Future Risks of Pest Species under Changing Climatic Conditions

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Abstract

Most agricultural pests are poikilothermic species expected to respond to climate change. Currently, they are a tremendous burden because of the high losses they inflict on crops and livestock. Smallholder farmers in developing countries of Africa are likely to suffer more under these changes than farmers in the developed world because more severe climatic changes are projected in these areas. African countries further have a lower ability to cope with impacts of climate change through the lack of suitable adapted management strategies and financial constraints. In this study we are predicting current and future habitat suitability under changing climatic conditions for Tuta absoluta, Ceratitis cosyra, and Bactrocera invadens, three important insect pests that are common across some parts of Africa and responsible for immense agricultural losses. We use presence records from different sources and bioclimatic variables to predict their habitat suitability using the maximum entropy modelling approach. We find that habitat suitability for B. invadens, C. cosyra and T. absoluta is partially increasing across the continent, especially in those areas already overlapping with or close to most suitable sites under current climate conditions. Assuming a habitat suitability at three different threshold levels we assessed where each species is likely to be present under future climatic conditions and if this is likely to have an impact on productive agricultural areas. Our results can be used by African policy makers, extensionists and farmers for agricultural adaptation measures to cope with the impacts of climate change.

Introduction

The combination of about 795 Million people suffering from undernourishment [1] and an expected population increase from 7 up to 9 billion by 2050 is projected to lead to an increase in the need for food between 70–100% compared to 2010 [2]. At the same time, the United Nations have agreed to aim at reaching the Sustainable Development Goals (SDGs) that include the eradication of hunger and the sustainable use of terrestrial ecosystems. Yet, it is very likely that competition for the limited arable land will increase under these, at least...
partially, conflicting goals and that efficient and sustainable use of already cultivated land will become even more important in the future than it is already today.

The global scale of climate change (CC) and the necessity for sustainable poverty mitigation strategies increases the need to quantify CC impacts. Especially the effects on agricultural production are of major importance for the local populations, in particular farmers, national governments, regional bodies like the African Union (AU), international development partners and other stakeholders [3]. Such data can increase awareness and help to develop coping strategies for the vulnerable, as well as enable them to adapt better to a changing environment. It would also allow national governments to design and implement changes in policies, necessary for mitigating the effects of climate change [4].

In Africa, agriculture is the source of income for many families and represents over two thirds of livelihoods of the poor. About 65% of full-time employment is in the agricultural sector and over half of the total export earnings derive from agricultural goods [5]. Large losses of agricultural production can be attributed to pests. In Africa alone 12.8 billion US$ were estimated to be lost to pathogens, insects and weeds between 1988 and 1990. Insects are the economically most relevant pest group and the cause for about 1/3 of the actual crop production equal to 4.4 billion US$ being lost [6]. Many people in sub-Saharan Africa (SSA) heavily rely on natural resources and have a relatively low tolerance towards climatic and economic stress because of high poverty levels and lack of alternative sources of income [4]. The often limited capacity to adapt to changes through a lack of knowledge and education, further increases the vulnerability of Africa’s poor to CC impacts [7]. Apart from the more general negative effects that CC is expected to have on agricultural production in many developing countries, it is also likely to have a profound impact on the abundance and distribution of many pest species.

CC impacts are expected for the whole planet, but farmers in the developed world are likely to be better prepared to deal with potentially increasing numbers of pests or invasive species than those in less developed countries. Main reasons for this include greater financial means of farmers in the developed world to utilize different pest control strategies, be they biological, synthetic pesticides or genetically modified crops, interventions that are most often economically not feasible for most of the producers in the developing world. By consequence, this then results in greater food insecurity in these regions.

Agricultural productivity strongly depends on continued innovations to control pests as they develop resistances to different control measures, such as synthetic pesticides, or disperse to new regions [2]. While most studies estimate increasing numbers and distribution for many pest species, responses of individual species may vary depending on, among others, the bioclimatic conditions under different CC scenarios. In this study we use species distribution modelling to evaluate the possible extent and change of the habitat suitability across Africa for three important pest species, *Tuta absoluta, Ceratitis cosyra and Bactrocera invadens*, under future CC.

*Tuta absoluta* (Meyrick) (Lepidoptera: Gelechiidae), also called the tomato leaf miner, is a key threat to European and African tomato production. The pest originates from South America and is spreading rapidly over southern Europe into northern Africa since its first detection in Spain in 2006. Its high reproductive capacity and rapid development of resistance to many different insecticides make conventional chemical control very challenging. Consequently yield losses of 80–100% have been reported [8]. *Ceratitis cosyra* (Walker) (Diptera: Tephritidae), or mango fruit fly, is a serious pest in smallholder and commercial mango plantations across SSA. It is native to afrotropical regions and is commonly intercepted in Europe as larvae in infested mangoes [9]. *Bactrocera invadens* (Drew, Tsura and White, 2005) (Diptera: Tephritidae) is a member of the oriental fruit fly species complex, possibly of Sri Lankan origin, and since 2003 has spread across East and West Africa. It has a very broad host range and feeds on a wide...
variety of unrelated wild and cultivated crops. Due to its highly destructive and invasive potential, *B. invadens* has become economically the most important fruit fly in Africa [10].

Many studies investigating the impact of CC on pest distribution show increasing densities or an expansion of the geographical range of pests [11–14]. Significant research focusing on the possible impacts of CC on insects has already been carried out in temperate areas of the world [15–17]. It was reported that warmer winters might advance the survival rate of insects and permit a faster population revival that will consequently build-up in spring [17]. Additionally, an increase in the length of the cultivating period is projected, permitting multivoltine species to generate a higher number of generations per year and therefore upsurge invasions by alien species [15,18].

In comparison to temperate areas, tropical countries of Africa are highly disposed to insect pest problems and outbreaks due to their year-round favorable environments for insects’ population growth and host plants availability [19]. Further studies revealed that warming in tropical areas, although tiny in scale, is expected to yield harmful consequences because tropical insects are very sensitive to small changes on the magnitude of climatic variables, such as temperature [20,21].

Species distribution modelling with presence-only data can be used to model the current habitat suitability of a species and project future suitabilities under changed climatic conditions. Models of this type have already been developed for several pest species in Asia [22–24], North America [25,26], several European countries [27], as well as for the global distribution of pests [28–30]. Fewer modelling approaches have been carried out with a focus on Africa [31,32]. Considering the importance of agriculture for the continent, more studies assessing pest species dispersal are needed to estimate potential future yield losses as a result of CC.

Climate change is likely to have negative impacts on food security and livelihoods of farmers in Africa through change of the number of generations and distributions of pest species [13,33,34]. Despite this, there are only few studies providing specific information on which species are likely to affect which regions under CC in Africa. This information, however, is vital for farmers, if they want to adapt to the impacts of CC.

**Methods**

**Pest species and presence records**

Presence records of the three important insect pest species were used from different sources including the Global Biodiversity Information Facility (GBIF, www.gbif.org) and published studies (Table 1, Fig 1).

Presence data were randomly split in two sets for model training (70%) and testing (30%). A number of presence records had to be excluded because of missing environmental information or other issues with the geographical coordinates (e.g. coordinates pointed towards locations in the ocean). All used presence records are provided in S1 File.

**Environmental variables**

Presence records (Fig 1) were used in combination with bioclimatic variables downloaded from the WorldClim database ([44] accessed through [45]) to assess current and future habitat suitability of the pest species using maximum entropy modelling. Current climatic data correspond to interpolations of observed data from 1950 to 2000, while future climatic conditions represent downscaled modelled data averaged for the years 2041 to 2060 according to the IPCC5 (CMIP5). Three different global circulation models (GCMs) and four representative concentration pathways (RCP) were used in a 2.5 arc minute (4.6 km) resolution.
Not all of the selected GCMs provide data for all RCPs (Table 2). The GCMs were selected based on their ability to reflect the dynamics of the West African monsoon [46,47] and their representation of modelled precipitation and temperature values from different CMIP5 models [48]. While the GFDL model represents colder and wetter values, the HadGEM model values are warmer and drier and the MPI-ESM values are close to the multi-model mean.

Habitat suitability was modelled using bioclimatic variables under each GCM and RCP. Bioclimatic variables are calculated based on monthly values of temperature and rainfall but are likely to be biologically more meaningful than simple average values since they represent both annual trends but also seasonality and extreme conditions (Table 3).

Mean values of future habitat suitability were calculated over all modelled habitat suitability datasets. They were calculated first over all models for all RCPs before they were calculated over all models. Therefore the lack of data for two RCPs for two GCMs only results in a lower confidence level for the results of the two higher RCPs. The overall results, however, are not skewed towards higher or lower emission scenarios.

Modelling approach and evaluation

The machine learning approach Maxent (www.cs.princeton.edu/~schapire/maxent/) was used to assess habitat suitability based on maximum entropy. Maxent has been shown to perform particularly well for modelling presence-only data [49]. Maxent is used to predict the environmental suitability for the species as a function of the given environmental variables. Hereby the distribution probability is estimated by finding the distribution of maximum entropy that satisfies a set of constraints from environmental variables. The results serve as an approximation of

Table 1. Species names, source of presence records, date of data access, total number of presence records found for Africa in the original source and number of presence records used for training and testing.

<table>
<thead>
<tr>
<th>Species Name</th>
<th>Source</th>
<th>Access Date</th>
<th>No. of presence records</th>
<th>from source</th>
<th>used for training</th>
<th>used for testing</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tuta absoluta</td>
<td>[35–40]</td>
<td>03/02/15</td>
<td>542</td>
<td>141</td>
<td>60</td>
<td></td>
</tr>
<tr>
<td>Ceratitis cosyra</td>
<td>[35,41,42]</td>
<td>13/02/15</td>
<td>595</td>
<td>175</td>
<td>75</td>
<td></td>
</tr>
<tr>
<td>Bactrocera invadens</td>
<td>[10,31,43]</td>
<td>05/01/15</td>
<td>226</td>
<td>118</td>
<td>50</td>
<td></td>
</tr>
</tbody>
</table>

Fig 1. Presence records for 3 important pest species accessed through GBIF and other literature (see Table 1 for an overview).
a species’ ecological niche under the studied environmental conditions [50]. Maxent belongs to the family of models relying solely on presence records of the investigated species.

Environmental variables were reduced to only the three most important variables, with the highest variable contributions calculated according to a first model run based on all variables.

A bias file was produced with the package kernSmooth in R [51] to correct for uneven distributions of sampling efforts across the study area. For this purpose coordinates from presence 18,108,111 records of all animals from GBIF were used to create a kernel density estimate map across Africa with a bandwidth of 5 in each direction. Since the sample size is very large, we chose a relatively small bandwidth which is still large enough to result in values greater than 0 in each pixel cell.

The performance of the model is characterized by the area under the curve (AUC). The AUC is obtained by the threshold independent receiver operating characteristic (ROC) analysis [50]. In the process of modelling, 70% of occurrence localities were randomly selected as training data, while the remaining 30% served for testing the resulting models. The ROC method is based on the AUC when model sensitivity is plotted against 1 minus model specificity. This method has been shown to be effective in evaluating model performance and being independent of prevalence compared to the more commonly used kappa statistic [52–54].

Table 2. Overview of global circulation models (GCMs) and representative concentration pathways (RCPs) that were available and accessed via the database WorldCim ([44] accessed through [45]).

<table>
<thead>
<tr>
<th>Abbrev.</th>
<th>Bioclimatic Variable Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>BIO1</td>
<td>Annual Mean Temperature</td>
</tr>
<tr>
<td>BIO2</td>
<td>Mean Diurnal Range (Mean of monthly (max temp—min temp))</td>
</tr>
<tr>
<td>BIO3</td>
<td>Isothermality (BIO2/BIO7) (* 100)</td>
</tr>
<tr>
<td>BIO4</td>
<td>Temperature Seasonality (standard deviation *100)</td>
</tr>
<tr>
<td>BIO5</td>
<td>Max Temperature of Warmest Month</td>
</tr>
<tr>
<td>BIO6</td>
<td>Min Temperature of Coldest Month</td>
</tr>
<tr>
<td>BIO7</td>
<td>Temperature Annual Range (BIO5-BIO6)</td>
</tr>
<tr>
<td>BIO8</td>
<td>Mean Temperature of Wettest Quarter</td>
</tr>
<tr>
<td>BIO9</td>
<td>Mean Temperature of Driest Quarter</td>
</tr>
<tr>
<td>BIO10</td>
<td>Mean Temperature of Warmest Quarter</td>
</tr>
<tr>
<td>BIO11</td>
<td>Mean Temperature of Coldest Quarter</td>
</tr>
<tr>
<td>BIO12</td>
<td>Annual Precipitation</td>
</tr>
<tr>
<td>BIO13</td>
<td>Precipitation of Wettest Month</td>
</tr>
<tr>
<td>BIO14</td>
<td>Precipitation of Driest Month</td>
</tr>
<tr>
<td>BIO15</td>
<td>Precipitation Seasonality (Coefficient of Variation)</td>
</tr>
<tr>
<td>BIO16</td>
<td>Precipitation of Wettest Quarter</td>
</tr>
<tr>
<td>BIO17</td>
<td>Precipitation of Driest Quarter</td>
</tr>
<tr>
<td>BIO18</td>
<td>Precipitation of Warmest Quarter</td>
</tr>
<tr>
<td>BIO19</td>
<td>Precipitation of Coldest Quarter</td>
</tr>
</tbody>
</table>

doi:10.1371/journal.pone.0153237.t003
The output of the model represents an area with conditions comparable to those in the species’ known occurrence range, whereas the values between 0 and 1 indicate regions with no or most suitable habitat conditions, respectively.

The response of the species distribution model to specific environmental variables was investigated through the permutation importance and response curves of each bioclimatic variable. The permutation importance is calculated based on the drop or increase of the AUC, when the respective environmental variable is altered.

Based on a threshold value, habitat suitability can be converted from probability maps to species distribution maps. In this study we used three different threshold values to create distribution maps and overlaid them with agricultural production intensity with current and future species presence polygons. We chose to use three commonly used threshold levels to display the variability in distribution maps dependent on the selected threshold level (Table 4). The agricultural intensity map is an extract of the data on global crop land published by Ramanikutty et al. [55] and represents the fraction of area being under use as cropland for each raster cell at a resolution of 5 min (10 km).

All maps are displayed using a simple Plate Carrée (WGS 84) projection. This projection is an equidistant projection, which was chosen as it is appropriate for large-scale studies to balance area distortion and shape.

Results

Statistical model evaluation

AUC values for training data of all species are >0.85 and AUC values for test data of all species are >0.8 (Table 5). This demonstrates that all models show a good predictive performance.

Habitat suitability under current and future climatic conditions

Current habitat suitability shows which area each species is likely to inhabit under current climatic conditions (Fig 2).

*Bactrocera invadens* (Fig 2A) and *C. cosyra* (Fig 2D) show high values of habitat suitability scattered across SSA. However, while *C. cosyra* seems to be able to inhabit also Central Africa, *B. invadens* is more restricted to the coastal areas. Suitable habitats for *T. absoluta* (Fig 2G) are located almost exclusively in North Africa (Algeria, Libya and Egypt), across the Sahelzone.

Table 4. Thresholds used to estimate species distribution maps for all three species. The "Balance" threshold minimizes 6 * training omission rate + .04 * cumulative threshold + 1.6 * fractional predicted area.

<table>
<thead>
<tr>
<th></th>
<th><em>B. invadens</em></th>
<th><em>C. cosyra</em></th>
<th><em>T. absoluta</em></th>
</tr>
</thead>
<tbody>
<tr>
<td>Equal training sensitivity and specificity</td>
<td>0.266</td>
<td>0.378</td>
<td>0.379</td>
</tr>
<tr>
<td>Maximum training sensitivity plus specificity</td>
<td>0.196</td>
<td>0.187</td>
<td>0.463</td>
</tr>
<tr>
<td>Balance training omission, predicted area and threshold value</td>
<td>0.063</td>
<td>0.095</td>
<td>0.088</td>
</tr>
</tbody>
</table>

Table 5. Area under curve (AUC) values of training and test data as an evaluation parameter of model performance for all investigated species.

<table>
<thead>
<tr>
<th>Species</th>
<th>AUC training data</th>
<th>AUC test data</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Bactrocera invadens</em></td>
<td>0.929</td>
<td>0.917</td>
</tr>
<tr>
<td><em>Ceratitis cosyra</em></td>
<td>0.874</td>
<td>0.861</td>
</tr>
<tr>
<td><em>Tuta absoluta</em></td>
<td>0.938</td>
<td>0.868</td>
</tr>
</tbody>
</table>
Fig 2. Habitat suitability under current and future climatic conditions as well as change of habitat suitability of *Bactrocera invadens* (a-c), *Ceratitis cosyra* (d-f), and *Tuta absoluta* (g-i) modelled as logistic outputs of Maxent.

doi:10.1371/journal.pone.0153237.g002
and on the Arabian Peninsula. Especially in Egypt, tomato, the host plant of *T. absoluta*, is an important cash crop. Egypt belongs to the top tomato producing countries worldwide (http://faostat.fao.org/). Therefore, we expect large negative economic impacts of *T. absoluta* for tomato producers under future CC.

The shown values represent mean values over model outputs with all available 10 bioclimatic variable datasets (3 GCMs and 4 RCPs, Table 2) using the three most important variables. Mean values over each RCP can be found for each species in S1–S3 Figs. Future habitat suitability of the three species appears to be overall similar to the current suitability (Fig 2). Although species react differently to projected climatic changes, in this study all three species showed, at least for parts of Africa, an increase in habitat suitability. Across large parts of the continent they show an either unchanged or even slightly increasing number of suitable habitat sites. Within the maps highlighting the change of habitat suitability an increase of suitability is indicated by the colours yellow to red, while the suitability of areas with the colours light-green to dark-green is likely to decrease.

### Environmental variable importance and impact

Permutation importance varies between species and for all environmental variables (Table 6). Response curves of the models for the three most important environmental variables of each species are shown in S4 Fig. For *B. invadens* temperature seasonality (BIO4), temperature annual range (BIO7) and precipitation of the driest quarter (BIO17) are the most important variables. The species is preferring for all of these variables rather low values. Especially temperature seasonality and annual range are unlikely to change dramatically in the future, which

<table>
<thead>
<tr>
<th>Variable Abbrev.</th>
<th>Permutation importance (%)</th>
<th><em>Bactrocera invadens</em></th>
<th><em>Ceratitis cosyra</em></th>
<th><em>Tuta absoluta</em></th>
</tr>
</thead>
<tbody>
<tr>
<td>BIO1</td>
<td>0.5</td>
<td>0</td>
<td>2.3</td>
<td></td>
</tr>
<tr>
<td>BIO2</td>
<td>1.6</td>
<td>0.7</td>
<td>4.9</td>
<td></td>
</tr>
<tr>
<td>BIO3</td>
<td>0.3</td>
<td>15</td>
<td>1.6</td>
<td></td>
</tr>
<tr>
<td>BIO4</td>
<td>14/63.3</td>
<td>6.4</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>BIO5</td>
<td>0</td>
<td>0.4</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>BIO6</td>
<td>3.4</td>
<td>0</td>
<td>28.9/31.2</td>
<td></td>
</tr>
<tr>
<td>BIO7</td>
<td>15.3/23.4</td>
<td>5.5</td>
<td>1.4</td>
<td></td>
</tr>
<tr>
<td>BIO8</td>
<td>4.4</td>
<td>8.1</td>
<td>8/20.8</td>
<td></td>
</tr>
<tr>
<td>BIO9</td>
<td>0.4</td>
<td>0.3</td>
<td>3.8</td>
<td></td>
</tr>
<tr>
<td>BIO10</td>
<td>0.5</td>
<td>1.4</td>
<td>1.2</td>
<td></td>
</tr>
<tr>
<td>BIO11</td>
<td>1.3</td>
<td>11.1/28.5</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>BIO12</td>
<td>13.5</td>
<td>8.8</td>
<td>40.5/48</td>
<td></td>
</tr>
<tr>
<td>BIO13</td>
<td>5.2</td>
<td>5.3</td>
<td>6</td>
<td></td>
</tr>
<tr>
<td>BIO14</td>
<td>0.5</td>
<td>8.9/13.2</td>
<td>1.2</td>
<td></td>
</tr>
<tr>
<td>BIO15</td>
<td>5.1</td>
<td>5.9</td>
<td>1.3</td>
<td></td>
</tr>
<tr>
<td>BIO16</td>
<td>4.2</td>
<td>10.2/58.3</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>BIO17</td>
<td>20.9/13.4</td>
<td>4.8</td>
<td>0.6</td>
<td></td>
</tr>
<tr>
<td>BIO18</td>
<td>3.5</td>
<td>2.2</td>
<td>3.8</td>
<td></td>
</tr>
<tr>
<td>BIO19</td>
<td>4.8</td>
<td>5</td>
<td>4.5</td>
<td></td>
</tr>
</tbody>
</table>

doi:10.1371/journal.pone.0153237.t006
is probably the reason why *B. invadens* does not respond strongly to changing climatic conditions.

The *C. cosyra* modelled suitability depends mainly on the mean temperature of the coldest quarter (BIO11), precipitation of the driest month (BIO14) and the precipitation of the wettest quarter (BIO16). Overall it prefers medium to high precipitation rates and temperatures >10°C throughout the year. Climate change projections indicate that in those areas of Africa, where we find increasing habitat suitability especially temperatures and precipitation are likely to increase under future CC. For *T. absoluta* the minimum temperature of the coldest month (BIO6), the mean temperature of the wettest quarter (BIO8) and annual precipitation (BIO12) are the most important parameters for suitable habitat conditions. The model indicates that it prefers higher temperatures and lower precipitation rates. Under CC more areas are projected to fulfill these requirements, especially in northern Africa.

**Pest impact on agricultural areas**

Assuming a threshold habitat suitability at different level shows that although habitat suitability in some areas might be increasing, under the suitability threshold equal training sensitivity and specificity the species is still not predicted to be present. However, under a threshold value of maximum training sensitivity plus specificity as well as balancing training omission, predicted area and threshold value much larger proportions of Africa are predicted to be inhabited by the species (Fig 3).

Comparing the areas where the species is predicted to be present under current and future climate shows that *B. invadens* and *T. absoluta* are unlikely to shift their habitat at all. For all three applied habitat suitability thresholds species distribution does not change under current and future climatic conditions even though habitat suitability generally is increasing over large parts of the continent (compare with Fig 2). For *C. cosyra* we find a decreasing number of areas in southern and Central Africa being inhabited under a higher threshold (Fig 3D). For the two higher thresholds mostly the same areas are being predicted as suitable under future as under current climatic conditions.

Comparing current and future distributions of the studied species with agricultural crop intensity indicates that those areas with high agricultural production are not under a higher threat under future CC than under current conditions. Habitat extent of all tested species is likely to remain constant, shift to less productive sites or decrease.

Overlaying the habitat of all three species for all three thresholds under current and future climate shows that especially for lower threshold levels CC impact seems to be deniable (Fig 4). Under these levels almost all areas are already under current conditions affected by at least one of the pest species (Fig 4E and 4F). For higher threshold levels, CC seems to have a rather positive impact since the distribution of *C. cosyra* is slightly decreasing, while the distributions of the two other species remain largely constant (Fig 4A and 4B).

**Discussion**

In this study we assessed the impact of CC on three important agricultural pests of different crops in Africa. We used environmental variables to assess CC effects in combination with different presence records obtained from multiple sources. Few other studies have investigated the distribution and potential for dispersal of pests in Africa [31,32]. Here we modelled the future distribution of three important pests in Africa under current and future climatic conditions. Area under the curve statistics showed high values for all species confirming a good model performance.
Climate Change impacts on agriculture poses a major threat to agricultural productivity in Africa [34,56,57]. Even without consideration of the impact of pests under CC decreasing yields for major crops between 5% for maize and 17% for wheat have been projected by 2050 across the entire African continent [56]. At the same time many studies indicate that productivity could also benefit under CC if suitable adaptation measures are implemented [58]. Adaptation through better agricultural management as well as decision making under consideration of CC risks, as suggested for example by Vermeulen et al. [59], is strongly influenced by the availability of information on CC impacts, such as the future distribution of important pests.

Our results show that species presence or absence depends strongly on the choice of a habitat suitability threshold. While in a certain area a species might not be predicted to be present under a higher threshold level, it might yet be under a lower threshold level. Using variable threshold levels, as we did in this study, shows this uncertainty, which can also be translated to risk levels of pest species invasion. Such information are useful for farmers, NGOs and policy makers as they give them a prioritisation list on which pest species to focus on first and which species are of lesser concern to them. Furthermore, it provides a guideline which crops are recommendable to be planted and which should be avoided, if risks are to be minimized.

Presence-only data models stand beside those based on presence and absence data usually obtained through systematic sampling, e.g. generalized linear or additive models. However, presence data are often easier to obtain than verified absence data for example from databases and museum collections. Therefore, presence-only models, such as Maxent, GARP (Genetic Algorithm for Rule-Set Production) or ENFA (Ecological Niche Factor Analysis) are usually used for predictions based on presence-only data. GARP models are based on the integration or rejection of rules that are being tested to improve or decrease the predictive performance of the model. ENFA based predictions are calculated from uncorrelated factors explaining the differences between the whole study area and the area inhabited by the species. Maxent, on the other hand, assumes that a species distribution would follow a maximum entropy without any environmental constraints. The model predicts habitat suitability by fitting a probability distribution for the occurrence of the species across the whole area. Based on data from the different environmental variables different constraints are being formulated and considered in the model. According to to Elith et al. [49] Maxent performs relatively well compared to other presence-only models. However, Maxent also seems to suffer from a higher tendency of overfitting at low threshold levels than e.g. GARP models [60]. For this reason we used three different threshold levels to display species distribution maps.

Comparing our results with other studies shows some discrepancies in the predictions of habitat suitability for individual species under current and projected climate. For instance Ton-nang et al. [30] modelled worldwide habitat suitability for *T. absoluta* using CLIMEX and found much larger areas of high suitability, especially across Central, eastern and southern Africa than in our study. De Meyer et al. [31] modelled habitat suitability for *B. invadens* in Asia, Africa and worldwide using two different approaches, i.e. GARP and Maxent with presence records from India, Sri Lanka and Bhutan. The Maxent modelling approach showed a much smaller area than the GARP approach. Suitable areas as predicted by De Meyer et al. [31] show some overlaps with highly suitable areas identified in our study. Yet for Central Africa we predicted lower suitability values than De Meyer et al. [31].

The discrepancies between the findings in our study and other publications may be the result from the level of uncertainty of methodologies currently used for species distribution.
modelling as well as different modelling input data. For example both, Tonnang et al. [30] and De Meyer et al [31] did not correct for sampling bias.

Another important fact is that the two species, *B. invadens* and *T. absoluta*, for which our modelling outcomes differ from those in other studies, are alien invasive species (AIS) in Africa. In Tonnang et al. [30] and De Meyer et al [31], both authors reported to have used records from the species native areas to develop their models; the obtained parameters were then projected to Africa for estimating areas of habitat suitability of the species. This approach differs from our modelling method, where we used only presence records from Africa. Being alien to Africa, the two species are likely to be still spreading and adapting to new environmental conditions to establish their final realized niches. When a species invades a new region its dispersal into new areas depends on environmental conditions and its ability to adapt can take a considerable amount of time. Hence, for model development using data of an AIS’ home might not be ideal for correctly predicting habitat suitability in a newly colonised region/area. Alternatively, native presence records of a species can be used to predict habitat suitability in different regions. Yet for species with high tolerance towards different environmental conditions this may also lead to lower predicted habitat suitability, especially in areas with environmental conditions not present in the species native area. If a species has for example a high tolerance to lower temperatures but presence points of areas with low temperatures were not included in the model because the species has not yet invaded such an area, predicted habitat suitability solely based on native records could be lower than the true suitability.

We used bioclimatic variables to characterize environmental conditions. However, other important environmental variables, such as soil properties, land cover and agricultural management interventions (e.g. use of pesticides or fertilizers) can influence species distribution but were not considered in this study. Including such variables has been described as a key challenge for modelling approaches [61]. While we agree that these variables are key to assess the final distribution of species, in this study, however, we emphasize on the necessity to assess only CC impact on three important agricultural pests in Africa. Since their distribution very much depends on which crops are grown where and how they are managed, projections under future climatic conditions are substantially influenced by individual decisions of farmers as well as market developments and thus difficult to model. Hence we decided to focus on assessing CC effects on the habitat suitability only.

In our study we used three different habitat suitability levels as an estimated threshold for species presence. This way we accounted for uncertainties related to the choice of a fixed threshold level. However, estimating species distributions from current and future habitat suitability is hampered by the fact that a species fundamental niche may be different from its realized niche, for example due to competition with other species or because of the effects the species itself has on its environment [62,63]. Furthermore, it should also be considered that simple presence of a pest species might not be harmful for agriculture if the density remains low. Here, we only looked at habitat suitability and the likelihood of presence or absence of a species. Nevertheless, it is reasonable to assume that higher densities are likely to occur if habitat suitability is high as long as the host species is present.

Furthermore, we were only using agricultural crop intensity as an estimation of agricultural productivity. Yet, areas which are currently not in use and/or less productive, might become so under future CC. Since pest species are highly dependent on their host species and agricultural...
production, this is likely to affect species distribution in a significant way. This aspect was, however, not considered in this study due to the lack of specific data.

Future research should include not only more important pest species in Africa, but also other environmental variables, and individual pest species should be linked to the cultivated areas of their respective host plants. Estimations of the impacts of pest species on agricultural production under projected CC would benefit from more species presence records as well as more detailed current and forecasted maps of crop production areas across the continent.

We believe that the results of this study can help policy makers, extension organisations and farmers to make adapted agricultural management decisions today while anticipating future CC impacts, for instance by choosing crops that are less susceptible to certain pest species. This can help to secure food production and livelihoods of farmers in the coming decades, when some pest species are likely to expand their distribution under CC. The maps can give farmers an orientation which crops are less likely to suffer under pests in the future and which they should avoid planting because of their association with higher risk of infestation. Using species distribution maps and climate scenarios and integrating them into land management decision systems can help to increase agricultural productivity, mitigate global hunger and thereby decrease competition for arable land.

**Supporting Information**

**S1 Fig.** Modelled habitat suitability for *Bactrocera invadens* for current climatic conditions (a) and future climatic conditions (b-e) as mean values over suitability modelled from bioclimatic data for 3 different Generalized Circulation Models (GCM) under each representative concentration pathways (RCP) scenario as well as mean change over all RCP scenarios.

(PDF)

**S2 Fig.** Modelled habitat suitability for *Ceratitis cosyra* for current climatic conditions (a) and future climatic conditions (b-e) as mean values over suitability modelled from bioclimatic data for 3 different Generalized Circulation Models (GCM) under each representative concentration pathways (RCP) scenario as well as mean change over all RCP scenarios.

(PDF)

**S3 Fig.** Modelled habitat suitability for *Tuta absoluta* for current climatic conditions (a) and future climatic conditions (b-e) as mean values over suitability modelled from bioclimatic data for 3 different Generalized Circulation Models (GCM) under each representative concentration pathways (RCP) scenario as well as mean change over all RCP scenarios.

(PDF)

**S4 Fig.** Response curves of environmental variables indicating their effect on the predicted habitat suitability of Maxent with the logistic model output on the y-axis and the environmental variable on the x-axis. Response curves show effect of each variable in isolation on the model which makes interpretation of strongly correlated variables easier.

(PDF)

**S1 File.** Presence records of *Bactrocera invadens* (Dataset A), *Ceratitis cosyra* (Dataset B) and *Tuta absoluta* (Dataset C) used in this study for distribution modelling.

(ZIP)

**Acknowledgments**

The authors acknowledge the help of our colleagues Dr. Dominikus Heinzeller (KIT) and Dr. Sunday Ekesi (*icipe*) as well as the valuable comments and suggestions provided by two anonymous reviewers.
Author Contributions
Conceived and designed the experiments: LBF HEZT. Analyzed the data: JZ LBF. Contributed reagents/materials/analysis tools: HEZT. Wrote the paper: LBF JZ HEZT CB.

References


36. EMBL. Geographically tagged INSDC sequences. European Molecular Biology Laboratory (EMBL); 2014.


45. Hijmans RJ, Cameron SE, Parra JL, Jones PG, Jarvis A. WORLDCLIM—a set of global climate layers (climate grids) [Internet]. Available: www.worldclim.com