



GUIDELINES
FOR THE
IMPLEMENTATION
**OF THE SPECIFIC
PHYTOSANITARY
SURVEILLANCE SYSTEM**
CASE STUDY
***Xanthomonas
oryzae pv. oryzae***



Inter-American Institute for Cooperation on Agriculture (IICA), 2018



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Xanthomonas oryzae pv. *oryzae*

1. PURPOSE

Detection surveillance of *Xanthomonas oryzae* pv. *oryzae* in rice production and host weeds in the COSAVE region.

2. SCOPE

COSAVE region, considering pest records and suitable environmental conditions for the pest.

3. TARGET PEST

Xanthomonas oryzae pv. *oryzae*, see the pest datasheet in Annex 1.

4. DURATION AND APPROPRIATE TIMING

The duration is one (1) growing season, during the active growing stage of the crop, with a weekly frequency between surveys.

5. SITE SELECTION

For the area or site selection process, the following information is required in advance:

- Cadastral map of the region
- Hydrography and geographical characteristics (forests, mountains, lakes, rivers, deserts) in the location
- Risk map for the region
- Area and production of the crop at the political and administrative level as detailed as possible
- International transport routes

Xanthomonas oryzae pv. *oryzae*

5.1. Environmental modeling for the development of risk maps for the region

The environmental niche modeling is commonly used to develop probabilistic maps of species distribution. Among the available modeling techniques, MaxEnt has become one of the most popular tools for modeling species distribution, with hundreds of peer-reviewed articles published each year. The popularity of MaxEnt is mainly due to the short running time, easy operation, small sample size, high simulation precision, the use of a graphical interface and automatic parameter configuration capacities (Morales et. al, 2017; Costa P., Holtz V. 2011; Wang R. et al. 2018).

Risk maps for the target pest can be managed for the risk characterization and the regional prioritization of surveillance activities. In this study case, we use the MaxEnt model, to identify the environmental risk for *Xanthomonas oryzae* pv. *oryzae* with worldwide locations that report the pest and their correlation with the bioclimatic variables from the Worldclim database (<http://www.worldclim.org/>). The details of the method appear in Annex 2 and the resulting regional environmental risk map in Figure 1. The areas in red and yellow mean higher and medium risk respectively, light blue means lower estimated risk. The risk estimation is based on MaxEnt results and the comparison of the presence of the pest and the bioclimatic variables where the pest is reported.

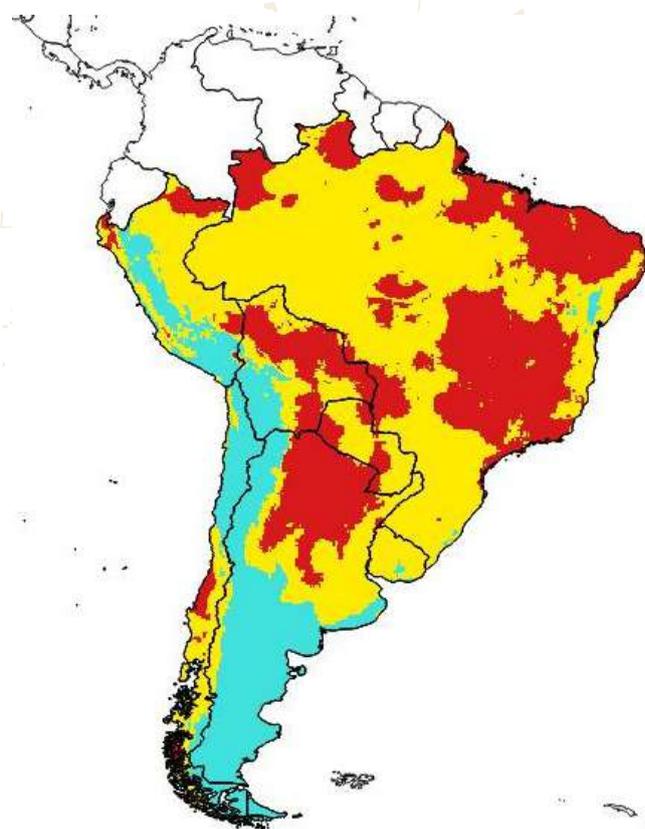


Fig. 1. Environmental risk map of *Xanthomonas oryzae* pv. *oryzae* for the COSAVE region. Higher risk in red, medium risk in yellow and lower risk in light blue. Scale 1:45,000,000. Source: Prepared for STDF/PG/502 COSAVE Project.

Xanthomonas oryzae pv. *oryzae*

5.2. Host area in the region

Xanthomonas oryzae pv. *oryzae* hosts belong to the genus *Oryza* as well as wild species from the Poaceae and Cyperaceae family, as described in the datasheet (Annex 1). Rice production is economically important in the region. It accounts for about 214,570 hectares in Argentina; 181,497 in Bolivia; 2,162,178 in Brazil; 20,937 in Chile; 167,088 in Peru; 131,740 in Paraguay, and 164,400 hectares in Uruguay. Based on rice production information at a first administrative-geopolitical level in each COSAVE country, it is possible to develop a host risk map. As Fig. 2 illustrates, the NPPO can determine the levels as high, medium and low based on the importance of production in each identified administrative-geopolitical level. The method for its preparation is discussed in Annex 2.

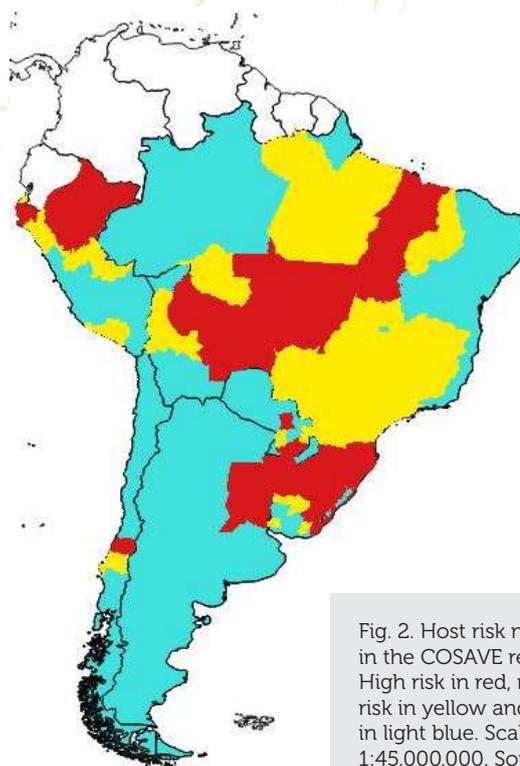


Fig. 2. Host risk map—rice in the COSAVE region. High risk in red, medium risk in yellow and low risk in light blue. Scale 1:45,000,000. Source: Prepared for STDF/PG/502 COSAVE Project.

5.3. Regional risk for *xanthomonas oryzae* pv. *oryzae*

It is possible to integrate the environmental and host risk maps in one regional risk map, as is illustrated in Fig. 3. The method is presented in Annex 2 and is based on the reclassification of high-risk category with a value of two (2), medium with one (1) and low with zero (0); and the use of the raster mathematical multiplication function of a geographic information system like QGIS. The method for its preparation is described in Annex 2.



Fig. 3. Regional risk map of *Xanthomonas oryzae* pv. *oryzae* for COSAVE. High risk in red, medium risk in yellow and low risk in light blue. Scale 1:45,000,000. Source: Prepared for STDF/PG/502 COSAVE Project.

*Xanthomonas oryzae pv. oryzae***5.4. Selected sites for the surveillance**

Considering pest reports, it is necessary to identify the potential considerations for the entry of the pest. In this regard, it is important to identify the main transporting routes and the main rivers that can spread the pest with rice seeds, as well as nearby places of production.

It is possible to integrate the regional risk of the pest and the location of routes and rivers consolidated by the Consejo Suramericano de Infraestructura y Planeamiento (COSIPLAN) (Available at: <http://www.sig.cosiplan.unasursg.org/node/15>, on January 5, 2018). For a better management of field actions, the NPPO can identify squares (grids), with the appropriate size to describe a uniform population for the sampling, as shown in Fig. 4 and 5, with 100 km grids. The method for preparation is also described in Annex 2.

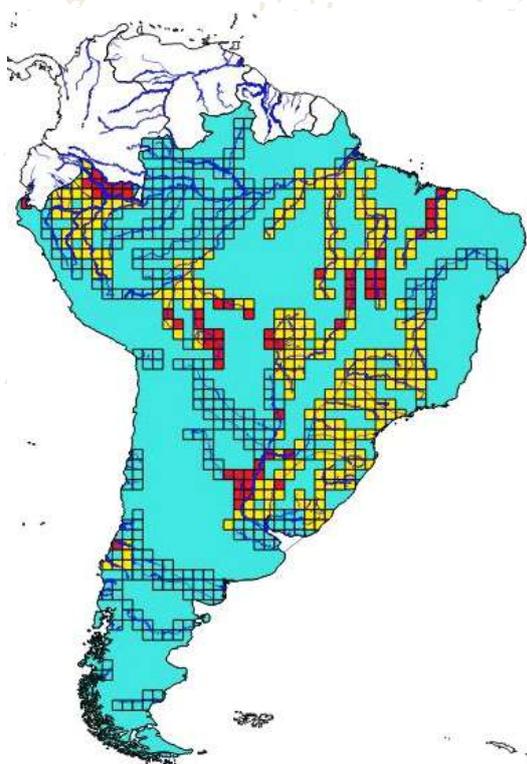


Fig. 4. 100 km squares used for the integration of main rivers location and the regional risk information of the pest (high risk in red, medium risk in yellow and low risk in light blue. Scale 1:45,000,000. Source: Prepared for STDF/PG/502 COSAVE Project.

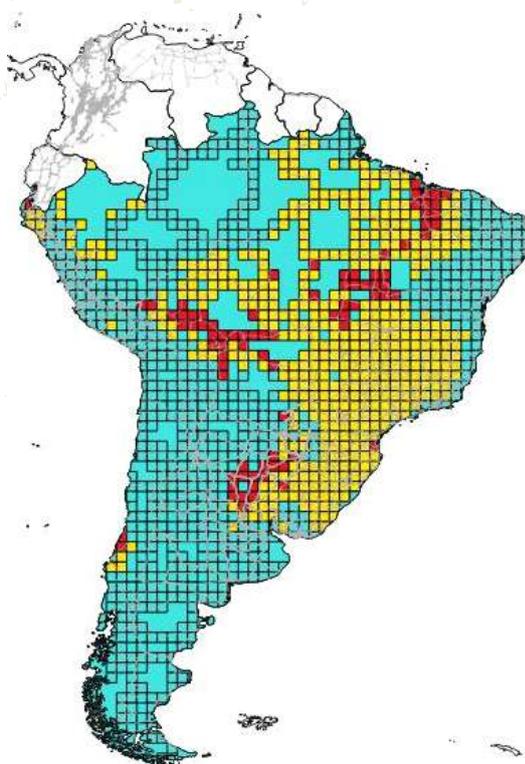


Fig. 5. 100 km squares used for the integration of main routes location and regional pest risk information (high risk in red, medium risk in yellow and low risk in light blue. Scale 1:45,000,000. Source: Prepared for STDF/PG/502 COSAVE Project.

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The NPPO can identify risk squares (grids) with a code and review it with additional information and then identify the appropriate place for the surveillance.

In addition, with the rivers and transport routes information, the NPPO can identify the high-risk squares (grids). The number of squares (grids) separated by risk category is presented in table 1 and 2. This identification should be complemented with additional information from the NPPO.

Table 1. Number of squares (grids) in major rivers by *Xanthomonas oryzae pv. oryzae* risk level.

COUNTRY	HIGH	MEDIUM	LOW	TOTAL
ARGENTINA	13	16	84	113
BOLIVIA	12	20	19	51
BRASIL	22	196	131	349
CHILE	1	3	23	27
PERÚ	11	37	28	76
PARAGUAY	1	3	14	18
URUGUAY		10	11	21
	60	285	310	655

Source: Prepared for STDF/PG/502 COSAVE Project.

Table 2. Number of squares (grids) in major routes by *Xanthomonas oryzae pv. oryzae* risk level.

COUNTRY	HIGH	MEDIUM	LOW	TOTAL
ARGENTINA	15	17	242	274
BOLIVIA	23	24	32	79
BRASIL	38	383	224	645
CHILE	2	4	75	81
PERÚ	3	18	76	97
PARAGUAY	2	9	14	25
URUGUAY	1	11	14	26
	84	466	677	1227

Source: Prepared for STDF/PG/502 COSAVE Project.

Xanthomonas oryzae pv. *oryzae*

6. PLANNING

6.1. Preliminary activities

For a better operational organization, it is necessary to recognize the national characteristics of the risk, places and supplies for surveillance implementation. This allows the NPPO to evaluate, manage and systematize the activity. In this regard, it is important to include the following:

- The annual operating plan with the inclusion of the budget, geographical distribution, task schedule and the implementation time.
- Coordination with the diagnostic laboratory, including the protocol and the number of samples to be submitted.
- Cadastral map of the region.
- Hydrography and geographical characteristics (forests, mountains, lakes, rivers or deserts) in the location.
- Hosts area and production at the administrative-geopolitical level as detailed as possible.
- Phenology of the involved hosts.
- Location of rice seedbeds, collection and storage centers, and risk areas.
- International transport routes.
- Collate pest information in a datasheet like Annex 1.
- Manage permissions to enter private property in advance.
- Ensure the required supplies and resources for the surveillance.
- Training for directly involved staff.

Xanthomonas oryzae pv. oryzae

6.2. Surveillance methodology

The visual inspection or survey can be an effective surveillance method when the characteristics or symptoms of the pest enable identification, which can be confirmed with the laboratory diagnosis. Moreover, it is important to consider:

- The surveillance of *Xanthomonas oryzae pv. oryzae* is performed with the visual inspection or survey in production locations, tissue sampling and pest isolation.
- The number of inspections or surveys can follow the recommendations of the “Guidelines for the implementation of the specific phytosanitary surveillance system”.
- Identification of the varieties and other pest susceptible species.
- Ecological conditions that encourage the presence of the pest.
- Rice seedbeds.

6.2.1. Required supplies

- Disposable cloths and gloves.
- Field knife for the sampling.
- Alcohol burner.
- Alcohol 70% for hand disinfection.
- Bleach 10% for sample collection.
- Identification labels.
- Plastic and paper bags for sampling.
- Entomological clamps.
- Writing pads.
- Pencil.
- Dishcloth for tool cleaning.
- Devices to capture geo-referenced field data.
- Templates for data collection.

6.2.2. Sample collection and submission

In case that the visual survey (inspection) can identify *Xanthomonas*, it is necessary to confirm its identification safeguarding its integrity and submitting the sample to the laboratory. This official diagnosis requires prior coordination of the number of the samples that can be identified.

*Xanthomonas oryzae pv. oryzae***6.3. Recording surveillance activities**

In order to collate and implement automated reports, the activities of the information record for the Surveillance System should be performed in a standardized and integrated manner. In this regard, we present a general template for the identification of the required information to record *Xanthomonas oryzae pv. oryzae* surveillance activities.

Item	COUNTRY AND ACTIVITY			PLACE		HOST AND RESULTS			INCIDENCE AND SEVERITY				
	País	Fecha (dd/mm/aaaa)	Actividad de Vigilancia Inspección, muestreo, trampeo, otro _____ (especificar)	Coordenadas geográficas decimales: LATITUD	Coordenadas geográficas decimales: LONGITUD	Hospedante (Citrus sinensis, Citrus reticulata, Citrus unshiu, Citrus aurantifolia, Oryza sativa, otro _____ (especificar)	Tipo de Predio (Comercial, vivero, traspatio, aislado, otro _____ (especificar)	Resultado (Ausente o Presente)	Incidencia encontrada	Descripción de la incidencia evaluada (% u otro _____ (especificar)	Severidad encontrada	Severidad (Grados u otro _____ (especificar)	Observación
1	2	3	4	5	6	7	8	9	10	11	12	13	14
1													
2													
3													
4													
5													
6													
7													
8													
9													
10													
11													
12													

Fig. 6. General template to record *Xanthomonas oryzae pv. oryzae* surveillance activities. Source: Prepared for STDF/PG/502 COSAVE Project.

In addition, this general template should have modeled fields in order to avoid mistakes in recording information and the use of computing platforms. For example, Open Data Kit is capable of collecting geo-referenced information with Android devices. The details, templates and explanations for its use are available in the ODK folder in the link: <https://goo.gl/WYFe6a>.

6.4. Biosafety

To achieve biosecurity in surveillance actions, it is necessary to include:

- The use of new disposable cloths during the entry to each surveillance site.
- The handling of plants or samples should be done with gloves to avoid contaminating nearby locations.
- Surveillance waste should be kept in properly closed plastic bags.
- Any waste material should follow NPPO waste recommendations.
- Hand disinfection with approved supplies is required after each activity.

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7. COMMUNICATION

It is important to produce reports with the results of the various level of decision making.

8. AUDIT

Through the central coordination, each NPPO will perform auditing activities at any stage of the process, analyzing the data in the system, and monitoring the quality of the field tasks performance, among others.

9. REFERENCES

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Sheldeman X & van Zonneveld M. 2011. Manual de Capacitación en Análisis Espacial de Diversidad y Distribución de Plantas. Bioversity International, Rome, Italy. 186 pp. Available on October 27, 2017, at:

<http://www.bioversityinternational.org/e-library/publications/detail/manual-de-capacitacion-en-analisis-espacial-de-diversidad-y-distribucion-de-plantas/>

Singh, R.A., and Rao, M.H.S. 1977. A simple technique for detecting *Xanthomonas oryzae* in rice seeds. *Seed Science and Technology* 5: 123-127. Available on October 27, 2017, at: <http://download.ceris.purdue.edu/file/289>

Wang R, Li Q, He S, Liu Y, Wang M, Jiang G. 2018. Modeling and mapping the current and future distribution of *Pseudomonas syringae* pv. *actinidiae* under climate change in China. *PLoS ONE* 13(2): e0192153. Available on February 26, 2018, at: <https://doi.org/10.1371/journal.pone.0192153>

10. ANNEXES

Annex 1. Data sheet of the pest

Annex 2. Modeling for *xanthomonas oryzae* pv. *oryzae*

Xanthomonas oryzae pv. *oryzae*

ANNEX 1

DATA SHEET OF THE PEST

***Xanthomonas oryzae* pv. *oryzae* (Ishiyama 1922) Swings et al. 1990**

Synonyms:

Bacterium oryzae (Uyeda & Ishiyama) Nak., 1928; *Phytomonas oryzae* (Ishiyama) Magrou 1937; *Pseudomonas oryzae* Ishiyama; *Xanthomonas campestris* pv. *oryzae* (Ishiyama 1922) Dye 1978; *Xanthomonas itoana* (Tachinai) Dowson; *Xanthomonas kresiek* Schure 1953; *Xanthomonas oryzae* (Ishiyama 1922) Swings et al. 1990; *Xanthomonas translucens* f. sp. *oryzae* Jones et al; Dowson; Ishiyama; Pordesimo.

Taxonomic rank:

Phylum: Proteobacteria

Class: Bacteria

Order: Xanthomodales

Family: Xanthomonadaceae

Genus: *Xanthomonas*

Species: *Xanthomonas oryzae* pv. *oryzae* (Ishiyama 1922) Swings et al. 1990

Common names:

Enfermedad bacteriana de las hojas del arroz (spanish); maladie bactérienne des feuilles du riz (french); bacterial leaf blight of rice; kresiek disease; rice bacterial leaf blight; rice kresiek disease (english).

Hosts:

Cenchrus ciliaris, *Cynodon dactylon*, *Cyperus difformis*, *Cyperus rotundus*, *Echinochloa crus-galli*, *Leersia hexandra*, *Leersia oryzoides*, *Leptochloa chinensis*, *Megathyrsus maximus*, *Oryza* spp., *Oryza longistaminata*, *Oryza sativa*, *Paspalum scrobiculatum*, *Urochloa mutica*, *Zizania aquatica*, *Zizania palustris*, *Zoysia japonica* (CABI, 2017).

Geographic distribution:

America: Mexico, United States, Costa Rica, El Salvador, Honduras, Nicaragua, Panama, Colombia, Ecuador, Venezuela (CABI, 2017).

Asia: Pakistan, Philippines, Sri Lanka, Taiwan, Vietnam (CABI, 2017).

Africa: Burkina Faso, Cameroon, Egypt, Gabon, Gambia, Madagascar, Mali, Niger, Nigeria, Tanzania, Togo (CABI, 2017).

Europe: Russia (CABI, 2017).

Oceania: Australia (restricted) (CABI, 2017).

Xanthomonas oryzae pv. *oryzae*

Biology:

Xanthomonas oryzae pv. *oryzae* survives during the off-season in seed, weed hosts, rice volunteers and infected rice straw and stubble. The evidence for significant seed transmission is contradictory. Survival in soil is limited to a few months at the most in relatively cool, moist conditions (the bacterium dies rapidly in hot, dry conditions) and is thus of little significance in single-crop rice systems. Transmission in irrigation water and floodwater is important during the cropping period, but survival in water is limited to a few days (CABI, 2017).

The bacterium invades rice plants through hydathodes on leaves, root-growth cracks and wounds. Seedling roots are wounded when pulled from the seedbed; and leaf tips are often cut before transplanting. When inside the plant, the bacterium enters the vascular system in which it spreads. Bacteria eventually ooze out of water pores on hydathodes (CABI, 2017).

Signs and symptoms:

The symptoms known as 'kresek' (wilting, desiccation of leaves and death, characteristic of systemic infection) occur with particular combinations of virulent isolates and susceptible cultivars, probably when the vascular system is blocked by bacterial cells and extracellular polysaccharide. Kresek is associated with tropical storms which spread the pathogen and also wound rice plants. High temperature (30°C) and humidity favor the disease (CABI, 2017).



Fig. 7. Symptoms of *Xanthomonas oryzae* pv. *oryzae* due to artificial inoculation (Source: Purdue University, 2017).

Symptoms appear on leaves of young plants, after planting out, as pale-green to grey-green, water-soaked streaks near the leaf tip and margins. These lesions coalesce and become yellowish-white with wavy edges. The whole leaf may eventually be affected, becoming whitish or greyish and then dying. Leaf sheaths and culms of more susceptible cultivars may be attacked. Systemic infection, known as kresek, results in wilting, desiccation of leaves and death, particularly of young transplanted plants. In older plants, the leaves become yellow and then die. In later stages, the disease may be difficult to distinguish from bacterial leaf streak (*X. oryzae* pv. *oryzicola*) (CABI, 2017).

CASE STUDY

Xanthomonas oryzae pv. *oryzae*

Entry pathway:

The bacteria can only move short distances in infected crops. The bacteria can only move long distances in infected rice seeds. The bacteria are usually found in the glumes, but may also penetrate the endosperm (EPPO, 2003).



Survey (inspection) and detection:

Symptoms appear on leaves of young plants, after planting out, as pale-green to grey-green, water-soaked streaks near the leaf tip and margins. These lesions coalesce and become yellowish-white with wavy edges. The whole leaf may eventually be affected, becoming whitish or greyish and then dying.

Fig. 8. Infected rice seedlings. Infected leaves wilt and roll up, turning grayish-green to yellow, until the whole seedling dies. Plants which have survived the disease are stunted and yellowish (Picture from EPPO, 2017).

Leaf sheaths and culms of more susceptible cultivars may be attacked. Systemic infection, known as kresek, results in wilting, desiccation of leaves and death, particularly of young transplanted plants. In older plants, the leaves become yellow and then die. In later stages, the disease may be difficult to distinguish from *X. oryzae* pv. *oryzicola* (CABI, 2017).

Pest impact:

Bacterial leaf blight is the most serious disease of rice in South-East Asia, particularly since the widespread cultivation of dwarf high-yielding cultivars (EPPO, 2003). In Japan, where figures are available, up to 400,000 ha may be affected annually, with losses of 20-30% and up to 50%. In Africa, losses of 2.7-41% in grain yield have been found (CABI, 2017).

In the Philippines, losses are estimated at 22.5% (in wet seasons) and 7.2% (in dry seasons) in susceptible varieties and 9.5 and 1.8%, respectively, in resistant crops (EPPO, 2003).

Pest control and mitigation measures:

Preventive phytosanitary control measures include:

- Use of healthy and treated seeds.
- Resistant varieties.
- Careful attention to crop management (for example, water control, avoidance of damage to seedlings) is very important.
- Restricting nitrogen fertilizer applications to about 80-100 kg N/ha.
- Chemical control is not recommended (CABI, 2017).

Xanthomonas oryzae pv. *oryzae*

ANNEX 2

Modeling for *Xanthomonas oryzae* pv. *oryzae*

A. Data:

This section will discuss the origin and type of data used in the case studies. All the working files, software, references, and other data are available in the link: <https://goo.gl/WYFe6a>.

A1. Environmental raster data (*.tif)

For geo-referenced climate data, go to <http://www.worldclim.org/> and then click on Version 2 (<http://worldclim.org/version2>) for updated climate data.



Fig. 9. Link to Version 2 of the web page: <http://www.worldclim.org>.



Fig. 10. Download the 10 minutes data.

NOTE: The folder with this data is named "WC10y1990tiff" and is located in the link: <https://goo.gl/WYFe6a>.

NOTE: The bioclimatic variables are: BIO1 = Annual Mean Temperature, BIO2 = Mean Diurnal Range (Mean of monthly (max temp - min temp)), BIO3 = Isothermality (BIO2/BIO7) (* 100), BIO4 = Temperature Seasonality (standard deviation *100), BIO5 = Max Temperature of Warmest Month, BIO6 = Min Temperature of Coldest Month, BIO7 = Temperature Annual Range (BIO5-BIO6), BIO8 = Mean Temperature of Wettest Quarter, BIO9 = Mean Temperature of Driest Quarter, BIO10 = Mean Temperature of Warmest Quarter, BIO11 = Mean Temperature of Coldest Quarter, BIO12 = Annual Precipitation, BIO13 = Precipitation of Wettest Month, BIO14 = Precipitation of Driest Month, BIO15 = Precipitation Seasonality (Coefficient of Variation), BIO16 = Precipitation of Wettest Quarter, BIO17 = Precipitation of Driest Quarter, BIO18 = Precipitation of Warmest Quarter, BIO19 = Precipitation of Coldest Quarter.

NOTE: In the link to Version 1.4, we share access to the projected future data for 2050 and 2070, under four (4) scenarios called RCP ("Representative Concentration Pathways").

*Xanthomonas oryzae pv. oryzae***A2. Vector data ("*.shp")**

The Consejo Suramericano de Infraestructura y Planeamiento (COSIPLAN) collated cartographic information about the following layers in the region:

Border control (CSP_AH070_N), Populated center (CSP_AL105_N), Railway Line (CSP_AN010_L), Railway station (CSP_AN070_N), Main road (CSP_AP030_L), Port (CSP_BB005_N), Lake (CSP_BH080_P), River (CSP_BH140_L, CSP_BH140_P), Administrative border (CSP_FA000_L), (CSP_FA001_L, CSP_FA001_P), 3er level administrative zone (CSP_FA002_P), Border crossing (CSP_FA125_N), Airport (CSP_GB001_N), and others.

This information is packed in a zip file, available on October 27, 2017 in the link <http://www.sig.cosiplan.unasursg.org/node/15>, unzip and save the file in a separated folder like DATA in the link: <https://goo.gl/WYFe6a>.

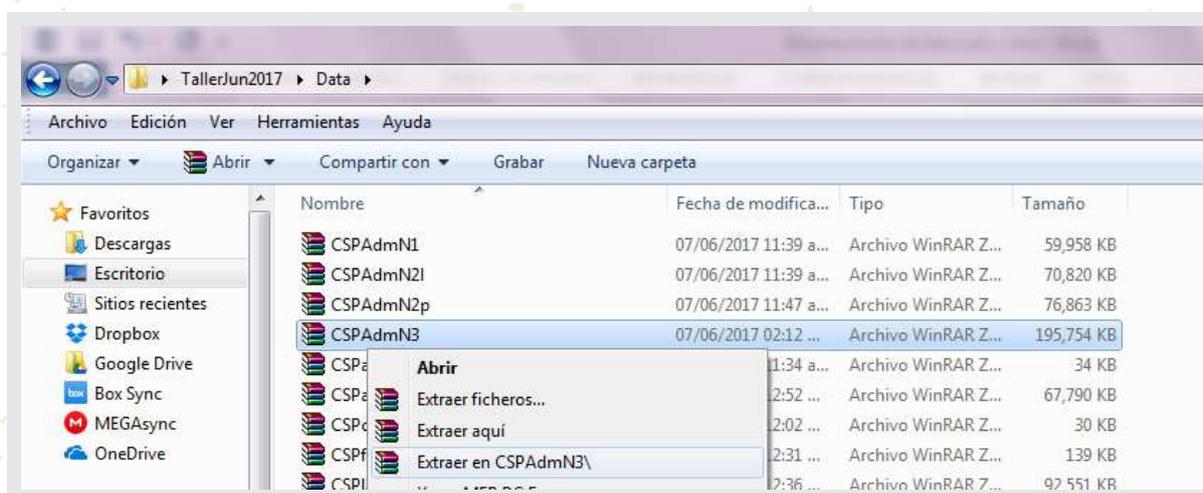


Figura 11. Descargar las capas vectoriales y descomprimirlas haciendo un clic con el botón izquierdo del mouse.

A3. Geo-referenced data of pests (use of geographic coordinates as an example)

In the CABI Invasive Species Compendium webpage, refer to: *Xanthomonas oryzae pv. oryzae* or go to the link: <http://www.cabi.org/isc/datasheet/56956>

Xanthomonas oryzae pv. *oryzae*

Go to the lower part of the world distribution map and to the link "Download CSV file" or "Comma separated values" that can be opened with Microsoft Excel.



Fig. 12. Download the CSV ("Comma separated values") files.

	A	B	C	D
1	species	Longitude	Latitude	
2	Xoryzaeoryza	93	10	
3	Xoryzaeoryza	79	16	
4	Xoryzaeoryza	117	32	
5	Xoryzaeoryza	93	26	
6	Xoryzaeoryza	134	-20	
7	Xoryzaeoryza	90	24	
8	Xoryzaeoryza	2.25	9.5	
9	Xoryzaeoryza	85.75	25.75	

Fig. 13. Delete the data with the "Absend" or "Restricted distribution" pest situation and arrange the elements as is shown.

Open the data in Microsoft Excel, delete the reports of the pest with the situation label as "Absent" or "Restricted distribution" and include first the column species, then longitude and after latitude. Delete the other columns.

Double-check the existence of the species column first, then the longitude and latitude columns respectively and save it as "Xoo.csv" ("Comma separated value"). This data is available in the "cXanthomonasoryzaoryzae" folder in the link: <https://goo.gl/WYFe6a>.



*Xanthomonas oryzae pv. oryzae***B. Software:****B1. QGIS**

QGIS (previously called Quantum GIS) is an open source Geographic Information System (GIS) for the GNU/Linux, Unix, Mac OS and Microsoft platforms, it supports many formats and functions for shapefiles, raster files and databases. In addition, it has defined extensions (<http://plugins.qgis.org/plugins/>) that make it one of the best of its kind and is in continuous development.

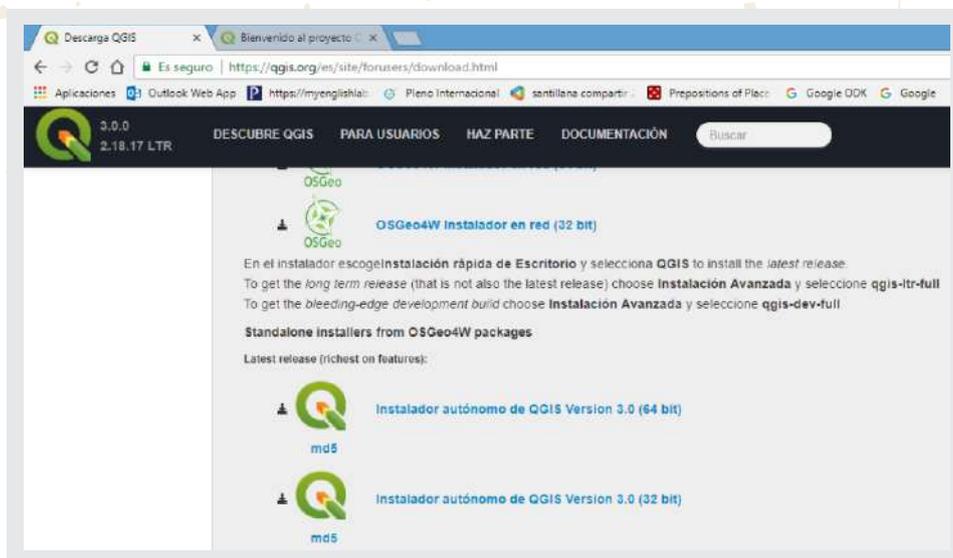


Fig. 14. Webpage to download QGIS 3.0.0 "Girona". Available on February 23, 2018 in: <https://www.qgis.org/es/site/forusers/download.html>Note. We reco

NOTE: We recommend reading and practicing with the training manual of the software. This is available on January 19, 2018, at: <https://www.qgis.org/en/site/forusers/index.html#download> and tutorial videos available on the Internet.

B2. MaxEnt**B2.1. Installing Java, if not already installed**

To verify if Java is installed on your computer, go to the link: <https://www.java.com/es/download/installed.jsp> with Internet Explorer

To install Java in Internet Explorer, go to the link: http://www.java.com/es/download/help/ie_online_install.xml

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To install Java in Mozilla Firefox, go to the link:
http://www.java.com/en/download/help/firefox_online_install.xml

NOTE: Java is not supported in Google Chrome

B.2.2. Installing MaxEnt

The MaxEnt software is available on January 19, 2018, at:
https://biodiversityinformatics.amnh.org/open_source/maxent/ (with small letters and a lower hyphen between open and source).

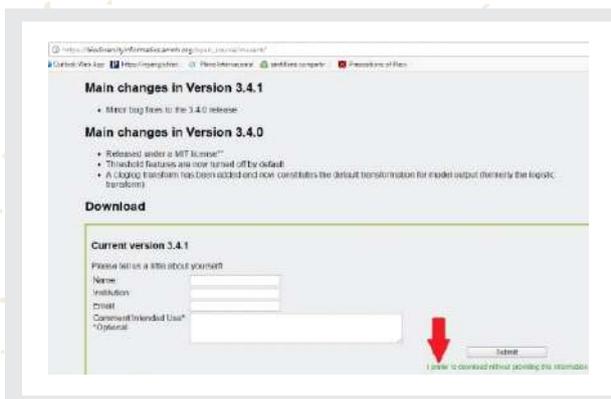


Fig. 15. Link to download MaxEnt software and optional requirement of user personal data.

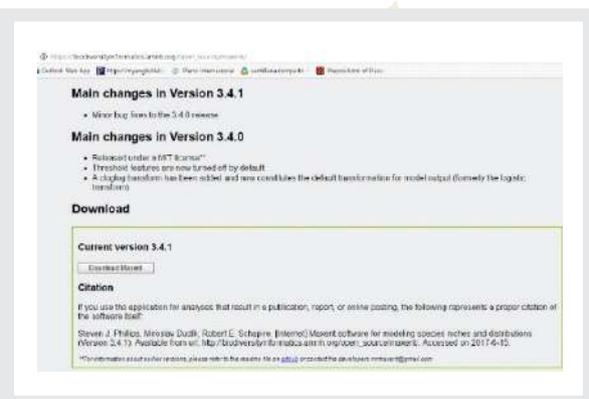


Fig. 16. Download Version 3.4.1. and the cited reference of the MaxEnt software.

C. Activities

C1. Opening a shapefile (or vector layer) and converting the raster data files ("*.tiff" to "*.asc") for the use of MaxEnt software in the modeling.

C1.1. Opening a shapefile in QGIS:

A shapefile has (at least) three files with the same name but with a different extension. The file with the SHP ("*.shp") extension is the main file and includes spatial features.

Open the QGIS with the short option: add vector layer (left margin) or the "Layer" tab, as shown in the below figure:

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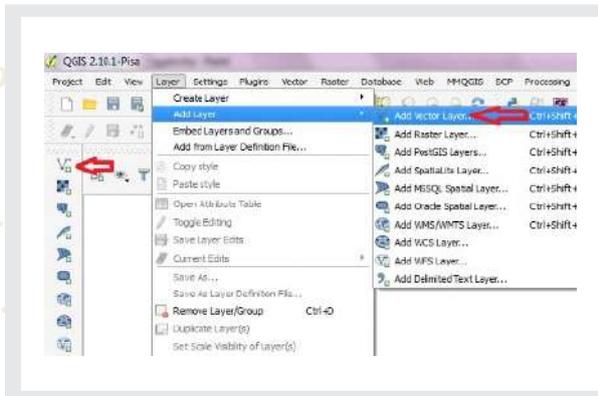


Fig. 17. Search for the DATA folder and unzip *.shp files.

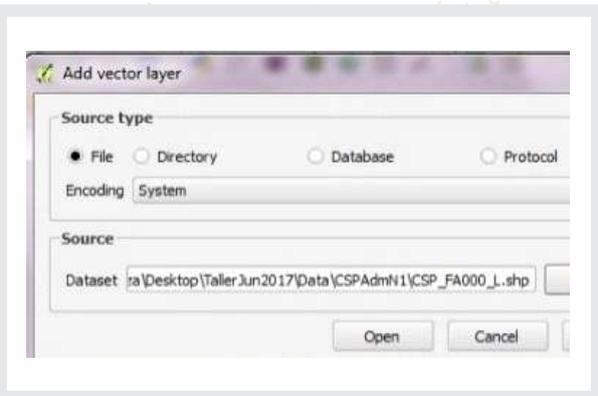


Fig. 18. Locate the shapefile (*.shp) and click on "Open".

In the "Project" tab in QGIS, select "Save as" to save the project.

NOTE: Check the right lower edge for the CRS "Coordinate Reference System" in World Geodetic System 1984 - WGS84 with the code: 4326, recommended because is a world-standardized system.

C.1.2. Converting raster data ("*.tif" to "*.asc"):

A raster is a set of pixels with the same size; each pixel has a different value for a variable (including temperature, type of soil). The size of the pixel is called "resolution", and its selection depends on the geographical scope and the objective of the project. To open the raster file, choose the "Add Raster Layer" option on the left side, as shown in the figure below. Locate, select and open files like "bio10m01.tif" from the "WC10y1990tiff" file.

Approximate size of geographic units (at the Equator), rounded (in km)	
Degrees	Size
1 degree	111 km
10 minutes	18 km
5 minutes	9 km
2.5 minutes	5 km
30 seconds	1 km

Fig. 19. Correspondencia aproximada entre grados y kilómetros en el Ecuador (de: Sheldeman X. & van Zonneveld M. 2011).

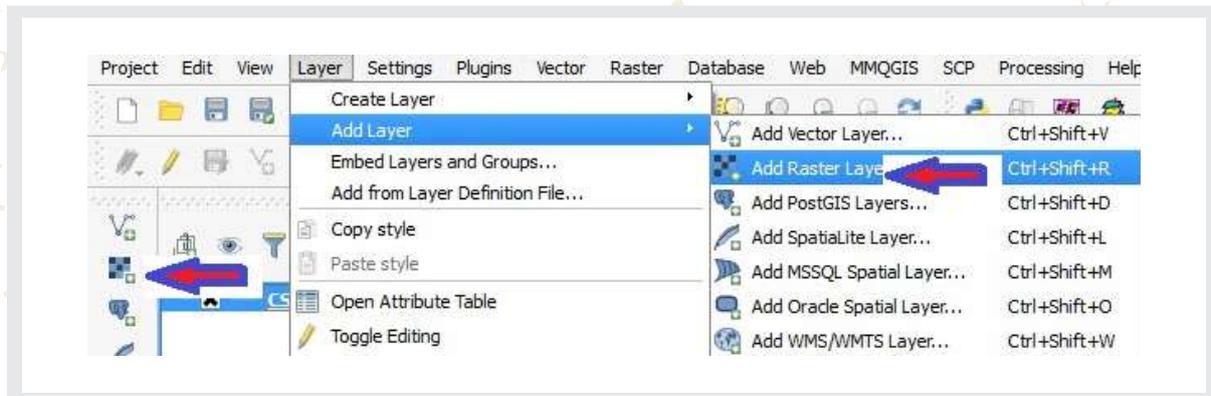
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Fig. 20. Command to open raster (*.tif) files, indicated with the red arrows in the Layer/Add layer/Add Raster Layer tab.

For the conversion, select the "Raster/Conversion/Translate" option, as shown in Fig. 20, in the box "Input Layer" select each file "*.tif" and in the box "Output file" the location of the "*.asc" files in a folder like WC10y1990asc (not the WC10y1990tiff folder) and select the Spatial Reference System (SRS) EPSG:4326, as illustrated in Fig. 22.

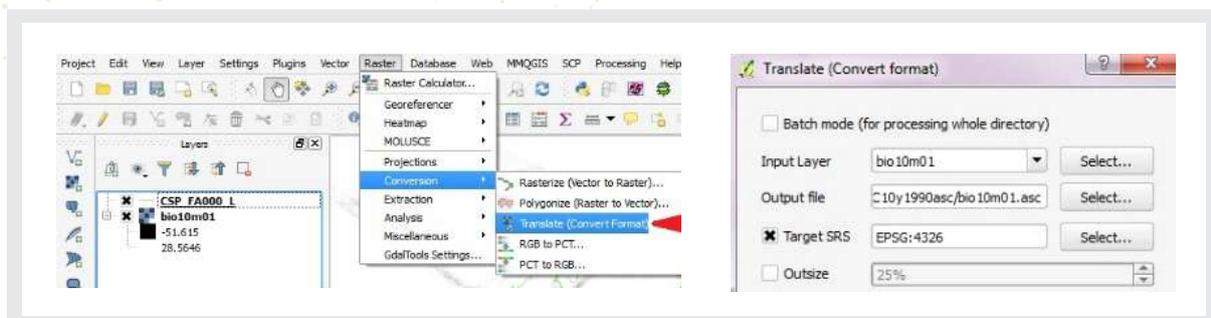


Fig. 21. Command "Translate" to convert raster "*.tif" files to "*.asc".

Fig. 22. Details of the conversion, specifying the "*.asc" extension.

Double check extension "*.asc" in the output file and the selection of the Spatial Reference System.

Repeat the same procedure with the 19 layer of the bioclimatic variables downloaded in the section A1, which are available in the link: <https://goo.gl/WYFe6a> and the WC10y1990tiff folder.

NOTE: The raster "*.asc" files are also available in the WC10y1990asc folder in the link: <https://goo.gl/WYFe6a>.

*Xanthomonas oryzae pv. oryzae***C2. Using MaxEnt for the modeling.**

To open the MaxEnt software, click on the “maxent.jar” file. Then open the “Settings” option, select “Basic” and include the standard option and the box “Random test percentage” write 25 for an additional software test with 25% of the samples. Then close the window.

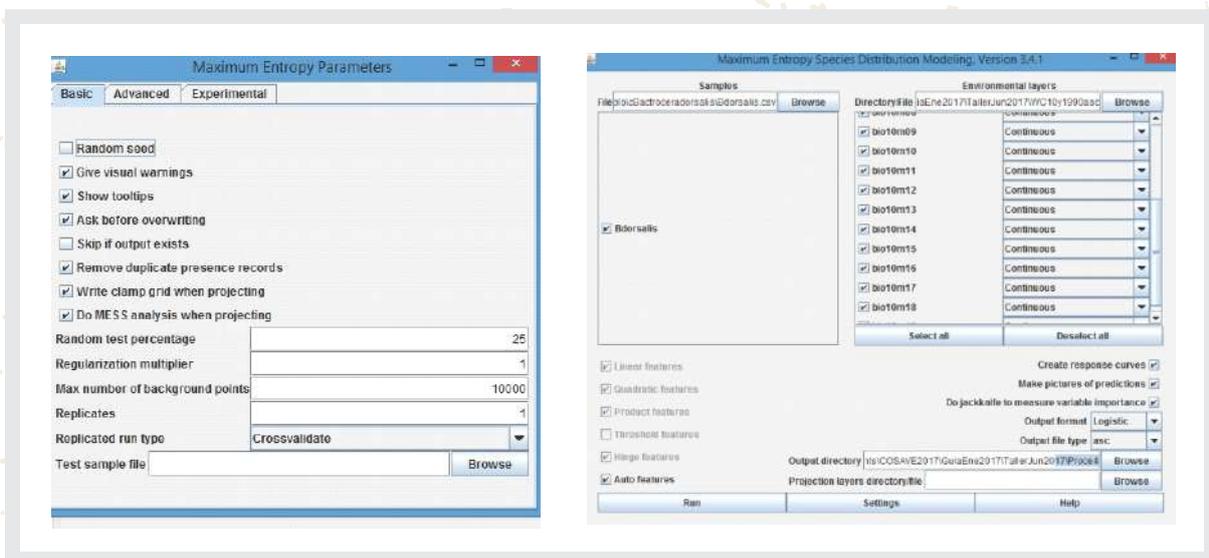


Fig. 23. In “Settings” and “Basic”, include the standard options, as illustrated here.

Fig. 24. Main screen where geo-referenced data and bioclimatic variables are included in “*.asc” format.

In the main screen, in the samples option, select the “*.csv” (“Comma separated values” not “*.xls”) file with the geo-referenced pest data. This was discussed in section A3.

Next, select “Auto features”, “Create response curves”, “Make pictures of predictions”, “Do jackknife to measure variable importance”, “Logistic format” and “asc file type”, and select a folder for the output of the files, thus ordering a complete modeling, with figures like “Do jackknife” that visually describe the contribution of each bioclimatic variable to the final model. In the other part with the environmental layers option, choose the folder with the converted variables to the “*.asc” format, i.e. a raster file like the one described in section A1.

For practical purposes, record warning messages like “Unused field” or “Missing environmental data” and click “Ok”.

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Fig. 25. Record messages like: "Unused field" o "Missing environmental data".

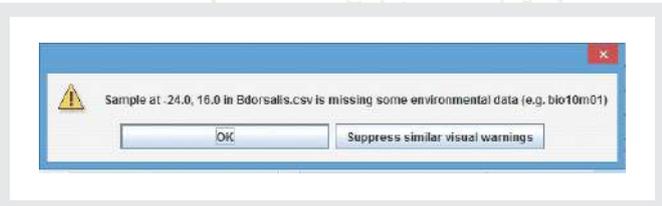


Fig. 26. After checking the repetition of messages, you can select the "Suppress similar visual warnings" option.

The results of the environmental modeling are available in the *Xoryzaeoryzae.html* file, which can be opened with any web browser. The raster file of the map is available in the *Xoryzaeoryzae.asc* file, as illustrated in the figure below.

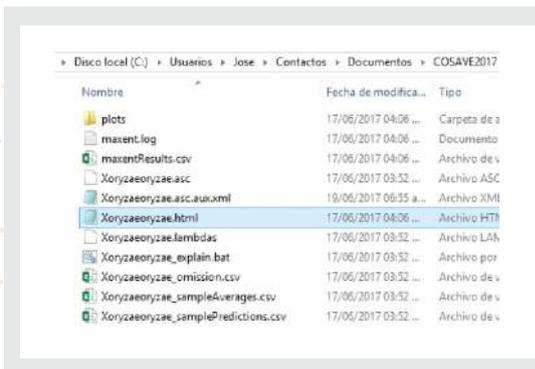


Fig. 27. File with the results of the MaxEnt modeling.

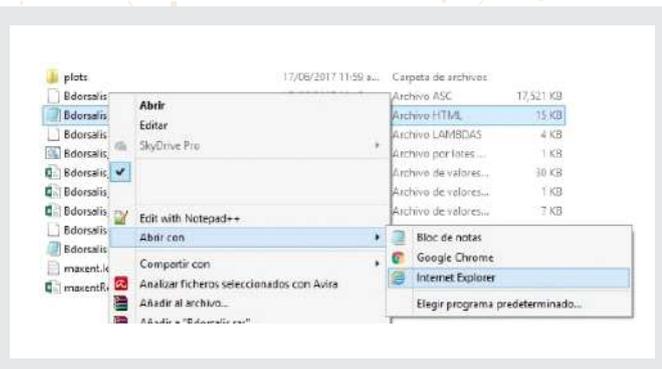


Fig. 28. Opening the "*.html" file with the modeling. The "*.asc" file is a raster, as those described in section A1.

C3. Open a map in geographical coordinates, creating a 100 km reference grid, and a centroid in each grid.

Develop a grid with a reference size of 100 Km or 0.9 decimal degrees, based on the equivalences of grades and kilometers from Sheldeman X. & van Zonneveld M. 2011, shown in the Fig. 19.

First, open a shape vector layer from COSIPLAN (<http://www.sig.cosiplan.unasursg.org/node/15>).

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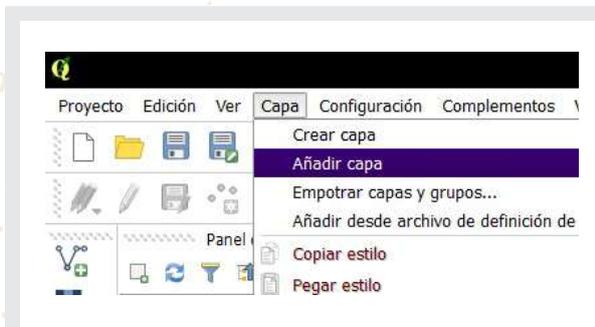


Fig. 29. Open a reference vector layer with the ".shp" extension.

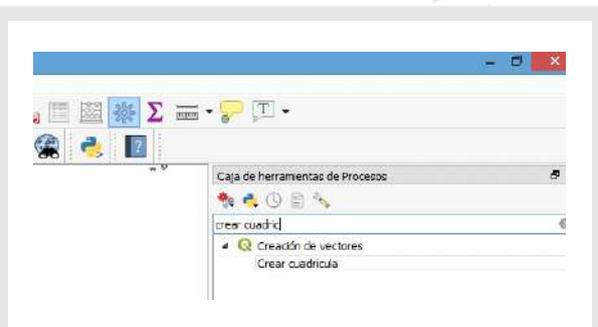


Fig. 30. Find the "Create Grid" function in the "Processing toolbox" (see the note below) and click it.

NOTE: If the Processing toolbox is not available, please select it in tab "View", "Panels" and then "Processing toolbox".

In the "Grid type" choose the "Rectangle (polygon)" option in the "Grid extent" choose "select extent on canvas" dragging the mouse on the map to create the grid. Complete "Vertical spacing" and "Horizontal spacing" with the 0.9 data. Check that EPSG 4326 – WGS84 is in "Grid CRS" and provide a route to save the grid file. Finally, click on "Run in Background" as illustrated in the figure below.

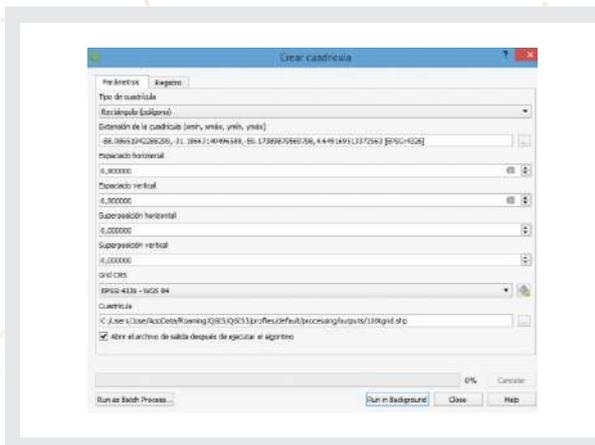


Fig. 31. Details included in the "Create grid" function.

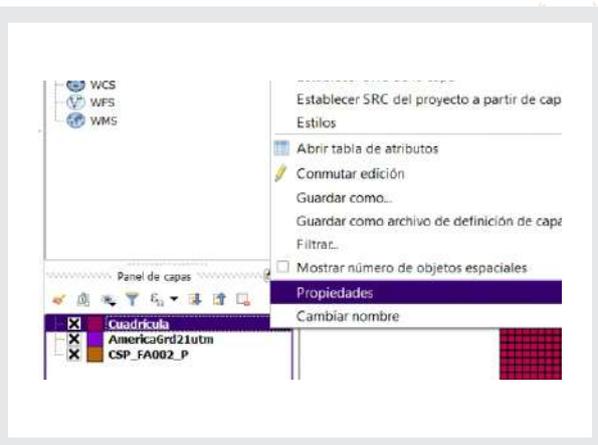


Fig. 32. This is the grid file in the "Layers" panel. To make it transparent, select "Properties" with the right button.

NOTE: If the Processing toolbox is not available, please select it in tab "View", "Panels" and then "Processing toolbox".

In the "Layers" panel, you can change the grid file appearance and its location in the map with a right click on its name. To make it transparent, select "Options", "Style", "Single symbol", "Fill", "Simple fill", and in "Fill style" select "No Brush". Front or background movement on the map is commanded with a change in file position in the "Layers" panel.

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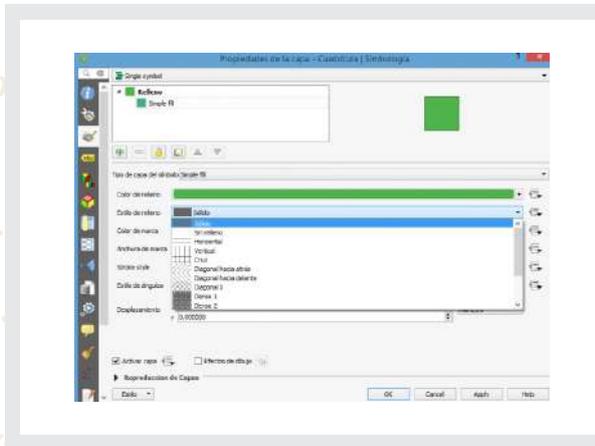


Fig. 33. To make the grid transparent, select "Single symbol", "Fill Style", then "No brush".

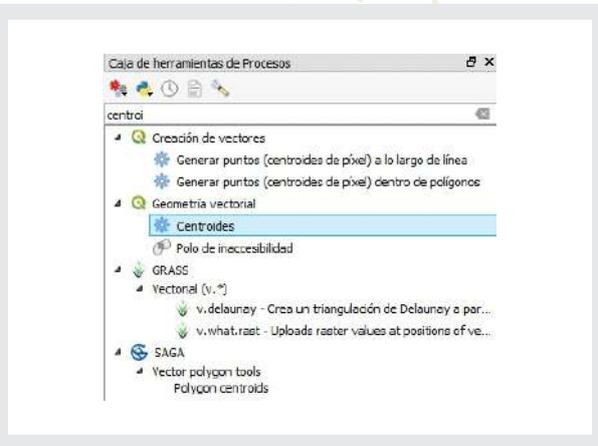


Fig. 34. To obtain centroids in each grid, look for the "Centroids" function ("Vector geometry" set) in "Processing Toolbox".

NOTE: If the Processing toolbox is not available, please select it in tab "View", "Panels" and then "Processing toolbox".

Using this vector "*.shp" file of centroids, you can extract values from the modeling.

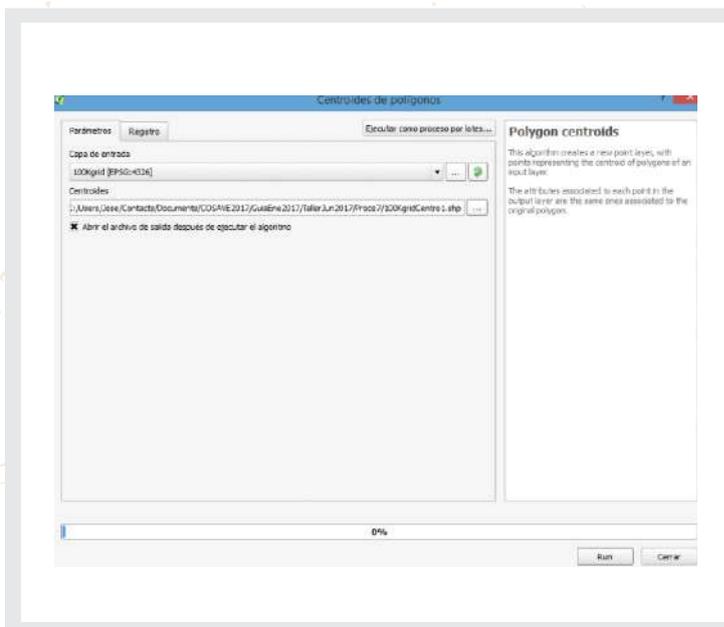


Fig. 35. Details in of the use of the "Centroids" function ("Vector geometry").



Fig. 36. Results of the "Centroids" functions.

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C4.2. Converting a vector shapefile in a raster.

In order to integrate the host risk index, first convert the vector “*.shp” file to raster with a function in the tap “Raster”, “Conversions” and then “Rasterize (vector to raster)”. To do this, use the column with the host risk index for the value of the raster. The illustration of this function is shown in the figure below.

For a better view of model raster values, with a right click on the layer, open the “Properties” then “Style”, “Singleband pseudocolor” with the “Spectral” option. Include the modification of the label as HIGH, MEDIUM or LOW, as illustrated in the figure below.

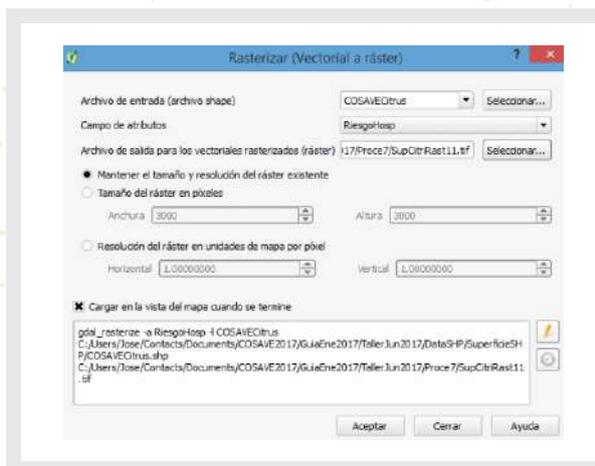


Fig. 41. With the “Rasterize” function, convert the “CSP_FA001_P.shp” file using the host risk index as a value.

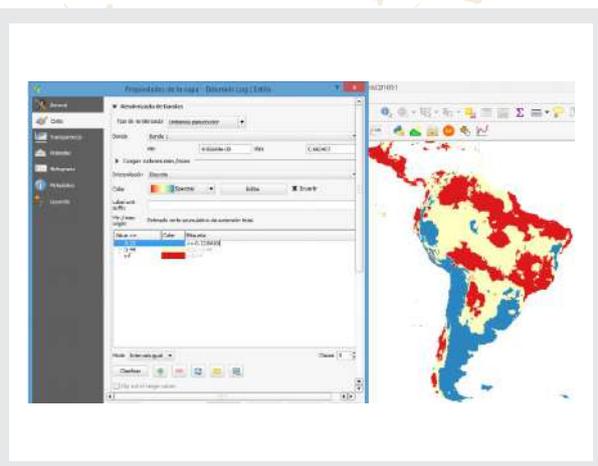


Fig. 42. For a better presentation of the bioclimatic risk model, change the properties with a right click on the layer.

C4.3. Reclassifying the raster of the model for its integration.

Similarly to the host risk index, index 0, 1 or 2 is required in the environmental modeling. For this purpose, use the “Reclassify values (simple)” function from SAGA (System for Automated Geoscientific Analysis) in the “Processing toolbox” on the right side of QGIS. In the function, select the “Fixed Table 3x3” option with the values 0, 1 or 2. These details are illustrated in the figures below.

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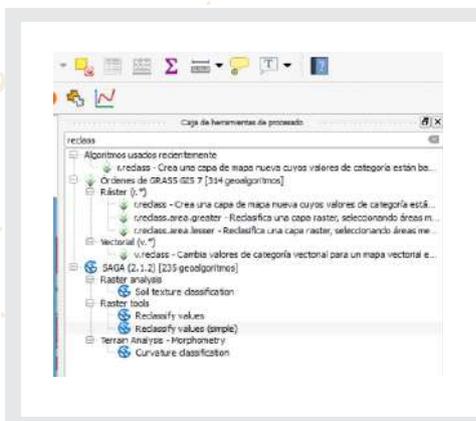


Fig. 43. Location of the “Reclassify values (simple)” function in SAGA.

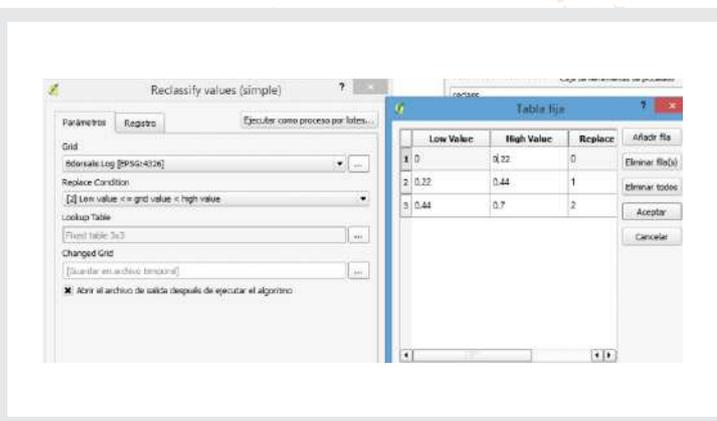


Fig. 44. Details of the “Reclassify values” function and the “Fix table 3x3”.

Moreover, it is possible to integrate the environmental risk with the host risk index or others with the “Raster calculator” function, which develops math between raster layers. With this tool, you can also integrate information on land cover or Normalized Vegetation Index (NDVI), which are raster files too. When raster layers are multiplied, the grids with a lower value are integrated with the zero (0) value, while higher categories are integrated with maximum values, as shown in Table 3.

Table 3. Integrating raster files with the reclassification of the value in the grid and the multiplication function in the “Raster calculator”.

RASTER INTEGRATION	LOW INDEX B = 0	MEDIUM INDEX B = 1	HIGH INDEX B = 2
LOW INDEX A = 0	LOW = 0	LOW = 0	LOW = 0
MEDIUM INDEX A = 1	LOW = 0	MEDIUM = 1	MEDIUM = 2
HIGH INDEX A = 2	LOW = 0	MEDIUM = 2	HIGH = 4

Source: Prepared for STDF/PG/502 COSAVE Project.

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The figure below illustrates the use and results of the "Raster calculator" function:

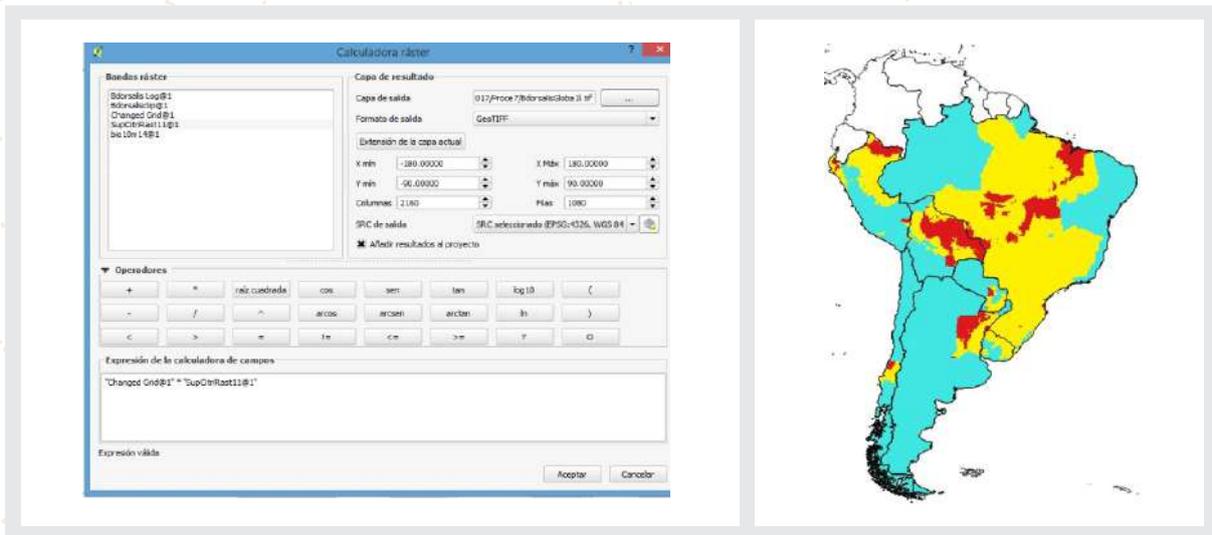


Fig. 45. Details in the "Raster calculator" function for the integration of raster layers.

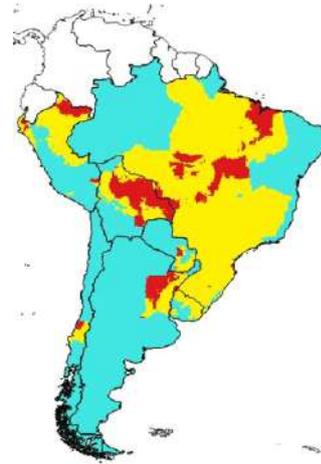


Fig. 46. Regional Risk map. Higher risk in red, medium risk in yellow and lower risk in light blue. Scale 1:45,000,000.

C4.4. Obtaining risk values with a vector shapefile

Open the centroid layer developed in section C3 to extract risk values to a vector (point) layer.

To include geographical coordinates, make a right click on the vector file and select "Open attributes table". Then use the "Field calculator" as shown with the red arrow in the figure below and the "Geometry" option selecting \$x for the field "longitude" and \$y for the field "latitude". It is important to check that the format of the field is "Decimal number (real)" and with 5 decimals of precision.

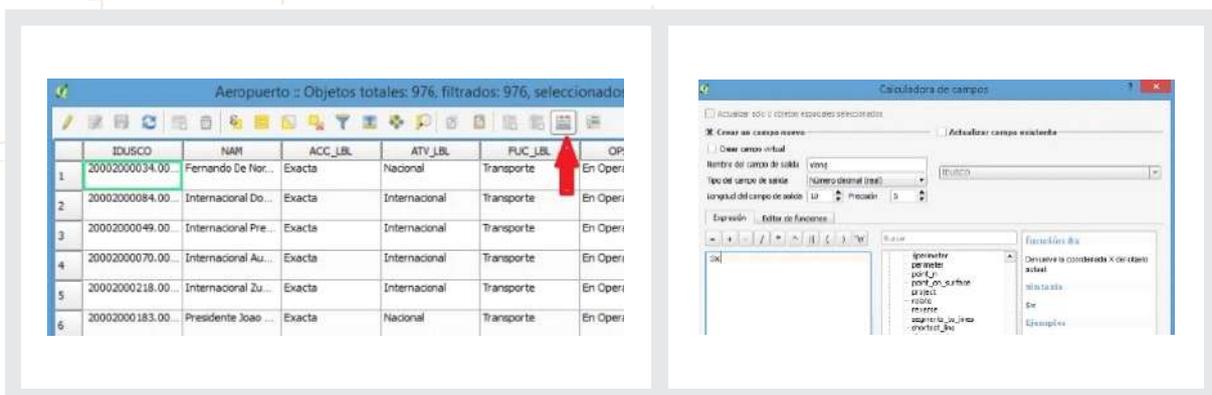


Fig. 47. With a right click on the vector shapefile, open the "Attributes table" and the "Field calculator", as indicated with the red arrow.

Fig. 48. Create a field called "longitude" as "Decimal number (real)" with 5 decimals and the "Geometry" function, then "\$x" and similarly with "\$y" for the "latitude".

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In the “Processing toolbox” on the right side, find the “v.sample” function and obtain raster values from the results of sections C2 or C4.3, as illustrated in the figures below.

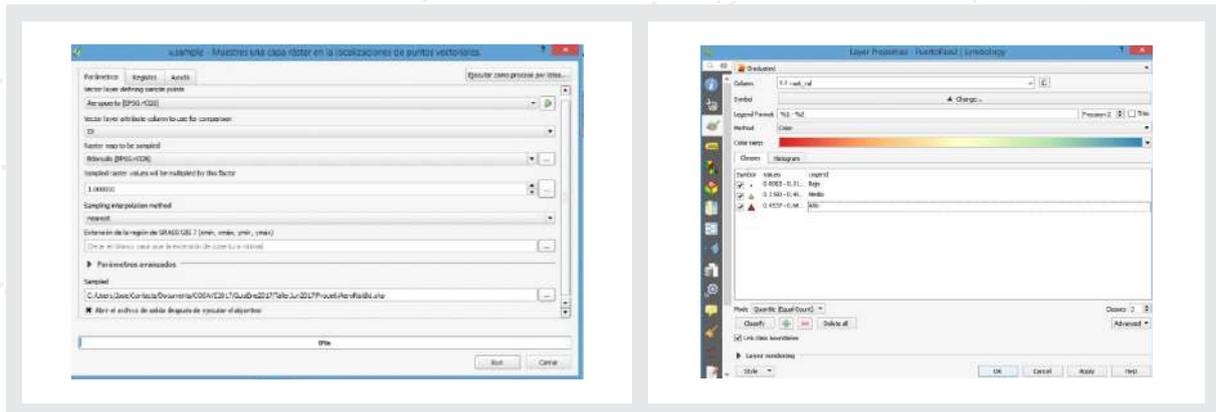


Fig. 49. Sampling risk values with a vector shapefile in a new file “Sampled” in a new route for the file.

Fig. 50. The developed shapefile has points and values that can be graphically shown, changing layer properties.

This layer “Sampled” can be integrated with another vector shapefile, like the 100 km grid that created the centroids. For this, find the “Join attributes by location” function in the “Processing toolbox”. Select the “Sampled” layers and the “Grid” shapefile and save it, as illustrated in the figure below.

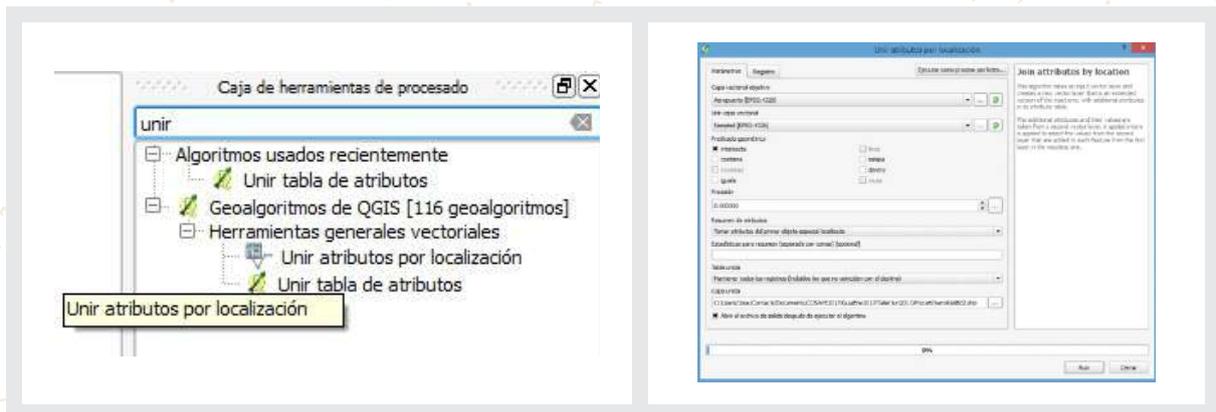


Fig. 51. Find the “Join attributes by location” function in order to integrate vector shapefiles.

Fig. 52. Select the layers to integrate with the “intercept” option and name this integrated layer.

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This integrated information can be shown in a map, selecting with a right click the "Layer properties", "Style" and "Categorize" in reference to the values of the modeling, as shown in the figures below.

It is also feasible to integrate other geo-referenced information, like the river layer "CSP_BH140_L.shp", available in <https://goo.gl/WYFe6a> using the "Intersection" function in the "Vector" tab and "Geoprocessing tools" group.

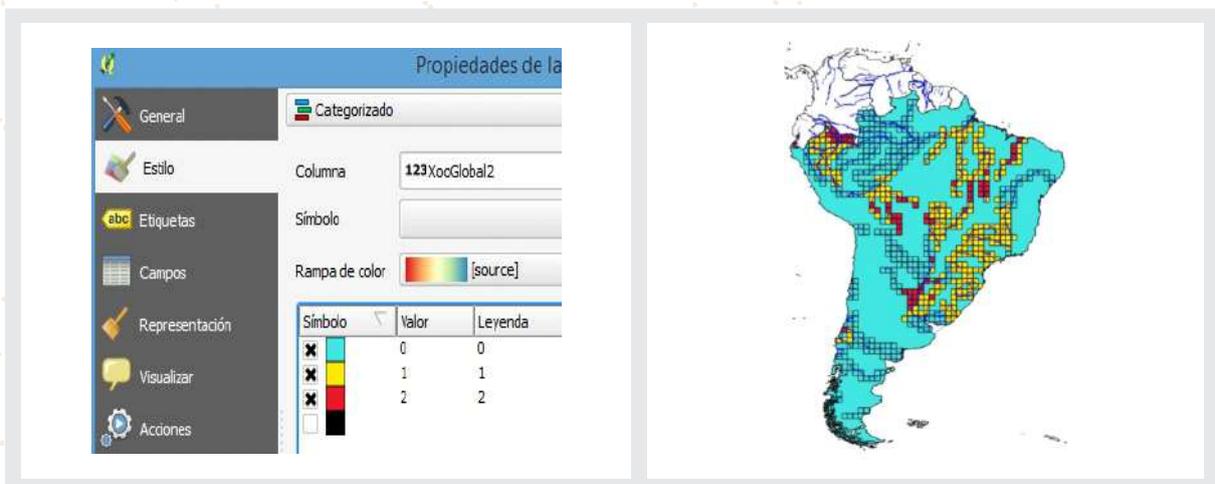


Fig. 53. With the "Layer properties" function you can change the style display of the 100 km grid for risk value (high risk in red, medium risk in yellow and low risk in light blue).

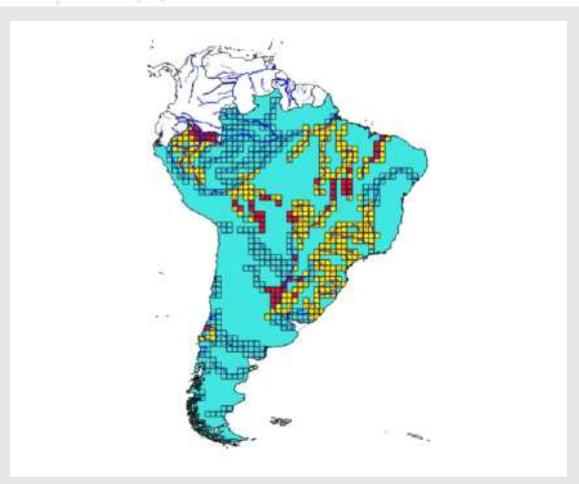


Fig. 54. With the tool "Intersection", located in the "Vector" tab, "Geoprocessing tools", you can integrate vector shapefile information from the river layers.

The vector shapefile resulting layer contains origin and modeled risk information. This information can be exported with the right click on the layer, as illustrated in the figures below:

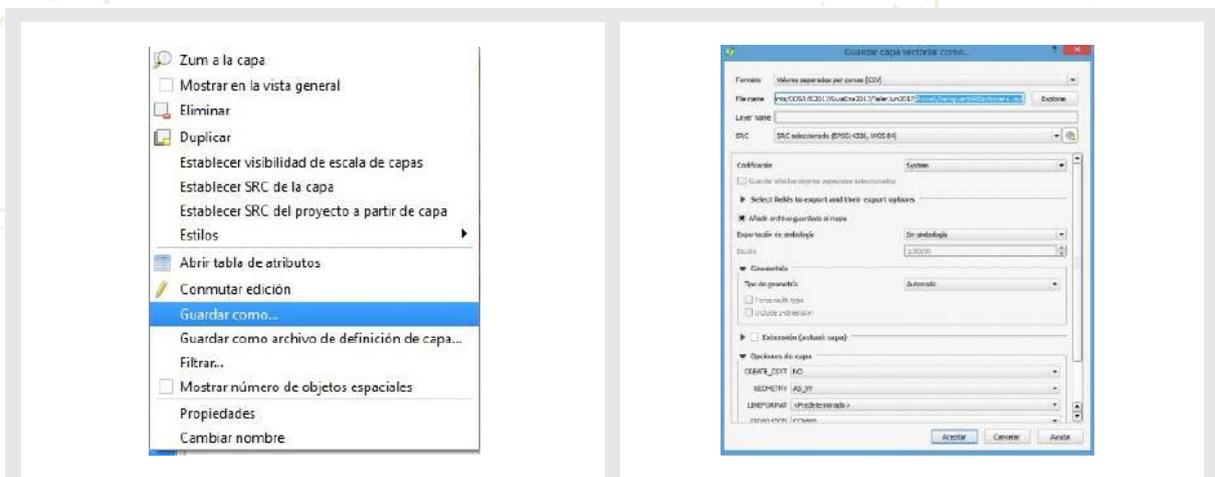


Fig. 55. To collect the information in another format, you can select the layer with a right click and choose "Save as".

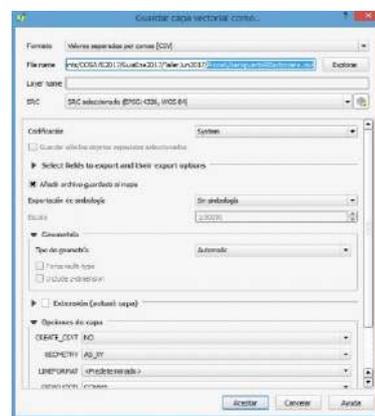
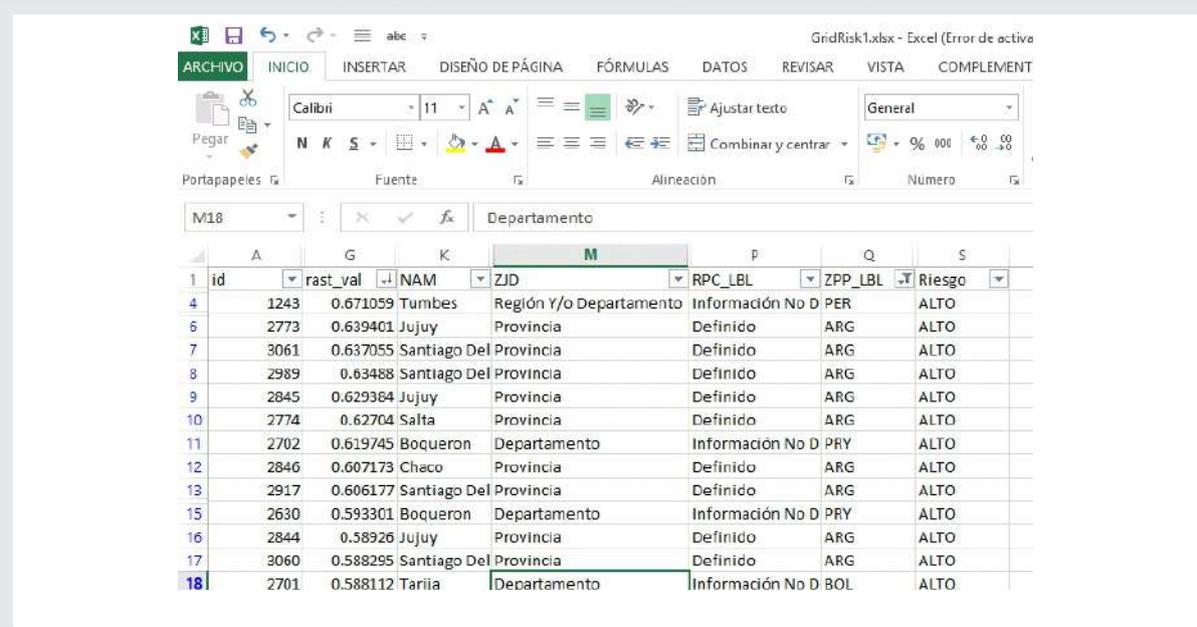


Fig. 56. Save the information in CSV ("Comma separated values") format, which can be opened in Microsoft Excel.

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In Microsoft Excel, the NPPO can assess the risk parameters in based on expert criteria. The figure below provides an example.



id	rast_val	NAM	ZJD	RPC_LBL	ZPP_LBL	Riesgo	
4	1243	0.671059	Tumbes	Región Y/o Departamento	Información No D PER	ALTO	
6	2773	0.639401	Jujuy	Provincia	Definido	ARG	ALTO
7	3061	0.637055	Santiago Del	Provincia	Definido	ARG	ALTO
8	2989	0.63488	Santiago Del	Provincia	Definido	ARG	ALTO
9	2845	0.629384	Jujuy	Provincia	Definido	ARG	ALTO
10	2774	0.62704	Salta	Provincia	Definido	ARG	ALTO
11	2702	0.619745	Boqueron	Departamento	Información No D PRY	ALTO	
12	2846	0.607173	Chaco	Provincia	Definido	ARG	ALTO
13	2917	0.606177	Santiago Del	Provincia	Definido	ARG	ALTO
15	2630	0.593301	Boqueron	Departamento	Información No D PRY	ALTO	
16	2844	0.58926	Jujuy	Provincia	Definido	ARG	ALTO
17	3060	0.588295	Santiago Del	Provincia	Definido	ARG	ALTO
18	2701	0.588112	Tariia	Departamento	Información No D BOL	ALTO	

Fig. 57. Raster risk value and the identification ID of each generated grid.

NOTE: All this case study information is available in the cXanthomonasoryzaeoryzae folder in the link: <https://goo.gl/WYFe6a>.

NOTE: The QGIS generated map, "Bdorsalis.qgs", is available in the "QBdorsalis" folder in the link: <https://goo.gl/WYFe6a>. To use it, download the "QBdorsalis" folder containing all the involved layers.



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Purdue University. 2017. Bacterial blight - *Xanthomonas oryzae* pv. *oryzae*. Available on October 27, 2017, at: <http://pest.ceris.purdue.edu/services/approvedmethods/sheet.php?v=681>
<http://download.ceris.purdue.edu/file/3055>
<http://pest.ceris.purdue.edu/pest.php?code=FBZAXBM>



GUIDELINES
FOR THE
IMPLEMENTATION
**OF THE SPECIFIC
PHYTOSANITARY
SURVEILLANCE SYSTEM**
CASE STUDY
***Xanthomonas
oryzae pv. oryzae***

