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# An AgMIP framework for improved agricultural representation in integrated assessment models

To cite this article: Alex C Ruane *et al* 2017 *Environ. Res. Lett.* **12** 125003

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# Environmental Research Letters



## LETTER

# An AgMIP framework for improved agricultural representation in integrated assessment models

### OPEN ACCESS

RECEIVED  
3 March 2017

REVISED  
6 September 2017

ACCEPTED FOR PUBLICATION  
19 September 2017

PUBLISHED  
24 November 2017

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**Keywords:** AgMIP, climate impacts, crop models, integrated assessment models, agricultural impacts, emulators, model development



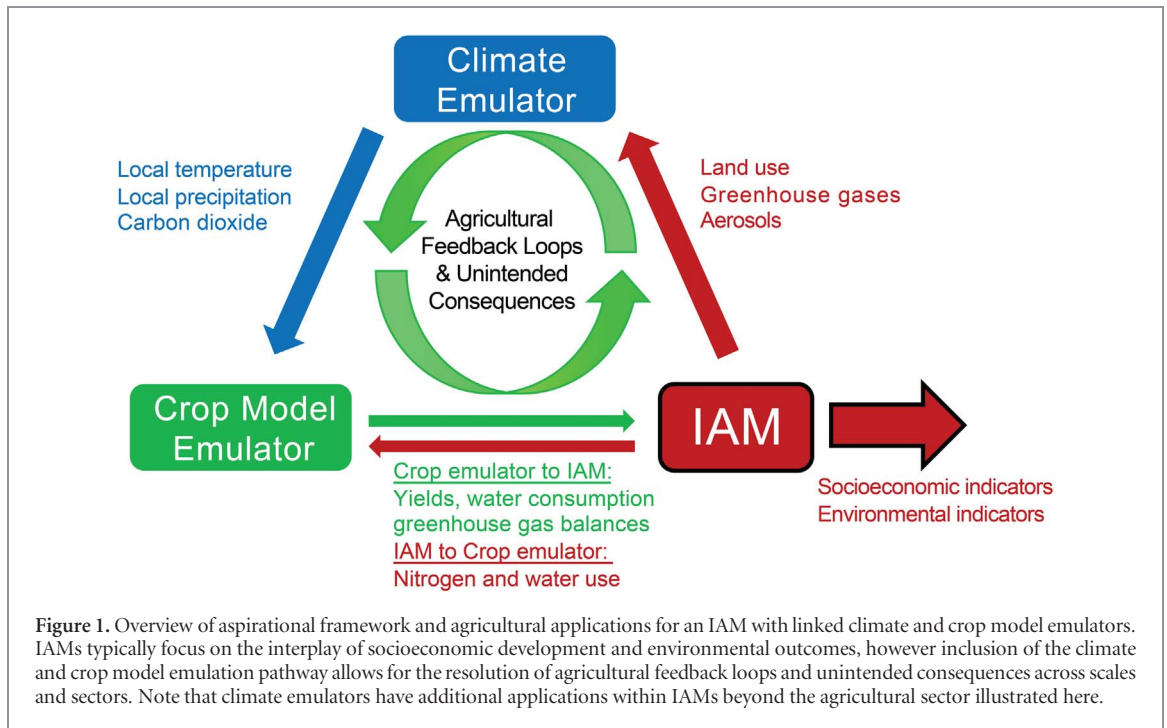
## Abstract

Integrated assessment models (IAMs) hold great potential to assess how future agricultural systems will be shaped by socioeconomic development, technological innovation, and changing climate conditions. By coupling with climate and crop model emulators, IAMs have the potential to resolve important agricultural feedback loops and identify unintended consequences of socioeconomic development for agricultural systems. Here we propose a framework to develop robust representation of agricultural system responses within IAMs, linking downstream applications with model development and the coordinated evaluation of key climate responses from local to global scales. We survey the strengths and weaknesses of protocol-based assessments linked to the Agricultural Model Intercomparison and Improvement Project (AgMIP), each utilizing multiple sites and models to evaluate crop response to core climate changes including shifts in carbon dioxide concentration, temperature, and water availability, with some studies further exploring how climate responses are affected by nitrogen levels and adaptation in farm systems. Site-based studies with carefully calibrated models encompass the largest number of activities; however they are limited in their ability to capture the full range of global agricultural system diversity. Representative site networks provide more targeted response information than broadly-sampled networks, with limitations stemming from difficulties in covering the diversity of farming systems. Global gridded crop models provide comprehensive coverage, although with large challenges for calibration and quality control of inputs. Diversity in climate responses underscores that crop model emulators must distinguish between regions and farming system while recognizing model uncertainty. Finally, to bridge the gap between bottom-up and top-down approaches we recommend the deployment of a hybrid climate response system employing a representative network of sites to bias-correct comprehensive gridded simulations, opening the door to accelerated development and a broad range of applications.

## 1. Introduction

Integrated assessment models (IAMs) examine the interactions between human systems and the natural

environment. IAMs thus explore how societal changes, such as global policies, population growth, socioeconomic development, greenhouse gas emissions, and technological advances affect land, air, and water



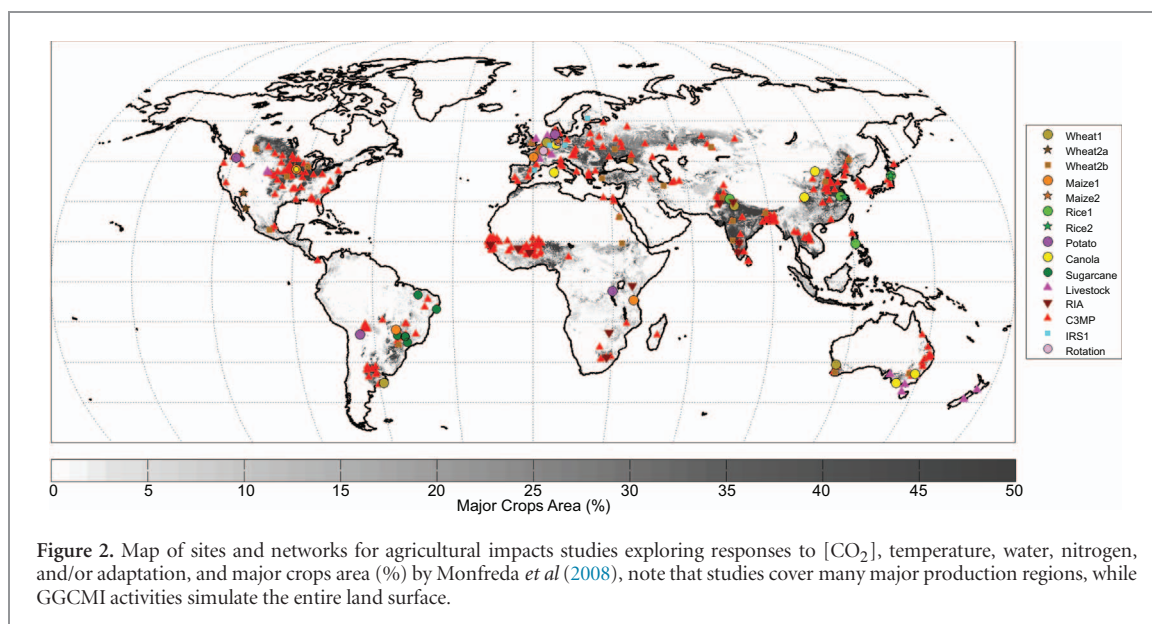
resources, as well as repercussions when these natural resources are strained (Füssel *et al* 2010, Clarke *et al* 2014). Agriculture has long been central to the relationship between society and natural systems, providing vital foods, fiber, and energy while drawing heavily on land and water resources.

IAMs have traditionally represented agricultural sector changes as exogenous yield changes provided via scenarios aggregated to national or regional level production using current harvested area weights (Müller and Robertson 2014, Nelson *et al* 2014, Wiebe *et al* 2015); however these only draw from a small subset of cutting-edge crop model assessments. A more direct coupling of agricultural responses within IAMs is facilitated by the application of crop model emulators, defined here as computationally-efficient representations of crop model results that capture fundamental responses to climate conditions. Crop model emulators may take the form of lookup tables (e.g. based upon response surfaces in Pirttioja *et al* 2015), simplified response functions (Howden and Crimp 2005, Crimp *et al* 2008, Ruane *et al* 2014, Makowski *et al* 2015), or complex statistical models (Blanc 2017, Mistry *et al* 2017, Moore *et al* 2017), each estimating yield as a function of climate variables with varying degrees of non-linearity and detail about the specific crop variety, farm environment, weather extremes, and crop model emulated. As these emulators get more complex the gain in computational efficiency (compared to just using the crop model itself) is reduced, and in the end a crop model emulator is limited by the performance of the crop model or crop model ensemble that it is emulating. Emulators are distinct from statistical crop models, which are trained upon observational data, with one advantage being that

they may use principles of biophysical process response to explore environments that have not been observed (such as future climate and land use change). The exact specifications and desired detail of a crop model emulator depends on the IAM to which it is coupled, the intended applications, and the capabilities and coverage of the underlying crop model assessments.

IAMs have a lot to gain by better incorporating crop responses to changes in carbon dioxide concentration ( $[CO_2]$ ), temperature, water, nitrogen, and adaptation (CTWNA). CTWNA sensitivity simulations can be more useful than projections driven by global climate models (GCMs) as they provide the information basis to construct crop model emulators for use in IAMs in conjunction with climate emulators (e.g. Meinshausen *et al* 2011, Castruccio *et al* 2014, Hartin *et al* 2015). Figure 1 illustrates how this powerful combination improves agricultural sector representation by allowing IAM land use changes and emissions of greenhouse gases and aerosols to influence regional temperature and precipitation changes (using the climate emulator), affecting crop production and requirements (using the crop model emulator) that feed back into the IAM. This also captures agricultural feedback loops, where societal or environmental changes alter the climate and shift agricultural production in a manner that reinforces or diminishes those changes, and unintended consequences when policies in another sector or region impact distant farming systems (potentially through climate responses or through independent mechanisms such as trade).

The Agricultural Model Intercomparison and Improvement Project (AgMIP; Rosenzweig *et al* 2013, 2015) was launched in 2010 to provide a common



framework and systematic approach for analysis of agricultural challenges. AgMIP connects climate, crop, livestock, and economic models at local, regional, and global scales, allowing multi-model, multi-discipline, multi-scale assessments of agricultural development and food security (Rosenzweig *et al* 2016, Antle *et al* 2015). AgMIP mainly utilizes process-based crop models that represent biophysical processes and their responses to genetics, environment, and management over the course of a growing season, with statistical models also included in some efforts. Integrated assessment modelers examining previous crop modeling studies have been challenged to make sense of differing assumptions, methods, and models in addition to the under-representation of agricultural systems beyond the mid-latitude, high-input breadbaskets (White *et al* 2011, Challinor *et al* 2014a). AgMIP facilitates more robust and transferable findings based on common simulation protocols, multi-model ensembles, the tracking of uncertainty, and an emphasis on under-simulated farm systems. Great strides in computational power are opening new doors for agricultural model development and application, raising the ceiling for multi-model analyses, new scales of decision support, and more accurate crop model emulators for IAM applications.

This article takes stock of the methods used by AgMIP to capture the response of agricultural productivity to changing climate conditions, examining the relative strengths and weaknesses of site, network, and gridded modeling approaches to inform IAMs and related crop model emulators. We then provide a framework for coordinated development and application of agricultural responses drawing value from local to global approaches and linking biophysical and integrated assessments. We conclude with recommendations for priority future work and applications.

## 2. Survey of crop model outputs germane to IAM emulators

Although AgMIP conducts more than 30 activities (Rosenzweig *et al* 2015), here we survey activities that (a) test for sensitivity to some or all of CTWNA factors and utilize (b) multiple agricultural models, (c) multiple sites, and (d) common protocols. These are described in table 1 along with related activities by the MACSUR project (Modelling European Agriculture with Climate Change for Food Security; Ewert *et al* 2015, Webber *et al* 2015). Figure 2 presents the geographic coverage of these site, network, and gridded activities.

### 2.1. Site-based approaches

The overwhelming majority of studies in the large literature on crop impacts are site-based studies (White *et al* 2011, Challinor *et al* 2014a), but inconsistent protocols, assumptions, geographic sampling, and methods make generalized interpretation of the results difficult. AgMIP's emphasis on model intercomparison and exploration of climate responses drove initial research activities toward species-based assessment at a small number of carefully selected sites. These 'pilot' projects organized around the application of multiple models on high-quality field datasets (Boote *et al* 2015, Kersebaum *et al* 2015) to expose differences in model structure, process responses, data requirements, and input/output formats.

The first crop pilot was organized by the AgMIP Wheat Team, in which 27 modeling groups ran historical simulations and 30 year sensitivity tests for [CO<sub>2</sub>], temperature, and nitrogen response at sites in the Netherlands, India, Argentina, and Australia (Asseng *et al* 2013, Martre *et al* 2015). The Wheat Pilot was open to all interested modeling groups as long as their models were published in peer-reviewed articles.

**Table 1.** Overview of multi-model, multi-site, protocol-based research activities sampling the carbon–temperature–water–nitrogen–adaptation (CTWNA) change space by AgMIP and related projects.

Research activity	Scope	C (ppm)	T (°C)	W (%)	N	A	Notes
AgMIP-Wheat Phase 1	Sites	360 to 720	−3 to +9	—	−50% to +50%	—	27 wheat models at each of four sites. <sup>a</sup>
AgMIP-Wheat Phase 2a	Sites	—	+0 to +16	—	—	—	30 wheat models simulated at two sites.
AgMIP-Wheat Phase 2b	Global network	—	+0 to +4	—	—	—	30 wheat models at 30 well-watered sites.
AgMIP-Wheat Phase 3	Global network	360 to 550	+0 to +4	—	—	—	32 wheat models at 60 sites (water-limited and well-watered sites).
AgMIP-Maize Phase 1	Sites	360 to 720	−3 to +9	—	—	—	23 maize models at each of four sites.
AgMIP-Maize Phase 2	Sites	387 and 550	—	RF and Irr.	—	—	21 maize models for Braunschweig, Germany, FACE site.
AgMIP-Rice Phase 1	Sites	360 to 720	−3 to +9	—	varied N	—	13 rice models at each of four sites. Two sites included N treatments ranging from 30–150 kg N ha <sup>−1</sup> .
AgMIP-Rice Phase 2	Sites	360 to 720	—	—	varied N	—	16 rice models at each of two FACE sites (Japan and China). <sup>b</sup>
AgMIP-Potato	Sites	360 to 720	−3 to +9	−30 to +30	—	—	Nine potato models at each of four sites.
AgMIP-Canola	Sites	360 to 720	−3 to +9	−25 to +25	0% to 150% of observed	—	Eight canola models at each of seven sites.
AgMIP-Sugarcane	Sites	350 to 750	−3 to +9	−30 to +30	—	—	Two sugarcane models at seven Brazilian sites.
AgMIP-Livestock and Grasslands Phase 2	Sites	330 to 900	−1 to +8	−50 to +50	—	—	Common protocols for single model tests at 14 sites. Seven models contributed yield and GHG balance results.
AgMIP Regional Integrated Assessments	Sites	360 to 720	−2 to +8	−75 to +100	0 to 210 kg N ha <sup>−1</sup>	—	Two models each for ten sites, multiple crops at many of the sites.
C3MP	Global network	330 to 900	−1 to +8	−50 to +50	—	—	1137 simulation sets in 56 countries; 18 crop species, 23 crop models.
MACSUR-IRS Phase 1	Sites	—	−2 to +9	−50 to +50	—	—	26 wheat models at four sites in Europe.
MACSUR-Crop Rotation	Sites	374 and 550	—	—	100% and 50% of observed	—	15 models with and without crop rotations. CN sensitivity tests performed at Braunschweig, Germany, and N sensitivity at Thibie, France.
GGCMI Phase 2	Global grid	360 to 810	−1 to +6	−50 to +30 plus Irr.	10 to 200 kg N ha <sup>−1</sup>	Fully reverse accelerated maturity	12 participating models. <sup>b</sup> Includes no water stress test and no nitrogen stress test. Adaptation adjusts cultivars to maintain planting to maturity duration.

Notes: RF = rainfed; Irr. = irrigated; GHG = greenhouse gas; varied N = multiple nitrogen treatments at each site.

<sup>a</sup> Nitrogen tests were only performed for 20 wheat models containing nitrogen dynamics.

<sup>b</sup> Ongoing project, final participation may change.

Similar multi-model crop pilots were conducted across selected sites by AgMIP Maize (CT responses; Bassu *et al* 2014), AgMIP Rice (CT responses; Li *et al* 2015), AgMIP Potato (CTW responses; Fleisher *et al* 2016), AgMIP Sugarcane (CTW responses; Marin *et al* 2015), and AgMIP Canola (CTWN responses; Wang *et al* 2017). MACSUR also analyzed TW responses at a transect of four European wheat

sites, providing continuous impacts response surfaces that characterize fundamental crop model properties (Pirttioja *et al* 2015, Fronzek *et al* 2017), and examined the CN response of crop rotations (Kollas *et al* 2015, Yin *et al* 2017). AgMIP's Livestock and Grasslands Team used individual models at a number of sites to create CTW response surfaces for yield and greenhouse gas balances (Erhardt 2017). Phase 2

**Table 2.** AgMIP-Wheat, AgMIP-Maize, and AgMIP-Rice Team Phase descriptions.

Phase ( <i>and key references</i> )	Description
AgMIP-Wheat	
Phase 1 (Asseng <i>et al</i> 2013, Martre <i>et al</i> 2015)	Protocol-based multi-model intercomparison at diverse, high-quality sites. Included limited information and full information calibration settings.
Phase 2a (Asseng <i>et al</i> 2015a, 2015b)	Protocol-based multi-model analysis of temperature response at Hot Serial Cereals artificial heating experiment in Arizona and temperature responses in Mexico.
Phase 2b (Asseng <i>et al</i> 2015a, 2015b, Liu <i>et al</i> 2016)	Intercomparison of temperature responses across 30 sites selected as a representative network of well-watered wheat production regions around the world.
Phase 3 (in progress)	Intercomparison of temperature responses across 60 sites selected to represent both well-watered and water-limited wheat production regions around the world.
AgMIP-Maize	
Phase 1 (Bassu <i>et al</i> 2014)	Protocol-based multi-model intercomparison at diverse, high-quality sites. Included limited information and full information calibration settings.
Phase 2 (Durand <i>et al</i> 2017)	Protocol-based multi-model intercomparison at Free-Air Carbon Enrichment (FACE) site in Germany.
AgMIP-Rice	
AgMIP-Rice Phase 1 (Li <i>et al</i> 2015)	Protocol-based multi-model intercomparison at diverse, high-quality sites. Included limited information and full information calibration settings.
AgMIP-Rice Phase 2 (in progress)	Protocol-based multi-model intercomparison at FACE Sites in Japan and China.

studies by AgMIP Wheat, Maize, and Rice teams have challenged models with field experiments that gauge climate sensitivity at test sites, utilizing ‘Hot Serial Cereal’ heat stress experiments for wheat (Asseng *et al* 2015a, Webber *et al* 2017) and Free-Air Carbon Enrichment data to explore [CO<sub>2</sub>] response in maize (Durand *et al* 2017) and rice (Hasegawa *et al* 2017) (table 2). CTWN sensitivity experiments also form a key component of AgMIP’s regional integrated assessments at sites across South Asia and Sub-Saharan Africa (Rosenzweig *et al* 2017).

### 2.1.1. Strengths and weaknesses of site-based approaches

Intensive, multi-model intercomparisons at high-quality pilot field sites are a critical first component of model evaluation, yielding valuable insight into process responses, structural biases, data requirements, and performance across contrasting systems. These analyses are anchored in field data that enable validation of state variables (e.g. leaf-area index; above-ground biomass and N contents; plant-available soil moisture) across a number of phenological stages as well as end-of-season characteristics (e.g. grain yield and protein content, harvest index). This allows evaluation of the mechanisms by which plants respond to environmental changes, highlighting sensitive biophysical processes and growth stages that in turn help focus climate projections on fundamental stresses (e.g. drought in reproductive stages; heat stress at anthesis). Results demonstrate that multi-model ensembles consistently outperform individual models when evaluated across variables and sites (Martre *et al* 2015, Asseng *et al* 2013, Bassu *et al* 2014, Li *et al* 2015, Fleisher *et al* 2016), although at any given site a subset of models may be preferred (Castañeda-Vera *et al* 2015).

Site-based assessments from the initial AgMIP Pilots are limited in their application to IAMs as they cover only a small number of sites and farming systems. As expected, crops responded differently at the selected sites owing to unique soils, weather, cultivars, and farm management. Additional careful sampling of interactions across the broader CTWNA space is needed, as shown by the benefits of elevated [CO<sub>2</sub>] for water use efficiency in recent AgMIP crop team activities (Cammarano *et al* 2016, Deryng *et al* 2016, Durand *et al* 2017).

### 2.2. Network-based approaches

As AgMIP protocols were developed and tested on individual sites, the next step scaled up these approaches through larger networks of sites coordinated to ensure adherence to a common protocol that enables direct comparison.

#### 2.2.1. Wide ad hoc network approach

AgMIP launched the Coordinated Climate-Crop Modeling Project (C3MP; Ruane *et al* 2014, McDerimid *et al* 2015a), to create CTW impact response surfaces at a range of sites around the world. C3MP samples the CTW space projected by GCMs in the 21st century, enabling the fitting of emulators and response surfaces that can be rapidly applied to estimate the agricultural impacts of new climate projections. C3MP created information technology tools and templates to facilitate the process and invited the agricultural modeling community to participate with their own calibrated sites. The resulting archive reflects submissions from 100 crop modelers, with 1137 simulation sets from 55 countries, including results from 19 crop model families and 18 crop species (McDerimid *et al* 2015a).

### 2.2.1.1. Strengths and weaknesses of wide ad hoc networks

C3MP's open call for crop model participation led to an unprecedented number and diversity of contributed simulation sets but also challenges in analyses. The result is a network of voluntary 'crowd-sourced' responses rather than a designed plan of geographic coverage, representative sites, or multi-model analyses. Nevertheless, C3MP's wide ad hoc network covers most major agricultural lands and features models calibrated with site-specific information (figure 2). Sampling across all submitted results for a given category of system (e.g. rainfed maize) provides CTW response surfaces isolating the common yield response across a broad sampling of sites and systems as well as uncertainty stemming from model, soil, baseline climate, cultivar, and farming system differences (McDermid *et al* 2015a). Recognizing that IAMs typically track major crops (wheat and rice) and commodity groups (e.g. oil seeds, coarse grains, sugar crops, fruits and vegetables), C3MP's relatively large number of crop species also reduces the amount of crop response mapping that is required to represent climate responses across the diversity of agricultural commodities. C3MP is particularly useful in distinguishing responses within a commodity group (for example, differentiating between millet, sorghum, and maize responses for coarse grains).

Aggregation of the C3MP archive to global production responses is challenging given geographic gaps and under-represented systems, and vetting is difficult given its reliance on prior model calibration and a skew toward common crop models (as were also challenges in the Challinor *et al* 2014a, meta-analysis). We recommend that C3MP analyses do not include simulation sets that use antiquated model versions and a small percentage of flagged sites where low historical yields indicate farming systems that are not presently viable. In some cases, these were conducted as tests of land uses that may become viable in wetter and high-[CO<sub>2</sub>] futures, but must be considered distinct from broader CTW analyses. C3MP remains an open process, and each new submission increases the robustness of ensemble statistics and analyses.

### 2.2.2. Representative network approach

AgMIP Wheat Phase 2b created a global network of 30 well-watered sites selected to represent major wheat systems and regional production areas (irrigated and high-rainfall wheat crops contribute ~70% of global production; see figure 2) (Asseng *et al* 2015a). 30 wheat models are configured for simulation of CT responses at each site, allowing robust ensemble projections and uncertainty analyses (Wallach *et al* 2015, 2016).

#### 2.2.2.1. Strengths and weaknesses of representative networks

The AgMIP Wheat Team network is distinct from C3MP's ad hoc network in that its design allows

multi-model assessment on major regional production systems that together generate the large majority of global wheat production (table 2). Simulated relative impacts are applied to recent FAO country production statistics associated with each simulated location to up-scale to global production impacts (Asseng *et al* 2015a, Liu *et al* 2016).

Even with 30 sites, the network is limited in its spatial coverage and individual sites may not reflect conditions in the broader production regions they represent. The network is concentrated in high-production zones and is likely to miss important responses in areas that were not simulated (AgMIP-Wheat Phase 3 will fill some of these gaps for water-stressed systems). As a simple metric of comprehensiveness of coverage, figure 3 shows how the rainfed and irrigated wheat networks from C3MP and AgMIP-Wheat Phase 2 cover wheat-growing climate conditions as compared with the global Monthly Irrigated and Rainfed Crop Area (MIRCA) year 2000 dataset (Portmann *et al* 2010), using AgMERRA climate data (Ruane *et al* 2015a) and growing seasons from the Global Gridded Crop Model Intercomparison (GGCMI; Elliott *et al* 2015). Both networks are most dense in climate zones that are prominent for wheat production; however the larger C3MP network also includes less common climates for rainfed wheat and samples more from the tails of the irrigated wheat distribution than does AgMIP-Wheat. By simulating more of the cool and wet tails it is likely that C3MP captures more farms that potentially benefit from increases in temperature or are less vulnerable to decreases in precipitation.

Regions with high levels of diversity are difficult to capture given limitations in representative site networks. Sentinel crop modeling sites are often calibrated with data from field experiment datasets designed to highlight potential genetic, fertilizer, water, or pest control treatments, and therefore may not be representative of prevailing agricultural systems within that production region. These site networks tend to be more useful when examining the percentage yield response to a given climate change; this metric has proven robust even in the face of persistent bias in mean regional yields (Challinor *et al* 2014b, Asseng *et al* 2015a).

### 2.3. Global approaches

Advances in high-performance computing have allowed crop models to enter a new phase of development that is nearly unconstrained by computational limitations. While IAMs are typically run on desktop computers or simple clusters, the 18 modeling groups participating in AgMIP's Global Gridded Crop Modeling Intercomparison (GGCMI; table 3) use parallel computing and advanced data processing pipelines to conduct protocol-based simulations on a 0.5° × 0.5° global grid (Rosenzweig *et al* 2014, Elliott *et al* 2015), with higher resolution gridded studies in the works. These outputs therefore form a desirable basis for more computationally-efficient IAM application through

Table 3. Overview of GGCM phases and model participation.

Phase (and key references)	Description [and # of models participating]
<b>Fast Track Rapid Assessment</b> (Rosenzweig et al 2014)	Conducted for AgMIP/ISIMIP using default versions of global gridded crop models, historical period and future scenarios from downscaled GCMs. Simulated maize, wheat, rice and soybean [seven GGCMs].
<b>Phase 1 Historical Intercomparison</b> (Elliott et al 2015, Müller et al 2017)	Default, harmonized, and no nitrogen stress versions of gridded crop models run over historical period using up to nine climate forcing datasets. Simulated maize, wheat, rice and soybean [15 GGCMs].
<b>Phase 2 CTWNA Sensitivity</b> [results submitted 2016–17]	Default simulations for historical period and sensitivity tests for [CO <sub>2</sub> ], temperature, water, nitrogen, and adaptation for all grid cells and crops. Simulated maize, spring wheat, winter wheat, rice and soybean [~12 GGCMs].
<b>Phase 3 Future Assessment</b> [planned for 2017–18]	Conducted for AgMIP/ISIMIP to assess future agricultural production under climate change scenarios [~12–20 GGCMs anticipated].

Note: ISIMIP = the Inter-Sectoral Impacts Model Intercomparison Project (Warszawski et al 2014).

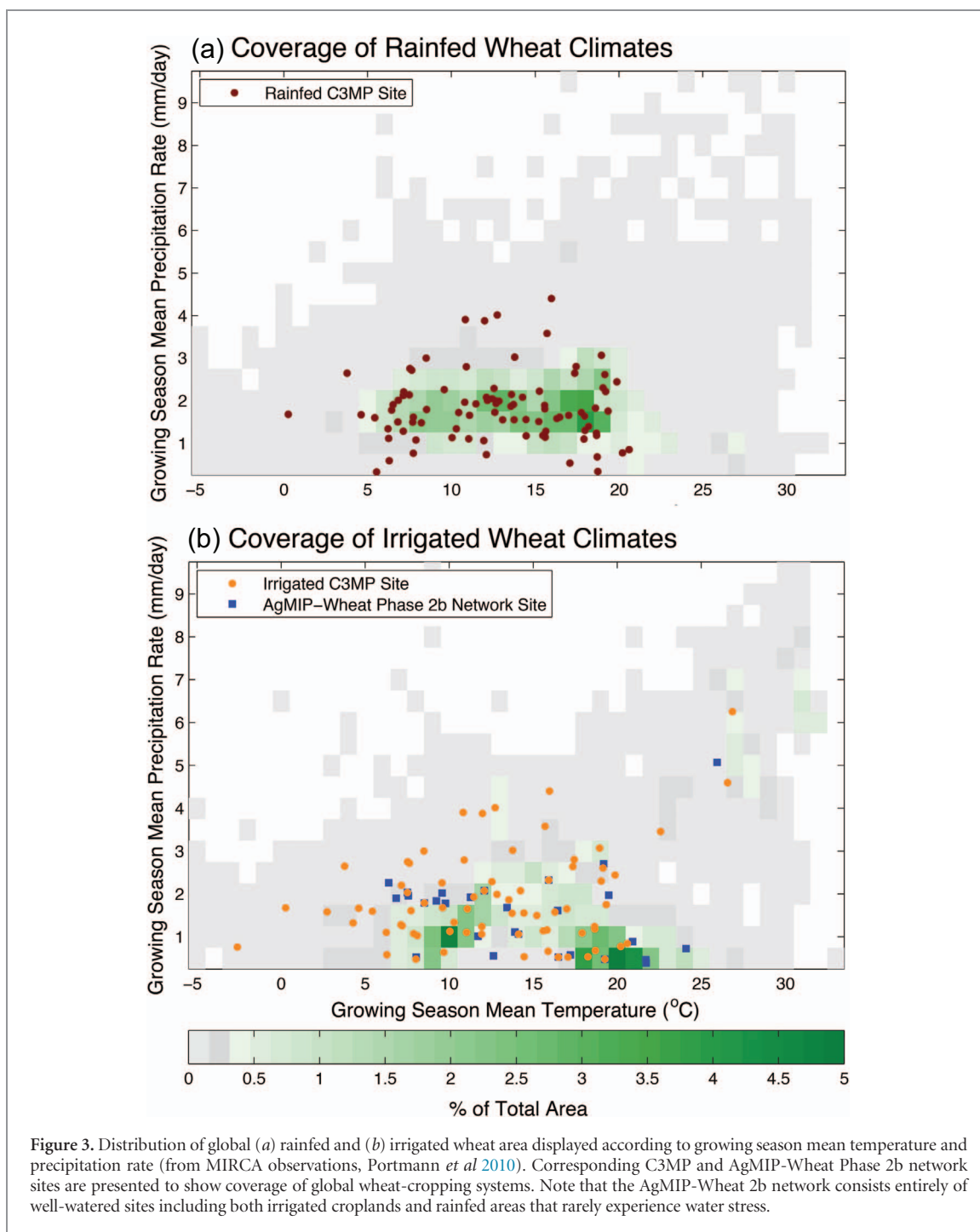
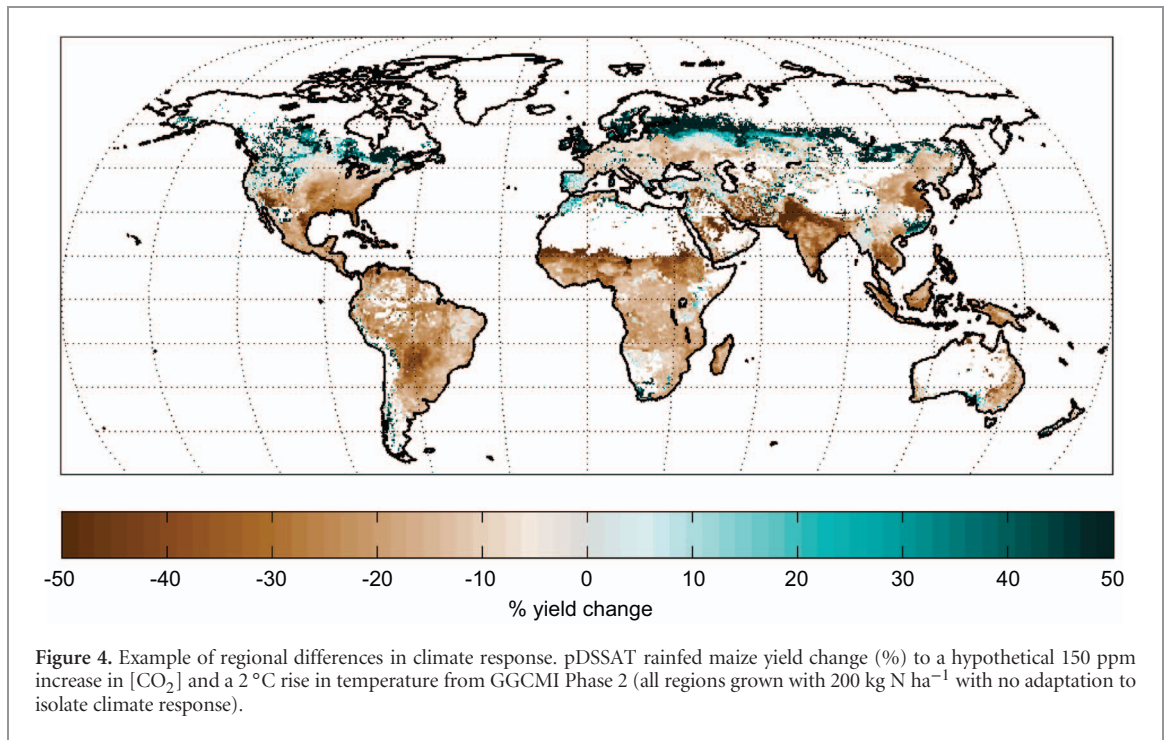


Figure 3. Distribution of global (a) rainfed and (b) irrigated wheat area displayed according to growing season mean temperature and precipitation rate (from MIRCA observations, Portmann et al 2010). Corresponding C3MP and AgMIP-Wheat Phase 2b network sites are presented to show coverage of global wheat-cropping systems. Note that the AgMIP-Wheat 2b network consists entirely of well-watered sites including both irrigated croplands and rainfed areas that rarely experience water stress.





emulators. GGCM Phase 2 performs a systematic analysis of CTWNA sensitivities for rainfed and irrigated maize, rice, wheat, and soybean with consistent climate information and harmonized planting dates. Adaptation is examined by shifting cultivars to maintain the growing period even as warmer temperatures accelerate phenologic development, thus offsetting some yield losses from climate change.

### 2.3.1. Strengths and weaknesses of global approaches

GGCM's fast-track results (table 3) provide biophysical impacts across emissions scenarios and five GCMs (Rosenzweig *et al* 2014), providing applications with ensemble mean impacts and uncertainty information from seven GGCMs for four crop species (maize, wheat, rice, and soybean) across the global grid (Nelson *et al* 2014, Wiebe *et al* 2015, Villoria *et al* 2016). It is difficult for crop model emulators to disentangle fundamental responses from these outputs, however, given the many types of changing and interacting climate conditions (e.g. mean temperatures and rainfall; sub-seasonal variations; extreme events). Emulation is also complicated by the inclusion of responsive adaptations allowing management to evolve with climate change in some participating models (Rosenzweig *et al* 2014, supplementary).

GGCM Phase 2 findings indicate considerable spatial variation in CTWNA response across different environments and farm systems, exemplified by the response of rainfed maize to higher  $[\text{CO}_2]$  and temperature in the parallel-DSSAT crop model (pDSSAT; Elliott *et al* 2014) (figure 4). These results provide a convenient basis for the construction of crop model emulators, and can also be connected to economic and/or resource availability drivers from IAMs to

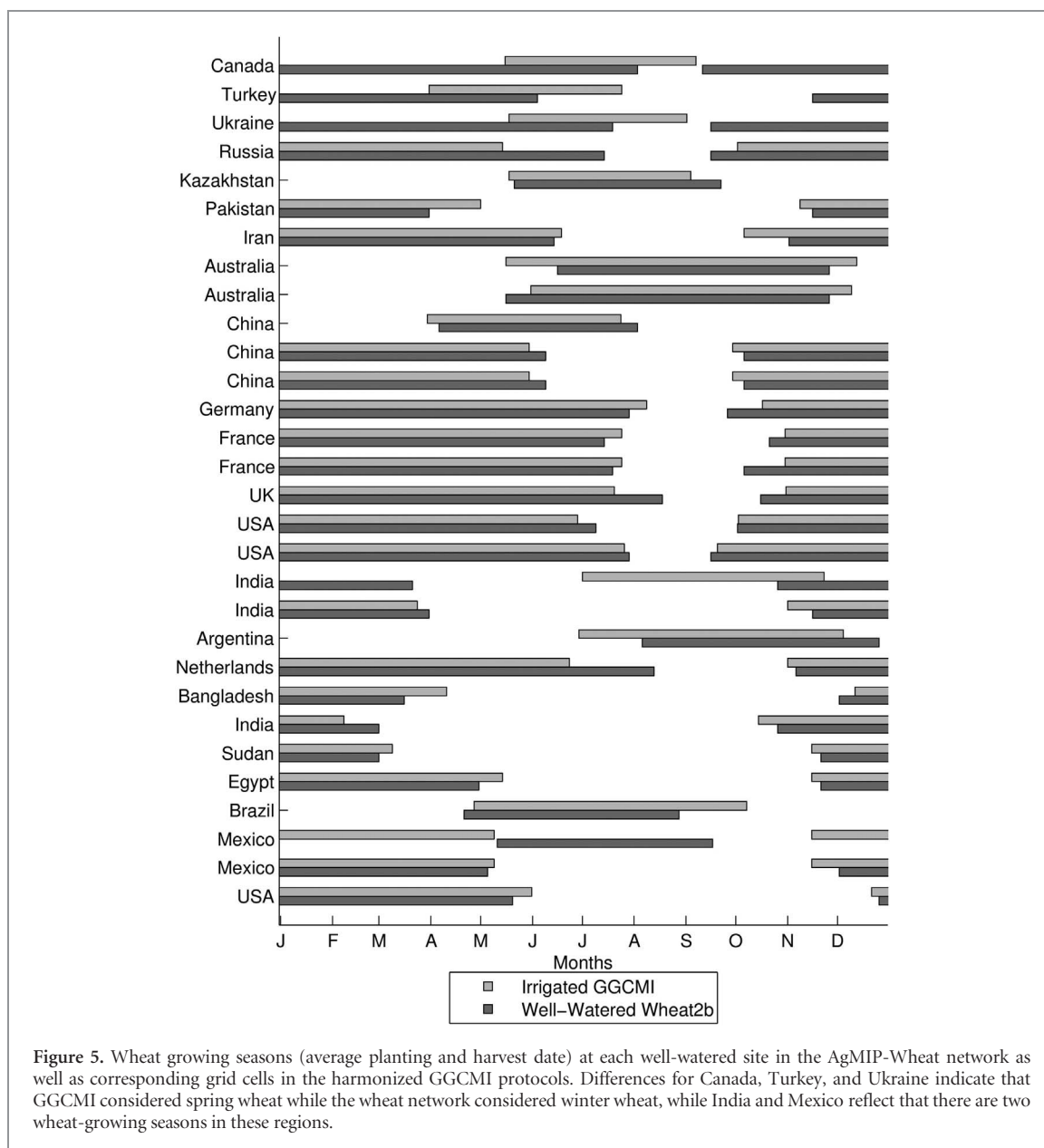
dynamically characterize the evolution of socioeconomic yield gap factors such as fertilizer use, irrigation, and adaptation potential.

In contrast to the site networks, GGCMs rely on gridded soil, genetic, management, and weather datasets designed to capture spatially-averaged conditions rather than conditions on a particular farm (Elliott *et al* 2015). While the  $0.5^\circ \times 0.5^\circ$  spatial resolution used within GGCM is finer than many GCMs, a grid cell on the equator represents  $>310\,000$  ha and thus poses a challenge for comprehensive farm system calibration.

GGCM results are often evaluated using regional yield and production reports, with trend adjustment recommended in recognition of technological development and processes that are not explicitly modeled such as pests, diseases, and widespread flooding (Müller *et al* 2017). Analogously, statistical crop response models are occasionally fitted to similar aggregate yield data that may reflect embedded abiotic factors (e.g. Lobell *et al* 2011). Bias-adjustment is recommended for GGCM application in IAMs, similar to common practices for climate model output applications (e.g. Wilby *et al* 2004, Ruane *et al* 2015b). Overall, GGCMs reflect that there are larger uncertainties in developing country and low-input farming systems, and stand to benefit from improved data collection and sharing in application regions (Kihara *et al* 2015, McDermid *et al* 2015b).

### 2.4. Emergent characteristics and opportunities from CTWNA simulations

AgMIP site, network, and gridded results demonstrate that multi-model ensembles outperform individual models when analyzed across multiple sites and evaluation variables (e.g. Asseng *et al* 2013,



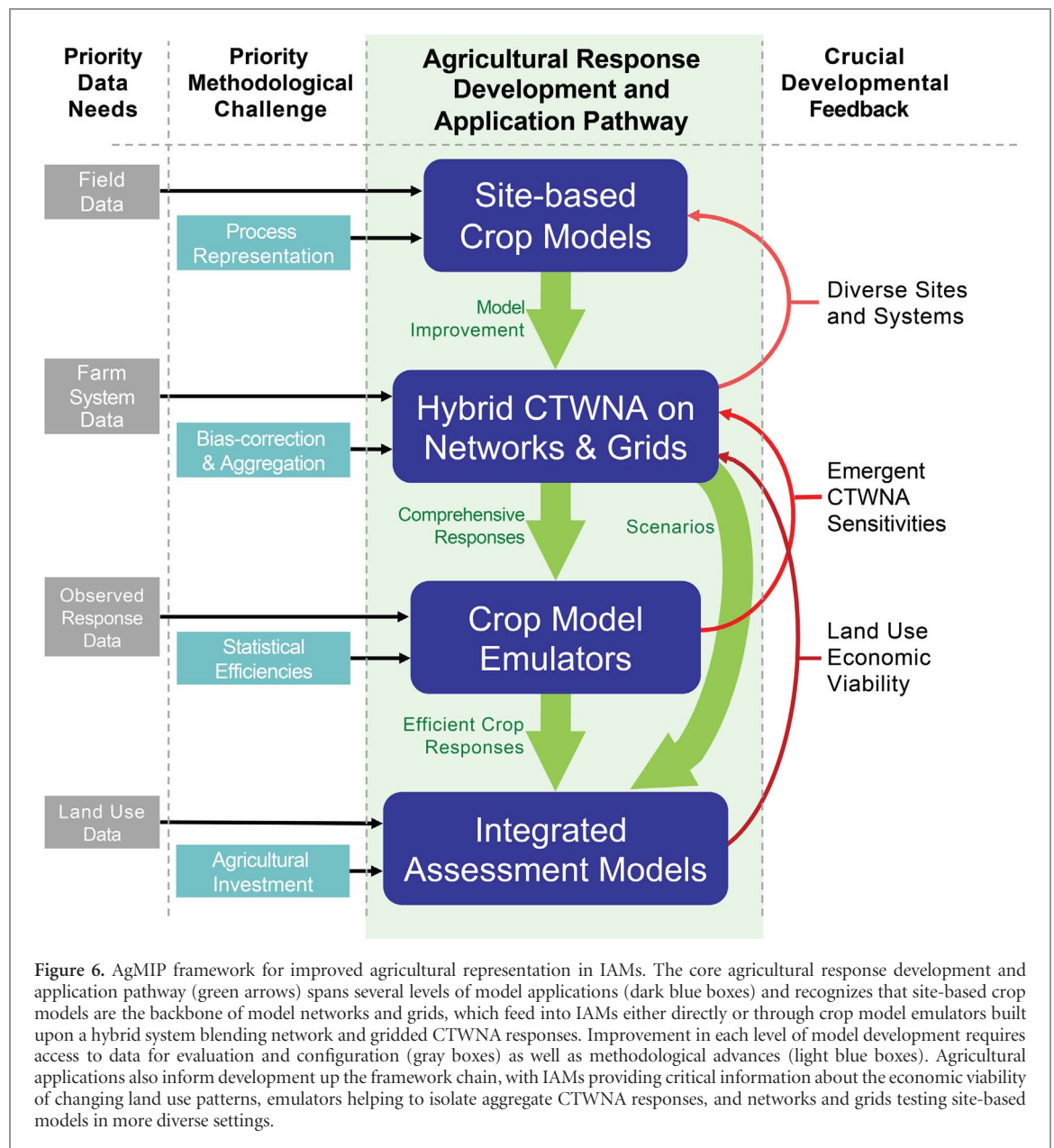
Rosenzweig *et al* 2014, Martre *et al* 2015, Li *et al* 2015, Bassu *et al* 2014, Wallach *et al* 2015, Ruane *et al* 2016, Fleisher *et al* 2016). Liu *et al* (2016) found relative agreement in wheat response to a 1 °C rise in global temperature, with multi-model ensembles in the well-watered AgMIP Wheat network, GGCMi's ISIMIP fast-track, and several statistical model approaches finding 4.1%–6.4% declines in global production.

Uncertainties in input data indicate that there is still room for harmonization that will improve consistency, as illustrated by a comparison of growing seasons at the well-watered AgMIP-Wheat network sites and corresponding GGCMi grid cells (figure 5). Uncertainty owing to model structure and parameters remains substantial, and differences in CTWNA responses by two modelers using the same DSSAT model within the MACSUR IRS and AgMIP-Wheat Phase 1 also highlights the potential role of modeler uncertainty stemming from assumptions and subjective decisions made in the absence of supporting data

(Pirttioja *et al* 2015, Confalonieri *et al* 2016). We therefore advise applications to recognize the uncertainty in model-based responses through the use of emulators derived from multiple models or an imposed error term scaled to model-based uncertainty. Evidence across AgMIP activities also recommends avoidance of universal yield functions in favor of yield response functions fitted to broad agro-ecological zones and farming systems (e.g. defined by fertilizer and irrigation inputs).

### 3. AgMIP framework for improved agricultural representation in IAMs

A cascading pathway of development underlies agricultural representation in IAMs, forming a framework that may be used to drive coordinated development of 'simulation levels', here defined as common communities of development including site-based crop models, network and gridded models, crop model emulators,



and eventual IAM applications (figure 6). Close collaboration and regular updates between site, network, and gridded crop modelers, emulation experts, and IAM groups are needed to keep agricultural impact applications on the cutting edge, to facilitate the use of multiple models, to incorporate understanding from multiple modeling groups, and to avoid the propagation of known biases.

Each simulation level in the AgMIP Framework benefits from improved data access and innovations in core methodologies. Investment in research and development is well served by matching the design, capabilities, and development priorities of models and tools at each level in figure 6. In particular, new biophysical process understanding is best developed within site-based models using field experiment data, particularly for under-sampled agro-ecological zones, crop species, and farming systems under various intensifications (Challinor *et al* 2015, Maiorano *et al* 2017).

Networks and gridded models gain from new datasets that allow extensive configuration for many sites and systems, and have tremendous potential to apply advanced bias-correction and aggregation approaches (Challinor *et al* 2014, van Bussel *et al* 2016, Hoffmann *et al* 2015, 2016, Zhao *et al* 2015, 2016). Crop model emulators are progressed with improved statistical efficiencies and the availability of observed agricultural response data for evaluating strengths and weaknesses. In addition to the potential benefit of adding improved crop model emulators, IAM simulations of long-term shifts in agricultural production are furthered by good data on current systems and advanced representation of the implications of agricultural investment and technological development.

The AgMIP framework for improved agricultural representation in IAMs is non-linear as lower simulation levels build upon advances higher up in the framework and high levels also receive critical

feedback from downstream simulation levels. Pathways of upstream improvements include that assessments of improved models on established grids and networks provide vital feedback for site-based model development on diverse sites. Likewise, emulators often spotlight key sensitivities and uncertainties that may spur further site-based model development and the creation of more representative networks. Network and gridded studies examine the biophysical viability of various simulated farm systems to determine land use pressures, but benefit tremendously by incorporating information on economic viability and resource constraints that IAMs can provide. It is also important to note that many of these simulation levels have extensive applications beyond agricultural representation in IAMs, and that the key bottleneck for one applications may differ from another's crucial development priority.

#### 4. Priority future development and applications

Analysis of the multi-model, multi-site climate sensitivity datasets reviewed in this study suggest that IAMs and other large-scale applications would be well served by the creation and systematic development of a hybrid CTWNA response system that blends the strengths of network and gridded approaches (as noted in figure 6). This hybrid response system would be rooted in (1) detailed process understanding across a representative network of well-calibrated field sites (ideally using field data from prevailing management systems) combined with (2) comprehensive CTWNA coverage from gridded models. Baseline responses generated by these gridded models could initially be compared against the corresponding representative network simulations to assess methodological uncertainty and calculate bias-correction factors. Bias-corrected gridded results could then provide an information basis for crop model emulators and IAM applications, characterizing different farming systems using nitrogen and water components of the CTWNA analysis.

Table 1 highlights that progress toward the creation of this hybrid response system is most advanced for wheat, given the AgMIP-Wheat Phase 2b representative network and spring and winter wheat simulations within GGCM Phase 2. In contrast, soybean is simulated in GGCM but has not yet been the focus of site- or network-based CTWNA analysis, and a number of other important commodities merit inclusion. Coordinated and systematic development of the hybrid response system would foster rapid iterative improvements, as research groups improve the hybrid framework by contributing new process understanding, field sites, model runs, regional configuration information, or statistical approaches. An expanded representative network of models and a fully configured high-resolution gridded (or geo-referenced polygon) model will eventually be interchangeable; however this

hybrid response system provides current state-of-the-art responses and a practical roadmap for applications.

Coordination across AgMIP activities supports the development of linked global and regional assessments to address agricultural sector challenges and food security (Rosenzweig *et al* 2016). Inclusion of IAMs would bring these to a new level, although it is critical that these account for lingering model uncertainty and data gaps even as these are addressed through the coordinated development of agricultural response in linked models.

#### Acknowledgements

We are grateful for the tremendous effort and contributions provided by all modelers and leaders of all projects listed in table 1. This letter was motivated by discussions with members of the Program on Integrated Assessment Model Development, Diagnostics, and Intercomparisons (PIAMDDI) and the Pacific Northwest National Laboratories, and benefitted from support by the National Aeronautics and Space Agency Science Mission Directorate (WBS 281945.02.03.06.79), the United Kingdom Department for International Development (GB-1-202108), and the Department of Energy (DE-AC5-76RL01830 and DE-SC0005171) as well as the Joint Research Centre of the European Commission. We appreciate the assistance of Meridel Phillips and Greg Reppucci in figure creation.

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