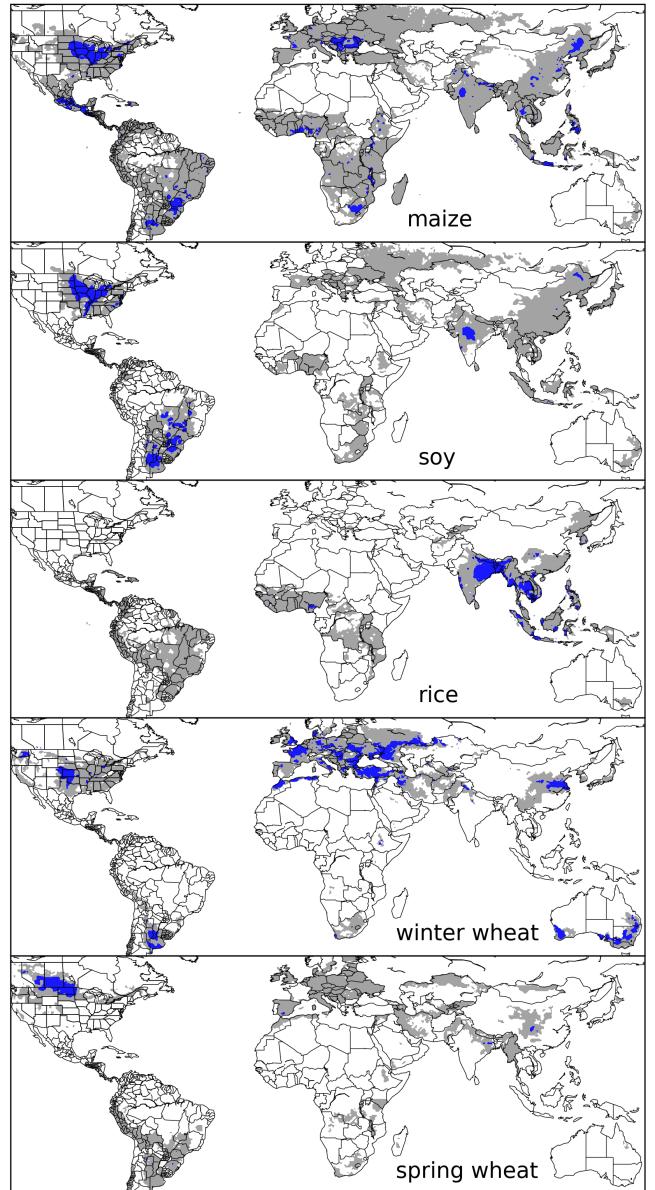


Figure 1: Presently cultivated area for rain-fed crops in the real world. Blue contour area indicates grid cells with more than 20,000 hectares (approx. 10% of the land in a grid cell near the equator for example) of crop cultivated. Gray contour shows area with more than 10 hectares cultivated. Cultivated areas for maize, rice, and soy are taken from the MIRCA2000 ("monthly irrigated and rain-fed crop areas around the year 2000") dataset (Portmann et al., 2010). Areas for winter and spring wheat areas are adapted from MIRCA2000 data and sorted by growing season. For analogous figure of irrigated crops, see Figure S1.

lators have begun to be used in the context of multi-model inter-
 comparisons, with Blanc & Sultan (2015), Blanc (2017), Ost-
 berg et al. (2018) and Mistry et al. (2017) using them to analyze
 the five crop models of the Inter-Sectoral Impacts Model Inter-
 comparison Project (ISIMIP) (Warszawski et al., 2014) (for
 maize, soy, wheat, and rice). Approaches differ: Blanc & Sultan

107 (2015) and Blanc (2017) used local weather variables (and CO₂¹³⁵
 108 values) and yields but emulate across soil types using historical¹³⁶
 109 simulations and a future climate scenario (RCP8.5 over mul-¹³⁷
 110 tiple climate models); Ostberg et al. (2018) used global mean¹³⁸
 111 temperature change (and CO₂) as regressors but pattern-scale¹³⁹
 112 to emulate local yields using multiple climate scenarios; Mis-¹⁴⁰
 113 try et al. (2017) used local weather and yields and a historical¹⁴¹
 114 simulation and compare with data.¹⁴²

115 Those emulation studies all utilized time-evolving runs from¹⁴³
 116 climate RCP runs and attempt to untangle correlations between¹⁴⁴
 117 the sensitivities to input drivers to crop models. The differences¹⁴⁵
 118 in year-to-year memory in the process-based crop models and¹⁴⁶
 119 the complexity of the changes in distributions in yearly weather¹⁴⁷
 120 under RCP scenarios in climate models are two complications¹⁴⁸
 121 we seek to control for with this study. Additional recent efforts¹⁴⁹
 122 have been made to generate datasets that allow more system-¹⁵⁰
 123 atic sampling of the input variable space (also the focus of this¹⁵¹
 124 study): Makowski et al. (2015) for temperature, CO₂, and nitro-¹⁵²
 125 gen, Pirttioja et al. (2015) and Snyder et al. (2018) for tempera-¹⁵³
 126 ture, water, and CO₂, and (Fronzek et al., 2018) for temperature¹⁵⁴
 127 and water, with all studies simulating selected sites for a limited¹⁵⁵
 128 number of crops. The parameter sweep approach simplifies the¹⁵⁶
 129 emulator development because it controls for higher frequency¹⁵⁷
 130 effects in the year-to-year variability, and isolate the broad cli-¹⁵⁸
 131 matological response. The use of limited input parameter space¹⁵⁹
 132 or restricted geographic scope in previous work may impede the¹⁶⁰
 133 ability to build future projections and to understand interaction¹⁶¹
 134 effects in global process-based crop models.¹⁶²

The Global Gridded Crop Model Intercomparison (GGCMI) Phase II experiment seeks to provide a comprehensive global dataset to allow systematically exploring how process-based crop models for the major crop respond to the main climate and management drivers and their interactions. The experiment involves running a suite of process-based crop models across historical conditions perturbed by a set of defined input parameters, and was conducted as part of the Agricultural Model Intercomparison and Improvement Project (AgMIP) (Rosenzweig et al., 2013, 2014), an international effort conducted under a framework similar to the Climate Model Intercomparison Project (CMIP) (Taylor et al., 2012, Eyring et al., 2016). The GGCMI protocol builds on the AgMIP Coordinated Climate-Crop Modeling Project (C3MP) (Ruane et al., 2014, McDermid et al., 2015) and will contribute to the AgMIP Coordinated Global and Regional Assessments (CGRA) (Ruane et al., 2018, Rosenzweig et al., 2018).

GGCMI Phase II is designed to allow addressing goals such as understanding where highest-yield regions may shift under climate change; exploring future adaptive management strategies; understanding how interacting parameters affect crop yield; quantifying uncertainties across models and major drivers; and testing strategies for producing lightweight emulators of process-based models. In this paper, we describe the GGCMI Phase II experiments, summarize output and present initial results, demonstrate that it is tractable to emulation, and present a simple climatological emulator as a potential tool for impacts assessments.

Input variable	Abbr.	Tested range	Unit
CO ₂	C	360, 510, 660, 810	ppm
Temperature	T	-1, 0, 1, 2, 3, 4, 5*, 6	°C
Precipitation	W	-50, -30, -20, -10, 0, 10, 20, 30, (and W _{inf})	%
Applied nitrogen	N	10, 60, 200	kg ha ⁻¹

Table 1: Phase II input variable test levels. Temperature and precipitation values indicate the perturbations from the historical, climatology. * Only simulated by one model. W-percentage does not apply to the irrigated (W_{inf}) simulations.

163 **2. Materials and Methods**

164 *2.1. GGCMI Phase II: experiment design*

165 GGCMI Phase II is the continuation of a multi-model com-
166 parison exercise begun in 2014. The initial Phase I compared
167 harmonized yields of 21 models for 19 crops over a historical
168 (1980-2010) scenario with a primary goal of model evalua-
169 (Elliott et al., 2015, Müller et al., 2017). Phase II compares sim-
170 ulations of 12 models for 5 crops (maize, rice, soybean, spring
171 wheat, and winter wheat) over hundreds of scenarios in which
172 individual climate or management inputs are adjusted from
173 their historical values. The reduced set of crops includes the
174 three major global cereals and the major legume and accounts
175 for over 50% of human calories (in 2016, nearly 3.5 billion tons
176 or 32% of total global crop production by weight (Food and
177 Agriculture Organization of the United Nations, 2018).

178 The major goals of GGCMI Phase II are to:

- Enhance understanding of how models work by character-
180 izing their sensitivity to input climate and nitrogen drivers.
- Study the interactions between climate variables and nitro-
182 gen inputs in driving modeled yield impacts.
- Explore differences in crop response to warming across the
184 Earth's climate regions.
- Provide a dataset that allows statistical emulation of crop
186 model responses for downstream modelers.
- Illustrate differences in potential adaptation via growing
188 season changes.

189 The guiding scientific rationale of GGCMI Phase II is to pro-
190 vide a comprehensive, systematic evaluation of the response
191 of process-based crop models to different values for carbon
192 dioxide, temperature, water, and applied nitrogen (collectively
193 known as "CTWN"). Phase II of the GGCMI project consists
194 of a series of simulations, each with one or more of the CTWN

195 dimensions perturbed over the 31-year historical time series
196 (1980-2010) used in Phase I. In most cases, historical daily cli-
197 mate inputs are taken from the 0.5 degree NASA AgMERRA
198 daily gridded re-analysis product specifically designed for agri-
199 cultural modeling, with satellite-corrected precipitation (Ruane
200 et al., 2015). Two models require sub-daily input data and use
201 alternative sources. See Elliott et al. (2015) for additional de-
202 tails.

203 The experimental protocol consists of 9 levels for precipita-
204 tion perturbations, 7 for temperature, 4 for CO₂, and 3 for ap-
205 plied nitrogen, for a total of 672 simulations for rain-fed agri-
206 culture and an additional 84 for irrigated (Table 1). For irri-
207 gated simulations, soil water is held at either field capacity or,
208 for those models that include water-log damage, at maximum
209 beneficial level. Temperature perturbations are applied as ab-
210 solute offsets from the daily mean, minimum, and maximum
211 temperature time series for each grid cell used as inputs. Pre-
212 cipitation perturbations are applied as fractional changes at the
213 grid cell level, and carbon dioxide and nitrogen levels are spec-
214 ified as discrete values applied uniformly over all grid cells.
215 Note that CO₂ changes are applied independently of changes
216 in climate variables, so that higher CO₂ is not associated with
217 higher temperatures. An additional, identical set of scenarios
218 (at the same C, T, W, and N levels) simulate adaptive agron-
219 omy under climate change by varying the growing season for
220 crop production. (These adaptation simulations are not shown
221 or analyzed here.) The resulting GGCMI data set captures a
222 distribution of crop responses over the potential space of future
223 climate conditions.

224 The 12 models included in GGCMI Phase II are all mecha-
225 nistic process-based crop models that are widely used in im-
226 pacts assessments (Table 2). Although some of the models
227 shares a common base (e.g. LPJmL and LPJ-GUESS and the
228 EPIC models), they have developed independently from this

Model (Key Citations)	Maize	Soy	Rice	Winter Wheat	Spring Wheat	N Dim.	Simulations per Crop
APSIM-UGOE , Keating et al. (2003), Holzworth et al. (2014)	X	X	X	–	X	Yes	37
CARAIB , Dury et al. (2011), Pirttioja et al. (2015)	X	X	X	X	X	No	224
EPIC-IIASA , Balkovi et al. (2014)	X	X	X	X	X	Yes	35
EPIC-TAMU , Izaurrealde et al. (2006)	X	X	X	X	X	Yes	672
JULES* , Osborne et al. (2015), Williams & Falloon (2015), Williams et al. (2017)	X	X	X	–	X	No	224
GEPIC , Liu et al. (2007), Folberth et al. (2012)	X	X	X	X	X	Yes	384
LPJ-GUESS , Lindeskog et al. (2013), Olin et al. (2015)	X	–	–	X	X	Yes	672
LPJmL , von Bloh et al. (2018)	X	X	X	X	X	Yes	672
ORCHIDEE-crop , Valade et al. (2014)	X	–	X	–	X	Yes	33
pDSSAT , Elliott et al. (2014), Jones et al. (2003)	X	X	X	X	X	Yes	672
PEPIC , Liu et al. (2016a,b)	X	X	X	X	X	Yes	130
PROMET*† , Mauser & Bach (2015), Hank et al. (2015), Mauser et al. (2009)	X	X	X	X	X	Yes†	239
Totals	12	10	11	9	12	–	3993 (maize)

Table 2: Models included in the GGCMI Phase II and the number of C, T, W, and N simulations performed for rain-fed crops (“Sims per Crop”), with 672 as the maximum. “N-Dim.” indicates if simulations include varying nitrogen levels; two models omit this dimension. All models provided the same set of simulations across all modeled crops, but some omitted individual crops in some cases. (For example, APSIM did not simulate winter wheat.) Irrigated simulations are provided at the level of the other covariates for each model (for an additional 84 simulations for the fully-sampled models). Geographic extent of simulation varies to some extent within a certain model for different scenarios (672 rain-fed simulations does not necessarily equal 672 climatological yields in all areas). This geographic variance only applies for areas far outside the area of currently cultivated crops. Two models (marked with *) use non-AgMERRA climate inputs. For further details on models, see Elliott et al. (2015). †PROMET provided simulations at only two nitrogen levels so is not emulated across the nitrogen dimension.

229 shared base, for more details on the genealogy of the mod-247 vated areas because cultivation will likely shift under climate
230 els see Figure S1 in Rosenzweig et al. (2014). Differences in248 change. See Figure 1 for the present-day cultivated area of
231 model structure does mean that several key factors are not stan-249 rain-fed crops, and Figure S1 in the supplemental material for
232 dardized across the experiment, including secondary soil nutri-250 irrigated crops. Some areas such as Greenland, far-northern
233 ents, carry over effects across growing years including residue251 Canada, Siberia, Antarctica, the Gobi and Sahara deserts, and
234 management and soil moisture, and extent of simulated area for252 central Australia are not simulated as they are assumed to re-
235 different crops. Growing seasons are identical across models,253 main non-arable even under an extreme climate change. Grow-
236 but vary by crop and by location on the globe. All stresses254 ing seasons are standardized across models with data adapted
237 except factors related to nitrogen, temperature, and water (e.g.,255 from several sources (Sacks et al., 2010, Portmann et al., 2008,
238 Alkalinity, salinity) are disabled. No additional nitrogen inputs,256 2010).
239 such as atmospheric deposition, are considered, but some mod-
240 els have individual assumptions on soil organic matter that may257 The participating modeling groups provide simulations at
241 release additional nitrogen through mineralization. See Rosen-258 any of four initially specified levels of participation, so the num-
242 zweig et al. (2014), Elliott et al. (2015) and Müller et al. (2017)²⁵⁹ ber of simulations varies by model, with some sampling only a
243 for further details on models and underlying assumptions. 260 part of the experiment variable space. Most modeling groups
261 Each model is run at 0.5 degree spatial resolution and covers262 simulate all five crops in the protocol, but some omitted one
244 all currently cultivated areas and much of the uncultivated land263 or more. Table 2 provides details of coverage for each model.
245 area. Coverage extends considerably outside currently culti-264 Note that the three models that provide less than 50 simulations
246 are excluded from the emulator analysis.

265 All models produce as output, crop yields ($\text{tons ha}^{-1} \text{ year}^{-1}$)
 266 for each 0.5 degree grid cell. Because both yields and yield
 267 changes vary substantially across models and across grid cells,
 268 we primarily analyze relative change from a baseline. We take
 269 as the baseline the scenario with historical climatology (i.e. T
 270 and P changes of 0). C of 360 ppm, and applied N at 200 kg
 271 ha^{-1} . We show absolute yields in some cases to illustrate geo-
 272 graphic differences in yields for a single model.

273 *2.2. Simulation model validation approach*

274 To verify the skill of the process-based models used, we re-
 275 peat the validation exercises presented in Müller et al. (2017)
 276 for GGCMI Phase I. Note however that the GGCMI Phase II
 277 simulations are designed for evaluating changes in yield but not
 278 absolute yields, and so omit the calibrations used in predict-
 279 ing modeling to account for cultivar, pest loss, and manage-
 280 ment differences. The Phase II simulations also do not repro-
 281 duce realistic nitrogen application levels for individual coun-
 282 tries, since nitrogen is one of the parameters systematically var-
 283 ied. The Müller et al. (2017) validation procedure evaluates re-
 284 sponse to year-to-year temperature and precipitation variations
 285 in a control run driven by historical climate and compares it
 286 to detrended historical yields from the FAO (Food and Agri-
 287 culture Organization of the United Nations, 2018) by calculat-
 288 ing the Pearson correlation coefficient. The procedure offers no
 289 means of assessing CO₂ fertilization, since CO₂ has been rel-
 290 atively constant over the historical data collection period. Ni-
 291 trogen data are limited for many countries, and as mentioned
 292 the GGCMI Phase II runs impose fixed and uniform nitrogen
 293 application, introducing some uncertainty into the analysis. We
 294 evaluate one or more control runs for each model, since some
 295 modeling groups provide historical runs for three different ni-
 296 trogen levels.

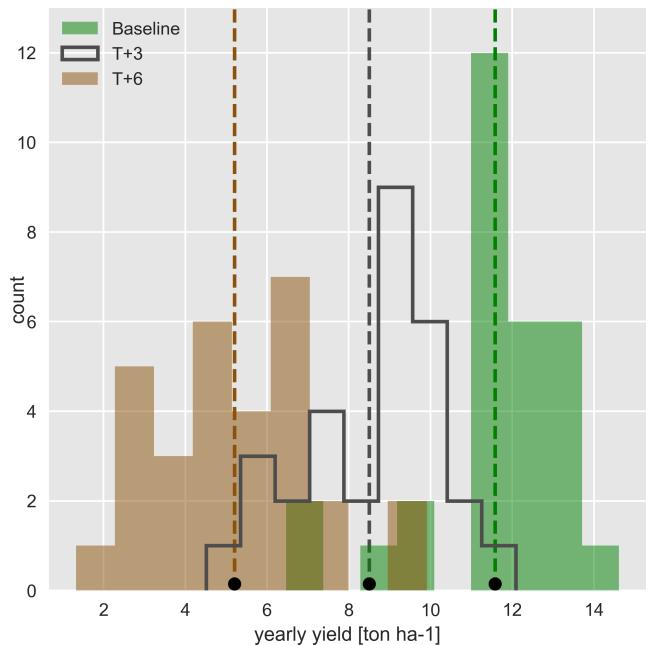


Figure 2: Example showing both climatological mean yields and distribution of yearly yields for three 30-year scenarios. Figure shows rain-fed maize for a grid cell in northern Iowa (a representative high-yield region) from the pDSSAT model, for the baseline climatology (1981–2010) and for scenarios with temperature shifted by (T) +3 and (T) +6 °C, with other variables held at baseline values. Dashed vertical lines and black dots indicate the climatological mean yield.

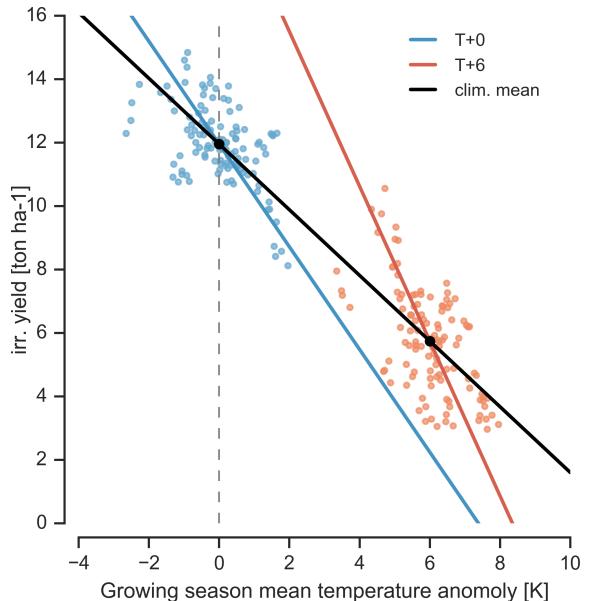


Figure 3: Example showing simple temperature relationship developed from year-to-year values vs. climatological mean values. Figure shows irrigated maize for four adjacent grid cell in northern Iowa (a representative high-yield region) from the pDSSAT model, for the baseline climatology (1981–2010) and for scenarios with temperature shifted (T) +6 °C, with other variables held at baseline values. Irrigated yields are shown to control for precipitation effects. Blue and red lines indicate total least squares linear regression across each temperature scenario. Black dots indicate the climatological mean yield values for each climatological temperature scenario.

297 2.3. Climatological-mean yield emulator design

298 To demonstrate the properties of the GGCMI Phase II
 299 dataset, we construct an emulator of 30-year climatological
 300 mean yields, which are of most interest to impact modelers.
 301 This approach differs from previous studies of crop model em-
 302 ultation, which have typically emulated at the annual level. An-
 303 nual emulation is required when the input training set consists
 304 of non-stationary projections of evolving yields (such as an
 305 RCP run). Recent studies (e.g. Fronzek et al., 2018, Snyder
 306 et al., 2018) that used a training set of stationary simulations
 307 with fixed variations in parameters allow emulating the clima-
 308 tological mean response instead. The two can differ for multiple
 309 reasons, including any year-to-year memory in the crop model,
 310 or if the distribution of growing-season daily temperatures as-
 311 sociated with interannual variability is different from that asso-
 312 ciated with long-term CO₂-driven changes. The confounding
 313 of year-to-year response with climatological response might be

314 considered a class of Simpson's paradox. Crop yields in pro-
 315 cess based models do not respond to the mean growing season
 316 temperature, they respond to the full distribution in temper-
 317 ature over the growing season (or, specifically the exact tem-
 318 perature time series). Much of the variance is left unexplained
 319 if one tries to fit a statistical model between yields and some
 320 aggregate temperature variable (mean growing season temper-
 321 ature, monthly temperature etc.). Application of relationships
 322 obtained from such statistical models to mean changes in cli-
 323 mate may provide problematic. The year-over-year yield re-
 324 sponse to individual factors in GGCMI Phase II do in fact often
 325 exceeds the climatological response (Figure 3). Note that the
 326 GGCMI Phase II datasets will not capture distributional shifts,
 327 because all simulations are run with fixed offsets from the his-
 328 torical climatology. (For methods to generate adjust histori-
 329 cal climate data inclusive of distributional changes, see Haugen
 330 et al. (2018) and Poppick et al. (2016)). Emulation approaches

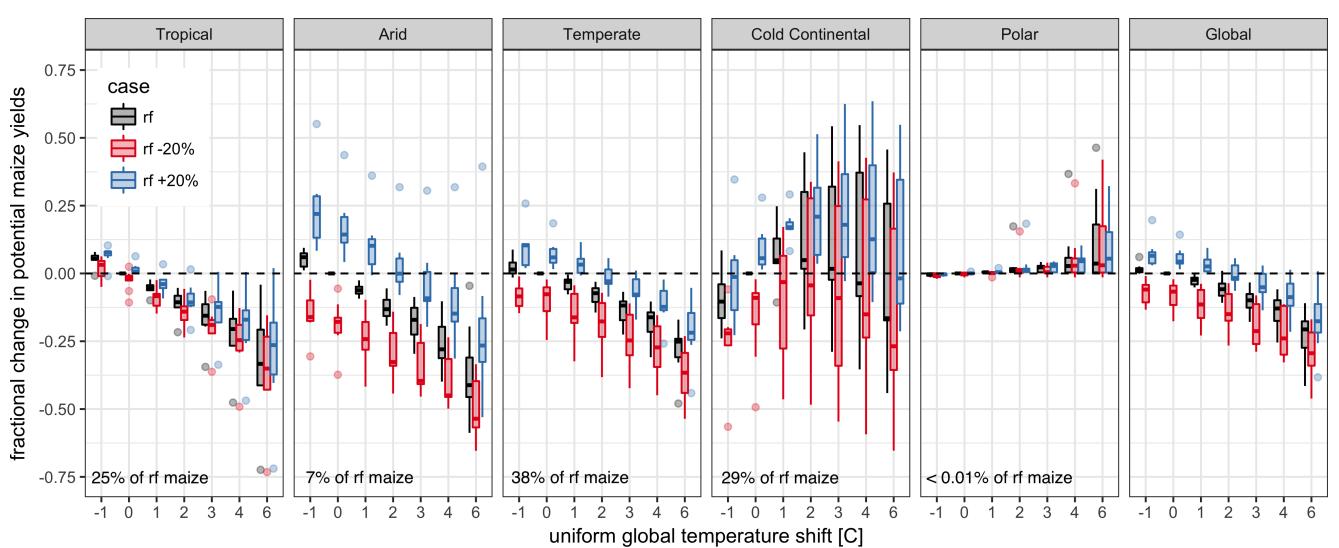


Figure 4: Illustration of the distribution of regional yield changes across the multi-model ensemble, split by Köppen-Geiger climate regions (Rubel & Kottek (2010)). We show responses of a single crop (rain-fed maize) to applied uniform temperature perturbations, for three discrete precipitation perturbation levels (rain-fed (rf) -20%, rain-fed (0), and rain-fed +20%), with CO₂ and nitrogen held constant at baseline values (360 ppm and 200 kg ha⁻¹ yr⁻¹). Y-axis is fractional change in the regional average climatological potential yield relative to the baseline. The figure shows all modeled land area; see Figure S6 in the supplemental material for only currently-cultivated land. Panel text gives the percentage of rain-fed maize presently cultivated in each climate zone (data from Portmann et al., 2010). Box-and-whiskers plots show distribution across models, with median marked. Box edges are first and third quartiles, i.e. box height is the interquartile range (IQR). Whiskers extend to maximum and minimum of simulations but are limited at 1.5-IQR; otherwise the outlier is shown. Models generally agree in most climate regions (other than cold continental), with projected changes larger than inter-model variance. Outliers in the tropics (strong negative impact of temperature increases) are the pDSSAT model; outliers in the high-rainfall case (strong positive impact of precipitation increases) are the JULES model. Inter-model variance increases in the case where precipitation is reduced, suggesting uncertainty in model response to water limitation. The right panel with global changes shows yield responses to an globally uniform temperature shift; note that these results are not directly comparable to simulations of more realistic climate scenarios with the same global mean change.

are an area of active ongoing study and one of the goals of the³⁶⁵
GGCMI Phase II dataset is to facilitate these efforts.

Emulation involves fitting individual regression models for³⁶⁷
each crop, simulation model, and 0.5 degree geographic pixel³⁶⁸
from the GGCMI Phase II data set. The regressors are the ap-³⁶⁹
plied constant perturbations in temperature, water, nitrogen and³⁷⁰
 CO_2 , we aggregate the simulation outputs in the time dimen-³⁷¹
sion, and regress on the 30-year mean yields. (See Figure 2 for³⁷²
illustration). The regression therefore omits information about³⁷³
yield responses to year-to-year climate perturbations, which are³⁷⁴
more complex. Emulating inter-annual yield variations would³⁷⁵
likely require considering statistical details of the historical cli-³⁷⁶
mate time series, including changes in marginal distribution and³⁷⁷
temporal dependencies. (Future work should explore this). The³⁷⁸
climatological emulation indirectly includes any yield response³⁷⁹
to geographically distributed factors such as soil type, insola-³⁸⁰
tion, and the baseline climate itself, because we construct sep-³⁸¹
arate emulators for each grid cell. The emulator parameter ma-³⁸²
trices are portable and the yield computations are cheap even at³⁸³
the half-degree grid cell resolution, so we do not aggregate in³⁸⁴
space at this time.

We regress climatological mean yields against a third-order³⁸⁶
polynomial in C, T, W, and N with interaction terms. The³⁸⁷
higher-order terms are necessary to capture any nonlinear re-³⁸⁸
sponses, which are well-documented in observations for tem-³⁸⁹
perature and water perturbations (e.g. Schlenker & Roberts³⁹⁰
(2009) for T and He et al. (2016) for W). We include inter-³⁹¹
action terms (both linear and higher-order) because past stud-³⁹²
ies have shown them to be significant effects. For example,³⁹³
Lobell & Field (2007) and Tebaldi & Lobell (2008) showed³⁹⁴
that in real-world yields, the joint distribution in T and W is³⁹⁵
needed to explain observed yield variance. (C and N are fixed³⁹⁶
in these data.) Other observation-based studies have shown the³⁹⁷
importance of the interaction between water and nitrogen (e.g.³⁹⁸

Aulakh & Malhi, 2005), and between nitrogen and carbon diox-
ide (Osaki et al., 1992, Nakamura et al., 1997). We do not fo-
cus on comparing different functional forms in this study, and
instead choose a relatively simple parametrization that allows
for some interpretation of coefficients. Some prior studies have
used more complex functional forms and larger numbers of pa-
rameters, e.g. 39 in Blanc & Sultan (2015) and Blanc (2017),
who borrow information across space by fitting grid points si-
multaneously across a large region in a panel regression. **We**
choose a simpler emulation at grid-cell level to avoid the re-
quirement of assuming responses are uniform across space and
to maximize interpretability.

The limited GGCMI variable sample space means that use
of the full polynomial expression described above, which has
34 terms for the rain-fed case (12 for irrigated), can be prob-
lematic, and can lead to over-fitting and unstable parameter es-
timations. We therefore reduce the number of terms through a
feature selection cross-validation process in which terms in the
polynomial are tested for importance. In this procedure higher-
order and interaction terms are added successively to the model;
we then follow the reduction of the the aggregate mean squared
error with increasing terms and eliminate those terms that do
not contribute significant reductions. See supplemental docu-
ments for more details. We select terms by applying the feature
selection process to the three models that provided the com-
plete set of 672 rain-fed simulations (pDSSAT, EPIC-TAMU,
and LPJmL); the resulting choice of terms is then applied for
all emulators.

Feature importance is remarkably consistent across all three
models and across all crops (see Figure S4 in the supplemental
material). The feature selection process results in a final poly-
nomial in 23 terms, with 11 terms eliminated. We omit the N^3
term, which cannot be fitted because we sample only three ni-
trogen levels. We eliminate many of the C terms: the cubic,

399 the CT, CTN, and CWN interaction terms, and all higher order
 400 interaction terms in C. Finally, we eliminate two 2nd-order in-
 401 teraction terms in T and one in W. Implication of this choice
 402 include that nitrogen interactions are complex and important,
 403 and that water interaction effects are more nonlinear than those
 404 in temperature. The resulting statistical model (Equation 1) is
 405 used for all grid cells, models, and crops:

$$Y = K_1 + K_2C + K_3T + K_4W + K_5N + K_6C^2 + K_7T^2 + K_8W^2 + K_9N^2 + K_{10}CW + K_{11}CN + K_{12}TW + K_{13}TN + K_{14}WN + K_{15}T^3 + K_{16}W^3 + K_{17}TWN + K_{18}T^2W + K_{19}W^2T + K_{20}W^2N + K_{21}N^2C + K_{22}N^2T + K_{23}N^2W \quad (1)$$

406 To fit the parameters K , we use a Bayesian Ridge probabilis-
 407 tic estimator (MacKay, 1991), which reduces volatility in pa-
 408 rameter estimates when the sampling is sparse, by weighting
 409 parameter estimates towards zero. The Bayesian Ridge method
 410 is necessary to maintain a consistent functional form across all
 411 models, and locations as the linear least squares fails to pro-
 412 vide a stable result in many cases. In the GGCMI Phase II
 413 experiment, the most problematic fits are those for models that
 414 provided a limited number of cases or for low-yield geographic
 415 regions where some modeling groups did not run all scenarios.
 416 Because we do not attempt to emulate models that provided
 417 less than 50 simulations, the lowest number of simulations em-
 418 ualized across the full parameter space is 130 (for the PEPIC
 419 model). We use the implementation of the Bayesian Ridge esti-
 420 mator from the scikit-learn package in Python (Pedregosa et al.,
 421 2011).

422 The resulting parameter matrices for all crop model emula-

tors are available on request, as are the raw simulation data and a Python application to emulate yields. The yield output for a single GGCMI model that simulates all scenarios and all five crops is ~ 12.5 GB; the emulator is ~ 100 MB, a reduction by over two orders of magnitude.

428 2.4. Emulator evaluation

429 Because no general criteria exist for defining an acceptable
 430 model emulator, we develop a metric of emulator performance
 431 specific to GGCMI. For a multi-model comparison exercise like
 432 GGCMI, a reasonable criterion is what we term the “normalized
 433 error”, which compares the fidelity of an emulator for a given
 434 model and scenario to the inter-model uncertainty. We define
 435 the normalized error e for each scenario as the difference be-
 436 tween the fractional yield change from the emulator and that in
 437 the original simulation, divided by the standard deviation of the
 438 multi-model spread (Equations 2 and 3):

$$F_{scn.} = \frac{Y_{scn.} - Y_{baseline}}{Y_{baseline}} \quad (2)$$

$$e_{scn.} = \frac{F_{em, scn.} - F_{sim, scn.}}{\sigma_{sim, scn.}} \quad (3)$$

Here $F_{scn.}$ is the fractional change in a model’s mean emulated or simulated yield from a defined baseline, in some scenario (scn.) in C, T, W, and N space; $Y_{scn.}$ and $Y_{baseline}$ are the absolute emulated or simulated mean yields. The normalized error e is the difference between the emulated fractional change in yield and that actually simulated, normalized by σ_{sim} , the standard deviation in simulated fractional yields $F_{sim, scn.}$ across all models. The emulator is fit across all available simulation outputs, and then the error is calculated across the simulation scenarios provided by all nine models (Figure 9 and Figures S12 and Figures S13 in supplemental documents). Note that the normalized error e for a model depends not only on the fidelity of its emulator in reproducing a given simulation but on

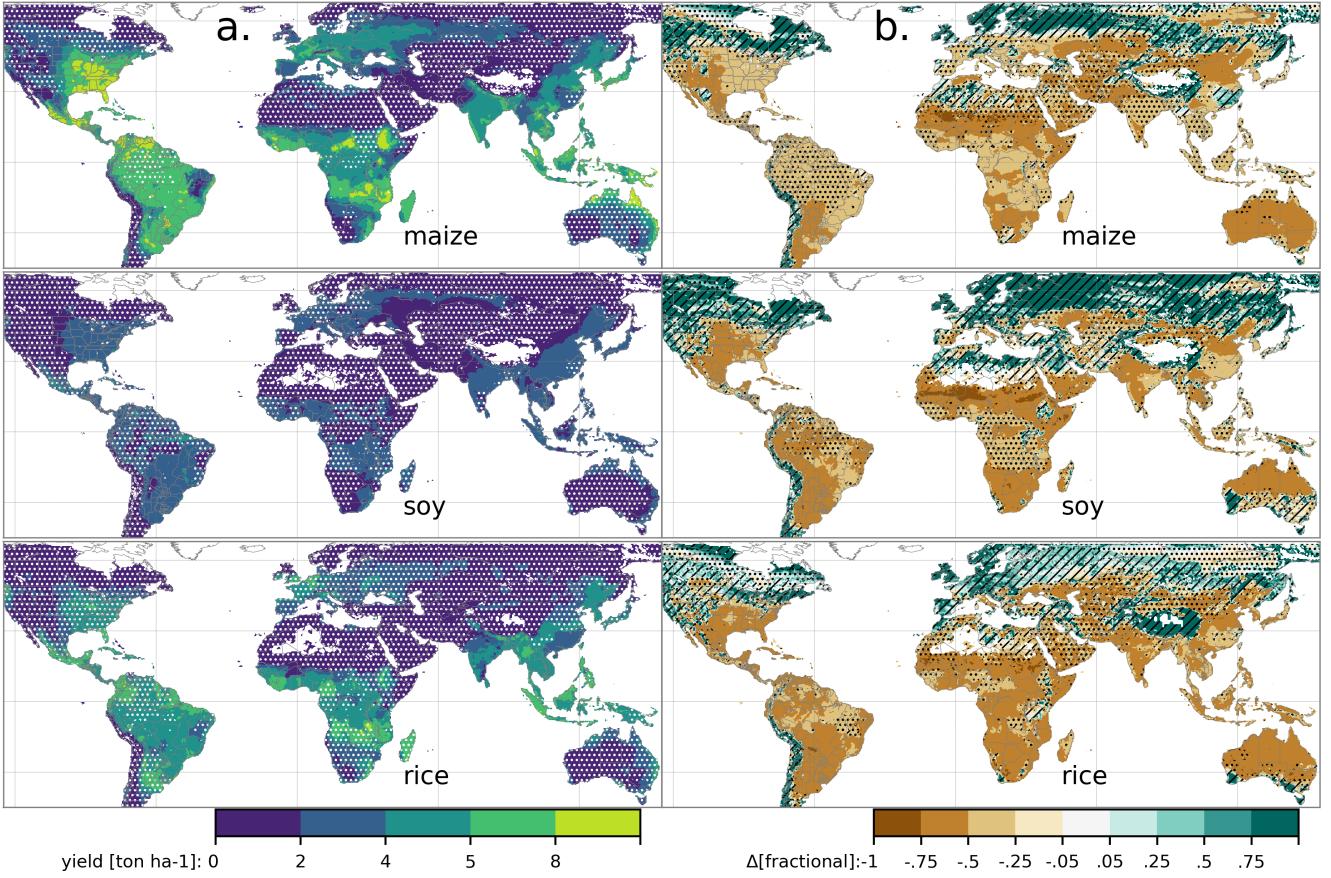


Figure 5: Illustration of the spatial pattern of potential yields and potential yield changes in the GGCMI Phase II ensemble, for three major crops. Left column (a) shows multi-model mean climatological yields for the baseline scenario for (top–bottom) rain-fed maize, soy, and rice. (For wheat see Figure S11 in the supplemental material.) White stippling indicates areas where these crops are not currently cultivated. Absence of cultivation aligns well with the lowest yield contour (0.2 ton ha^{-1}). Right column (b) shows the multi-model mean fractional yield change in the extreme $T + 4 \text{ }^{\circ}\text{C}$ scenario (with other inputs at baseline values). Areas without hatching or stippling are those where confidence in projections is high: the multi-model mean fractional change exceeds two standard deviations of the ensemble. ($\Delta > 2\sigma$). Hatching indicates areas of low confidence ($\Delta < 1\sigma$), and stippling areas of medium confidence ($1\sigma < \Delta < 2\sigma$). Crop model results in cold areas, where yield impacts are on average positive, also have the highest uncertainty.

the particular suite of models considered in the intercomparison exercise. The rationale for this choice is to relate the fidelity of the emulation to an estimate of true uncertainty, which we take as the multi-model spread.

3. Results

3.1. Simulation results

Crop models in the GGCMI ensemble show a broadly consistent responses to climate and management perturbations in most regions, with a strong negative impact of increased temperature in all but the coldest regions. We illustrate this result for rain-fed maize in Figure 4, which shows yields for the primary Köppen-Geiger climate regions (Rubel & Kottek, 2010).

In warming scenarios, models show decreases in maize yield in the temperate, tropical, and arid regions that account for nearly three-quarters of global maize production. These impacts are robust for even moderate climate perturbations. In the temperate zone, even a 1 degree temperature rise with other variables held fixed leads to a median yield reduction that outweighs the variance across models. A 6 degree temperature rise results in median loss of $\sim 25\%$ of yields with a signal to noise of nearly three. A notable exception is the cold continental region, where models disagree strongly, extending even to the sign of impacts. Model simulations of other crops produce similar responses to warming, with robust yield losses in warmer locations and high inter-model variance in the cold continental regions (Figures

477 S7).

478 The effects of rainfall changes on maize yields are also as ex-⁵¹²
479 pected and are consistent across models. Increased rainfall mit-⁵¹³
480 igates the negative effect of higher temperatures, most strongly⁵¹⁴
481 in arid regions. Decreased rainfall amplifies yield losses and⁵¹⁵
482 also increases inter-model variance more strongly, suggesting⁵¹⁶
483 that models have difficulty representing crop response to water⁵¹⁷
484 stress. We show only rain-fed maize here; see Figure S5 for the⁵¹⁸
485 irrigated case. As expected, irrigated crops are more resilient to⁵¹⁹
486 temperature increases in all regions, especially so where water⁵²⁰
487 is limiting.

488 Mapping the distribution of baseline yields and yield changes⁵²¹
489 shows the geographic dependencies that underlie these results.⁵²²
490 Figure 5 shows baseline and changes in the T+4 scenario for⁵²³
491 rain-fed maize, soy, and rice in the multi-model ensemble mean,⁵²⁴
492 with locations of model agreement marked. Absolute yield po-⁵²⁵
493 tentials are have strong spatial variation, with much of the⁵²⁶
494 Earth's surface area unsuitable for any given crop. In general,⁵²⁷
495 models agree most on yield response in regions where yield⁵²⁸
496 potentials are currently high and therefore where crops are cur-⁵²⁹
497 rently grown. Models show robust decreases in yields at low⁵³⁰
498 latitudes, and highly uncertain median increases at most high⁵³¹
499 latitudes. For wheat crops see Figure S11; wheat projections⁵³²
500 are both more uncertain and show fewer areas of increased yield⁵³³
501 in the inter-model mean.

502 3.2. Simulation model validation results

503 Figure 6 shows the Pearson time series correlation between⁵³⁷
504 the simulation model yield and FAO yield data. Figure 6 can be⁵³⁸
505 compared to Figures 1,2,3,4 and 6 in Müller et al. (2017). The⁵³⁹
506 results are mixed, with many regions for rice and wheat be-⁵⁴⁰
507 ing difficult to model. No single model is dominant, with each⁵⁴¹
508 model providing near best-in-class performance in at least one⁵⁴²
509 location-crop combination. The presence of very few vertical⁵⁴³
510 dark green color bars clearly illustrates the power of a multi-⁵⁴⁴

511 model intercomparison project like the one presented here. The
ensemble mean does not beat the best model in each case, but
shows positive correlation in over 75% of the cases presented
here. The EPIC-TAMU model performs best for soy, CARIAB,
EPIC-TAMU, and PEPIC perform best for maize, PROMET
performs best for wheat, and the EPIC family of models per-
form best for rice. Reductions in skill over the performance
illustrated in Müller et al. (2017) can be attributed to the nitro-
gen levels or lack of calibration in some models.

512 *** or harmonization *** Christoph

513 Soy is qualitatively the easiest crop to represent (except in
Argentina), which is likely due in part to the invariance of the
response to nitrogen application (soy fixes atmospheric nitrogen
very efficiently). Comparison to the FAO data is therefore easier
than the other crops because the nitrogen application levels do
not matter. US maize has the best performance across models,
with nearly every model representing the historical variability
to a reasonable extent. Especially good example years for US
maize are 1983, 1988, and 2004 (top left panel of Figure 6),
where every model gets the direction of the anomaly compared
to surrounding years correct. 1983 and 1988 are famously bad
years for US maize along with 2012 (not shown). US maize
is possibly both the most uniformly industrialized (in terms of
management practices) crop and the one with the best data col-
lection in the historical period of all the cases presented here.

514 The FAO data is at least one level of abstraction from ground
truth in many cases, especially in developing countries. The
failure of models to represent the year-to-year variability in rice
in some countries in southeast Asia is likely partly due to model
failure and partly due to lack of data. It is possible to speculate
that the difference in performance between Pakistan (no suc-
cessful models) and India (many successful models) for rice
may reside at least in part in the FAO data and not the mod-
els themselves. The same might apply to Bangladesh and In-

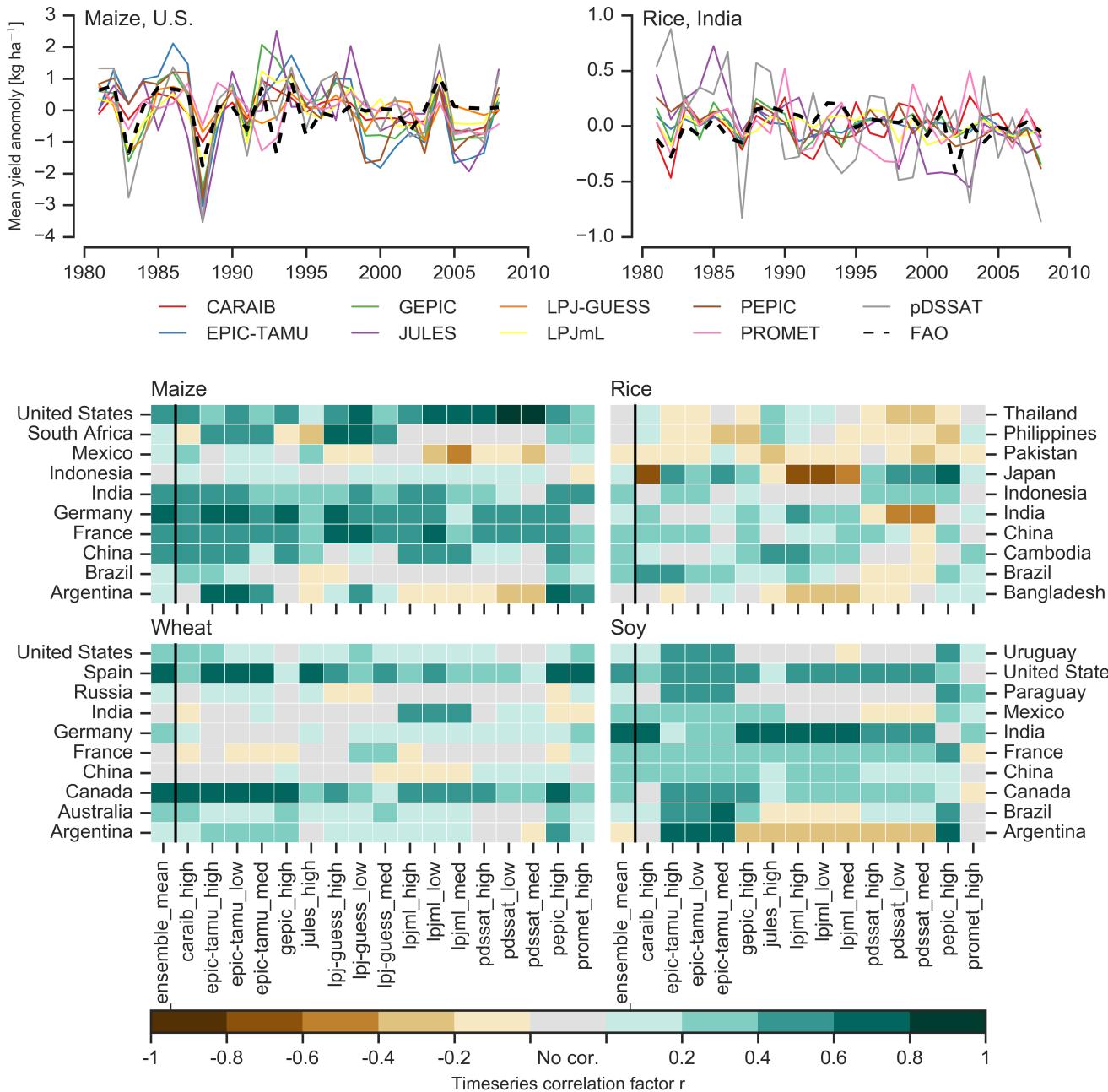


Figure 6: Time series correlation coefficients between simulated crop yield and FAO data (Food and Agriculture Organization of the United Nations, 2018) at the country level. The top panels indicate two example cases: US maize (a good case), and rice in India (mixed case), both for the high nitrogen application case. The heatmaps illustrate the Pearson r correlation coefficient between the detrended simulation mean yield at the country level compared to the detrended FAO yield data for the top producing countries for each crop with continuous FAO data over the 1980–2010 period. Models that provided different nitrogen application levels are shown with low, med, and high label (models that did not simulate different nitrogen levels are analogous to a high nitrogen application level). The ensemble mean yield is also correlated with the FAO data (not the mean of the correlations). Wheat contains both spring wheat and winter wheat simulations.

dia for rice. Partitioning of these contributions is impossible at this stage. Additionally, there is less year-to-year variability in rice yields (partially due to the fraction of irrigated cultivation). Since the Pearson r metric is scale invariant, it will tend to score the rice models more poorly than maize and soy. An example of very poor performance can be seen with the pDSSAT model for rice in India (top right panel of Figure 6).

3.3. Emulator performance

Emulation provides not only a computational tool but a means of understanding and interpreting crop yield response

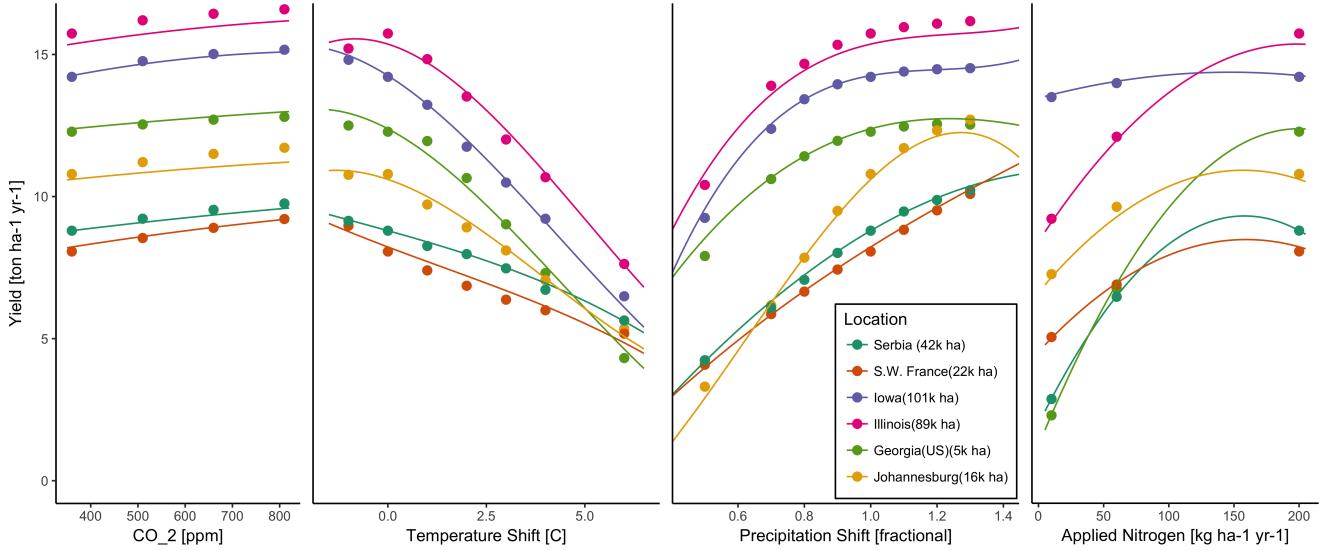


Figure 7: Example illustrating spatial variations in yield response and emulation ability. We show rain-fed maize in the pDSSAT model in six example locations selected to represent high-cultivation areas around the globe. Legend includes hectares cultivated in each selected grid cell. Each panel shows variation along a single variable, with others held at baseline values. Dots show climatological mean yield, solid lines a simple polynomial fit across this 1D variation, and dashed lines the results of the full emulator of Equation 1. The climatological response is sufficiently smooth to be represented by a simple polynomial. Because precipitation changes are imposed as multiplicative factors rather than additive offsets, locations with higher baseline precipitation have a larger absolute spread in inputs sampled than do drier locations (not visible here).

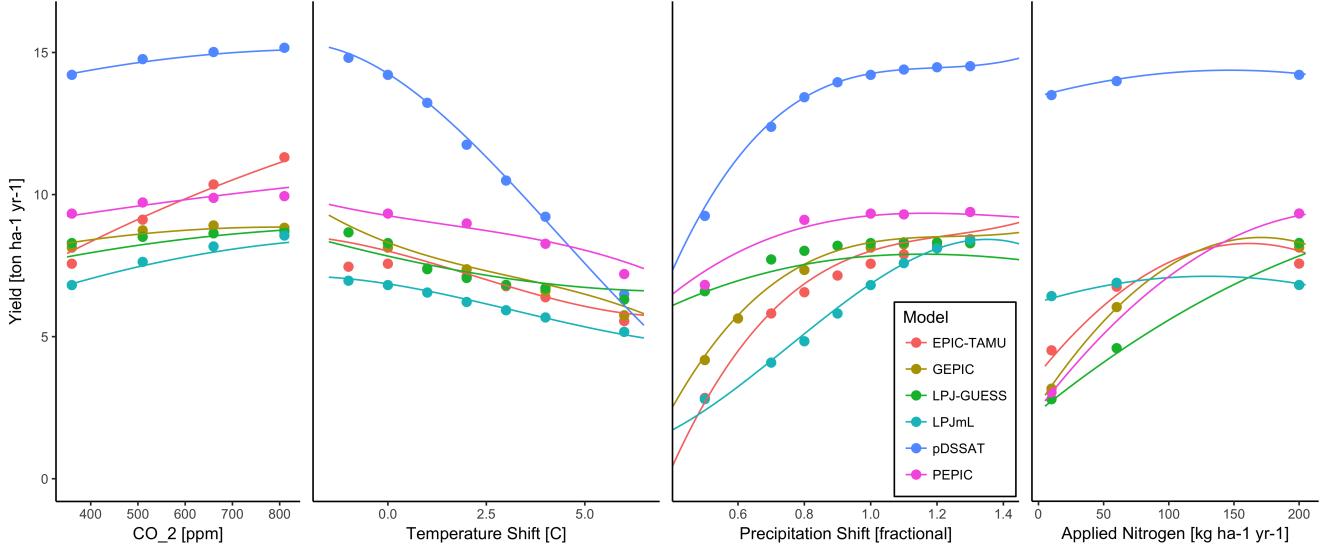


Figure 8: Example illustrating across-model variations in yield response. We show simulations and emulations from six models for rain-fed maize in the same Iowa location shown in Figure 7, with the same plot conventions. Models that did not simulate the nitrogen dimension are omitted for clarity. The inter-model standard deviation is larger for the perturbations in CO₂ and nitrogen than for those in temperature or precipitation illustrating that the sensitivity to weather is better constrained than to management at elevated CO₂. While most model responses can readily be emulated with a simple polynomial, some response surfaces are more complicated and lead to emulation error.

across the parameter space. Emulation is only possible, however, when crop yield responses are sufficiently smooth and continuous to allow fitting with a relatively simple functional form. In the GGCMI simulations, this condition largely but not always holds. Responses are quite diverse across locations,

crops, and models, but in most cases local responses are regular enough to permit emulation. Figure 7 illustrates the geographic diversity of responses even in high-yield areas for a single crop and model (rain-fed maize in pDSSAT for various high-cultivation areas). This heterogeneity validates the choice

565 of emulating at the grid cell level.

566 Each panel in Figure 7 shows model yield output from sce-⁶⁰⁰
567 narios varying only along a single dimension (CO_2 , tempera-⁶⁰¹
568 ture, precipitation, or nitrogen addition), with other inputs held⁶⁰²
569 fixed at baseline levels; in all cases yields evolve smoothly⁶⁰³
570 across the space sampled. For reference we show the results⁶⁰⁴
571 of the full emulation fitted across the parameter space. The⁶⁰⁵
572 polynomial fit readily captures the climatological response to⁶⁰⁶
573 perturbations.

574 Crop yield responses generally follow similar functional⁶⁰⁸
575 forms across models, though with a spread in magnitude. Fig-⁶⁰⁹
576 ure 8 illustrates the inter-model diversity of yield responses⁶¹⁰
577 to the same perturbations, even for a single crop and location⁶¹¹
578 (rain-fed maize in northern Iowa, the same location shown in⁶¹²
579 the Figure 7). The differences make it important to construct⁶¹³
580 emulators separately for each individual model, and the fidelity⁶¹⁴
581 of emulation can also differ across models. This figure illus-⁶¹⁵
582 trates a common phenomenon, that models differ more in re-⁶¹⁶
583 sponse to perturbations in CO_2 and nitrogen perturbations than⁶¹⁷
584 to those in temperature or precipitation. (Compare also Figures⁶¹⁸
585 4 and S18.) For this location and crop, CO_2 fertilization effects⁶¹⁹
586 can range from $\sim 5\text{--}50\%$, and nitrogen responses from nearly⁶²⁰
587 flat to a 60% drop in the lowest-application simulation.

588 While the nitrogen dimension is important and uncertain, it⁶²²
589 is also the most problematic to emulate in this work because⁶²³
590 of its limited sampling. The GGCMI protocol specified only⁶²⁴
591 three nitrogen levels (10, 60 and 200 $\text{kg N y}^{-1} \text{ha}^{-1}$), so a third-⁶²⁵
592 order fit would be over-determined but a second-order fit can⁶²⁶
593 result in potentially unphysical results. Steep and nonlinear de-⁶²⁷
594 clines in yield with lower nitrogen levels means that some re-⁶²⁸
595 gressions imply a peak in yield between the 100 and 200 kg N_{629}
596 $\text{y}^{-1} \text{ha}^{-1}$ levels. While there may be some reason to believe⁶³⁰
597 over-application of nitrogen at the wrong time in the growing⁶³¹
598 season could lead to reduced yields, these features are almost⁶³²

599 certainly an artifact of under sampling. In addition, the polyno-
600 mial fit cannot capture the well-documented saturation effect
601 of nitrogen application (e.g. Ingestad, 1977) as accurately as
602 would be possible with a non-parametric model.

To assess the ability of the polynomial emulation to capture the behavior of complex process-based models, we evaluate the normalized emulator error. That is, for each grid cell, model, and scenario we evaluate the difference between the model yield and its emulation, normalized by the inter-model standard deviation in yield projections. This metric implies that emulation is generally satisfactory, with several distinct exceptions. Almost all model-crop combination emulators have normalized errors less than one over nearly all currently cultivated hectares (Figure 9), but some individual model-crop combinations are problematic (e.g. PROMET for rice and soy, JULES for soy and winter wheat, Figures S14–S15). Normalized errors for soy are somewhat higher across all models not because emulator fidelity is worse but because models agree more closely on yield changes for soy than for other crops (see Figure S16, lowering the denominator). Emulator performance often degrades in geographic locations where crops are not currently cultivated. Figure 10 shows a CARAIB case as an example, where emulator performance is satisfactory over cultivated areas for all crops other than soy, but uncultivated regions show some problematic areas.

It should be noted that this assessment metric is relatively forgiving. First, each emulation is evaluated against the simulation actually used to train the emulator. Had we used a spline interpolation the error would necessarily be zero. Second, the performance metric scales emulator fidelity not by the magnitude of yield changes but by the inter-model spread in those changes. Where models differ more widely, the standard for emulators becomes less stringent. Because models disagree on the magnitude of CO_2 fertilization, this effect is readily seen

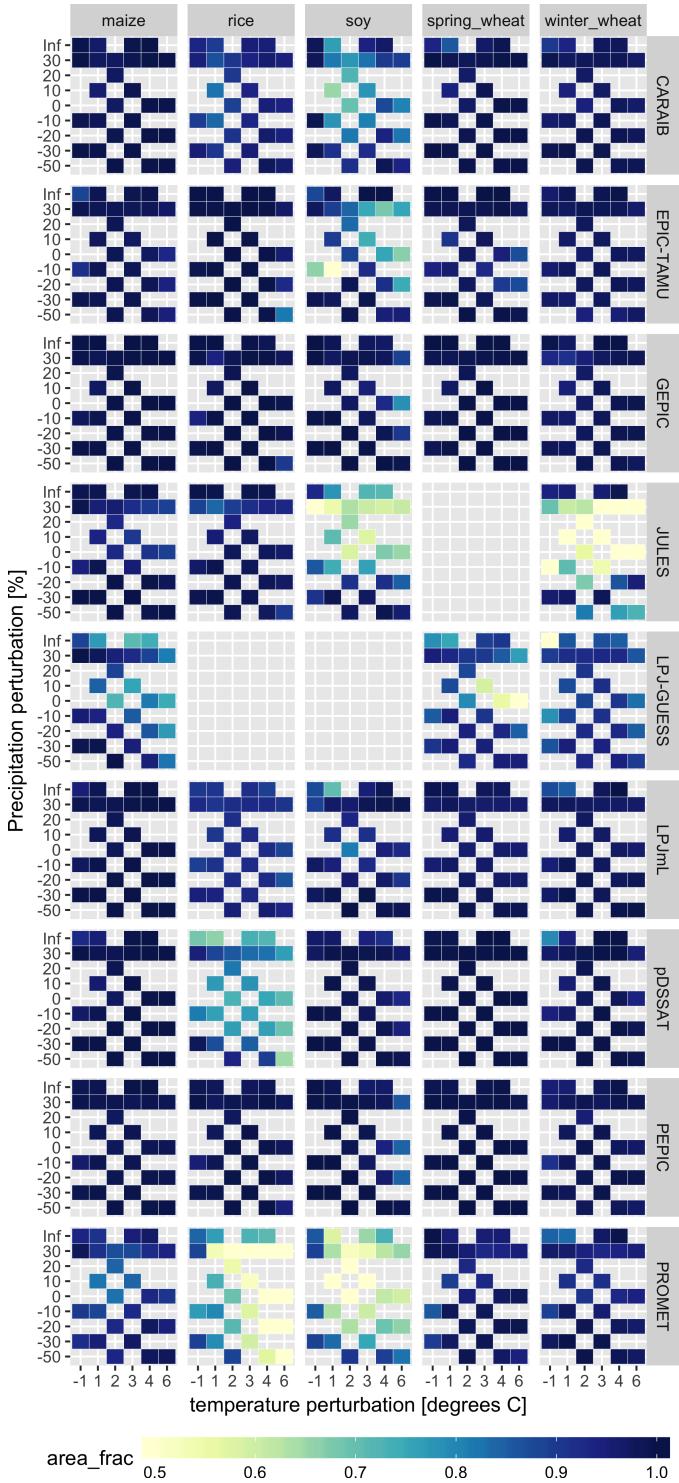


Figure 9: Assessment of emulator performance over currently cultivated areas based on normalized error (Equations 3, 2). We show performance of all 9 models emulated, over all crops and all sampled T and P inputs, but with CO₂ and nitrogen held fixed at baseline values. Large columns are crops and large rows models; squares within are T,P scenario pairs. Colors denote the fraction of currently cultivated hectares with $e < 1$. Of the 756 scenarios with these CO₂ and N values, we consider only those for which all 9 models submitted data. JULES did not simulate spring wheat and LPJ-GUESS did not simulate rice and soy. Emulator performance is generally satisfactory, with some exceptions. Emulator failures (significant areas of poor performance) occur for individual crop-model combinations, with performance generally degrading for hotter and wetter scenarios.

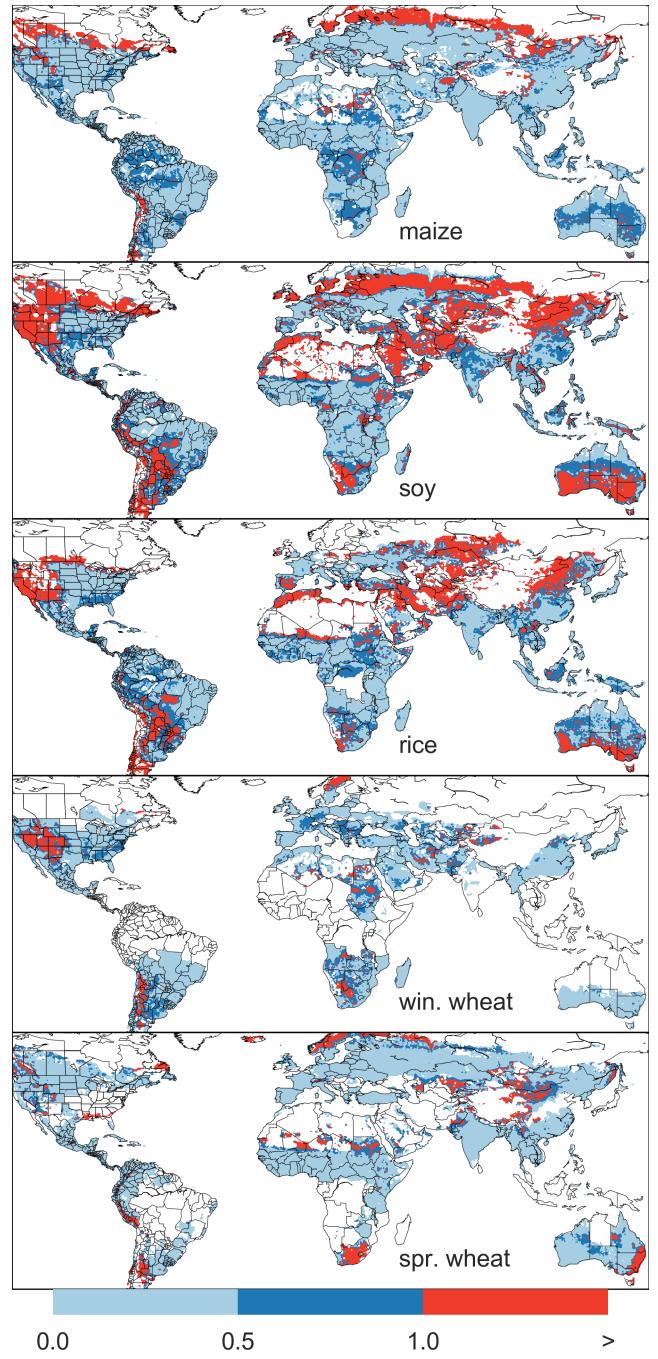


Figure 10: Illustration of our test of emulator performance, applied to the CARAIB model for the T+4 scenario for rain-fed crops. Contour colors indicate the normalized emulator error e , where $e > 1$ means that emulator error exceeds the multi-model standard deviation. White areas are those where crops are not simulated by this model. Models differ in their areas omitted, meaning the number of samples used to calculate the multi-model standard deviation is not spatially consistent in all locations. Emulator performance is generally good relative to model spread in areas where crops are currently cultivated (compare to Figure 1) and in temperate zones in general; emulation issues occur primarily in marginal areas with low yield potentials. For CARAIB, emulation of soy is more problematic, as was also shown in Figure 9.

when comparing assessments of emulator performance in simulations at baseline CO₂ (Figure 9) with those at higher CO₂ levels (Figure S13). Widening the inter-model spread leads to an apparent increase in emulator skill.

3.4. Emulator applications

Because the emulator or “surrogate model” transforms the discrete simulation sample space into a continuous response surface at any geographic scale, it can be used for a variety of applications. Emulators provide an easy way to compare a ensembles of climate or impacts projections. They also provide a means for generating continuous damage functions. As an example, we show a damage function constructed from 4D emulations for aggregated yield at the global scale, for maize on currently cultivated land, with simulated values shown for comparison. (Figure 11; see Figures S16- S19 in the supplemental material for other crops and dimensions.) The emulated values closely match simulations even at this aggregation level. Note that these functions are presented only as examples and do not represent true global projections, because they are developed from simulation data with a uniform temperature shift while increases in global mean temperature should manifest non-uniformly. The global coverage of the GGCMI simulations allows impacts modelers to apply arbitrary geographically-varying climate projections, as well as arbitrary aggregation mask, to develop damage functions for any climate scenario and any geopolitical or geographic level.

4. Conclusions and discussion

The GGCMI Phase II experiment provides a database targeted to allow detailed study of crop yields from process-based models under climate change. The experiment is designed to facilitate not only comparing the sensitivities of process-based crop yield models to changing climate and management inputs

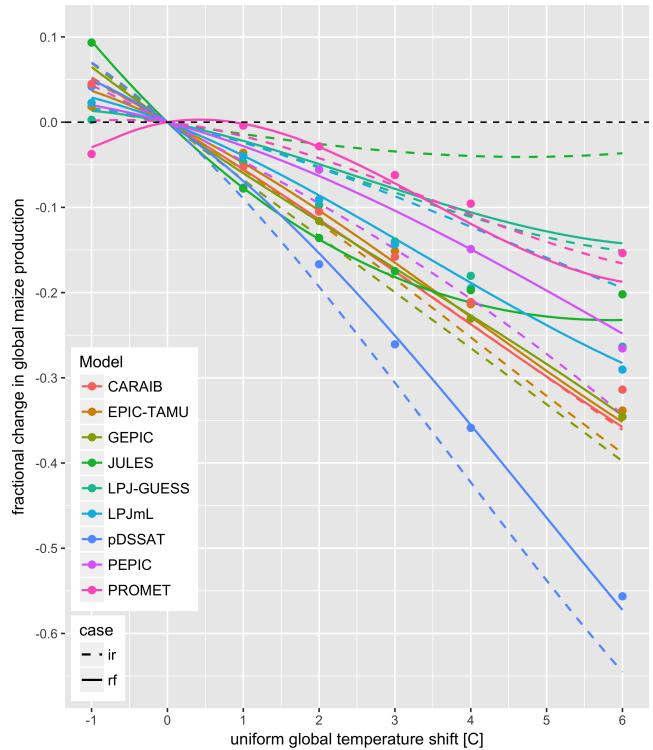


Figure 11: Global emulated damages for maize on currently cultivated lands for the GGCMI models emulated, for uniform temperature shifts with other inputs held at baseline. (The damage function is created from aggregating up emulated values at the grid cell level, not from a regression of global mean yields.) Lines are emulations for rain-fed (solid) and irrigated (dashed) crops; for comparison, dots are the simulated values for the rain-fed case. For most models, irrigated crops show a sharper reduction than do rain-fed because of the locations of cultivated areas: irrigated crops tend to be grown in warmer areas where impacts are more severe for a given temperature shift. (The exceptions are PROMET, JULES, and LPJmL.) For other crops and scenarios see Figures S16- S19 in the supplemental material.

but also evaluating the complex interactions between driving factors (CO₂, temperature, precipitation, and applied nitrogen). Its global nature also allows identifying geographic shifts in high yield potential locations. We expect that the simulations will yield multiple insights in future studies, and show here a selection of preliminary results to illustrate their potential uses.

First, the GGCMI Phase II simulations allow identifying major areas of uncertainty. Across the major crops, inter-model uncertainty is greatest for wheat and least for soy. Across factors impacting yields, inter-model uncertainty is largest for CO₂ fertilization and nitrogen response effects. Across geographic regions, projections are most uncertain in the high latitudes where yields may increase, and most robust in low latitudes

678 where yield impacts are largest.

679 Second, the GGCMI Phase II simulations allow understand-⁷¹³
680 ing the way that climate-driven changes and locations of cul-⁷¹⁴
681 tivated land combine to produce yield impacts. One coun-⁷¹⁵
682 terintuitive result immediate apparent is that irrigated maize⁷¹⁶
683 shows steeper yield reductions under warming than does rain-⁷¹⁷
684 fed maize when considered only over currently cultivated land.⁷¹⁸
685 The effect results from geographic differences in cultivation. In⁷¹⁹
686 any given location, irrigation increases crop resiliency to tem-⁷²⁰
687 perature increase, but irrigated maize is grown in warmer loca-⁷²¹
688 tions where the impacts of warming are more severe (Figures⁷²²
689 S5–S6). The same behavior holds for rice and winter wheat,⁷²³
690 but not for soy or spring wheat (Figures S8–S10). Irrigated⁷²⁴
691 wheat and maize are also more sensitive to nitrogen fertiliza-⁷²⁵
692 tion levels than are analogous non-irrigated crops, presumably⁷²⁶
693 because those rain-fed crops are limited by water as well as⁷²⁷
694 nitrogen availability (Figure S19). (Soy as an efficient atmo-⁷²⁸
695 spheric nitrogen-fixer is relatively insensitive to nitrogen, and⁷²⁹
696 rice is not generally grown in water-limited conditions).

697 Third, we show that even the relatively limited GGCMI⁷³¹
698 Phase II sampling space allows emulation of the climatologi-⁷³²
699 cal response of crop models with a relatively simple reduced-⁷³³
700 form statistical model. The systematic parameter sampling in⁷³⁴
701 the GGCMI Phase II procedure provides information on the in-⁷³⁵
702 fluence of multiple interacting factors in a way that single pro-⁷³⁶
703 jections cannot, and emulating the resulting response surface⁷³⁷
704 then produces a tool that can aid in both physical interpretation⁷³⁸
705 of the process-based models and in assessment of agricultural⁷³⁹
706 impacts under arbitrary climate scenarios. Emulating the cli-⁷⁴⁰
707 matological response isolates long-term impacts from any con-⁷⁴¹
708 founding factors that complicate year-over-year changes, and⁷⁴²
709 the use of simple functional forms offer the possibility of phys-⁷⁴³
710 ical interpretation of parameter values. Care should be taken in⁷⁴⁴
711 applying relationships developed at the yearly level to shifts in⁷⁴⁵

712 the mean climatology. We anticipate that systematic parameter
sampling will become the norm in future model intercompari-
son exercise.

While the GGCMI Phase II database should offer the foun-
dation for multiple future studies, several cautions need to be
noted. Because the simulation protocol was designed to fo-
cus on change in yield under climate perturbations and not
on replicating real-world yields, the models are not formally
calibrated so cannot be used for impacts projections unless in
used in conjunction with historical data (or data products). Be-
cause the GGCMI simulations apply uniform perturbations to
historical climate inputs, they do not sample changes in higher
order moments, and cannot address the additional crop yield
impacts of potential changes in climate variability. Although
distributional changes in model projections are fairly uncertain
at present, follow-on experiments may wish to consider them.
Several recent studies have described procedures for generating
simulations that combine historical data with model projections
of not only mean changes in temperature and precipitation but
changes in their marginal distributions (e.g. Chang et al., 2016)
or temporal dependence (e.g. Leeds et al., 2015).

The GGCMI phase II output dataset invites a broad range
of potential avenues of analysis. A major target area involves
studying the models themselves with a detailed examination of
interaction terms between the major input drivers, a more robust
quantification of the sensitivity of different models to the input
drivers, and comparison with field-level experimental data. The
parameter space tested in GGCMI phase II will allow detailed
investigations into yield variability and response to extremes
under changing management and CO₂ levels. As mentioned
previously, the database allows study of geographic shifts in op-
timal growing regions for different crops and studying the via-
bility of switching crop types in some areas. The output dataset
also contains other runs and variables not analyzed or shown

here. Runs include several which allowed adaptation to climate⁷⁷⁹
changes by altering growing seasons, and additional variables⁷⁸⁰
include above ground biomass, LAI, and root biomass (as many⁷⁸¹
as 25 output variables for some models). Emulation studies that⁷⁸²
are possible include study of year-over-year vs climatological⁷⁸³
emulation, and more systematic evaluation of different statisti-
cal model specifications and formal calculation of uncertainties⁷⁸⁴
in derived parameters.

The future of food security is one of the larger challenges⁷⁸⁵
facing humanity at present. The development of multi-model⁷⁸⁶
ensembles such as GGCMI Phase II provides a way to begin⁷⁸⁸
to better understand crop responses to a range of potential cli-⁷⁸⁹
mate inputs, improve process based models, and explore the po-⁷⁹⁰
tential benefits of adaptive responses included shifting growing⁷⁹²
season, cultivar types and cultivar geographic extent.⁷⁹³

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