



marxan.io user guide

A web app for systematic conservation planning

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About marxan.io

Marxan is the most widely used decision support software for conservation planning globally. It's used in over 180 countries to build marine and terrestrial conservation plans and national parks systems. It's the global leader in conservation land-use planning software and new extensions are making it even more popular. URL http://marxan.org

What is marxan.io?

The marxan.io web app is a graphical user interface for Marxan users that runs on the Nectar research cloud. You can upload your own Marxan datasets, edit targets, SPF and BLM, conduct parameter testing and analysis, and visualise output maps, figures, and tables in an easy to use web interface.

The instructions relate to revision 34 of the marxan.io web app.

Get an account

We don't yet have an automated account management system. It is planned to integrate the app with Australian Access Federation (AAF) so you can use your federated access credentials from Australian Universities and research institutions.

Currently, you need to send an email to m.watts@uq.edu.au to receive your access credentials. It's not an automated process so please be patient and we'll endeavor to get back to you as quickly as we can.

The open source project

The app is written in R. It's open source software that you can freely use and modify subject to the conditions of the AGPLv3 open source software license.

You can access the source code on Github here:

https://github.com/mattwatts/marxan.io

You're welcome to participate in the evolution of the software by contributing changes to the Github repository. Lots more changes and improvements are planned.

Preparing your data

To use your own data with marxan.io, you first need to prepare your dataset and ensure Marxan is running ok on your local computer. If Marxan doesn't run ok, there is likely something wrong with your data. That is, one or more files malformed and not in compliance with the specification as detailed in the Marxan user manuals and course manuals.

There are functions in marxan.io that check for common mistakes in data creation and attempt to fix them, although some mistakes might stop your data from importing. More improvements to the error checking and correction code are planned.

Introduction to Marxan course materials

Day 1 of the Introduction to Marxan course materials describes in detail how to create a Marxan dataset from your own GIS layers. Handbooks are provided with the materials explaining this procedure using an open source GIS (Quantum GIS) and a commercial GIS (ESRI ArcMap). URL http://marxan.net, click "Teaching and Learning", "Course Materials".

Qmarxan plugin for Quantum GIS

An easy way to create a Marxan dataset is to use the Qmarxan plugin for Quantum GIS, described in the introduction to Marxan day 1 course materials.

Marxan workflow in R

We've created an entire workflow in R to create Marxan datasets that is available here:

https://github.com/mattwatts/CoESRA-Marxan

Other ways to create a Marxan dataset

There are many ways users create Marxan datasets and the methods multiply as time goes by. GRASS is a great way to intersect raster layers to create input files. Similarly PostGIS is a great way to intersect vector layers to create input files. You should choose the method that works best for you.

Creating a zip file

Once you have a Marxan dataset, assemble all the required files to prepare them for upload. You need these files:

- The input.dat input parameter file,
- The input folder with all the input files,
- The planning unit layer shapefile files.

The planning unit shapefile must have a planning unit id field called either: PUID, PU_ID, puid, or pu_id.

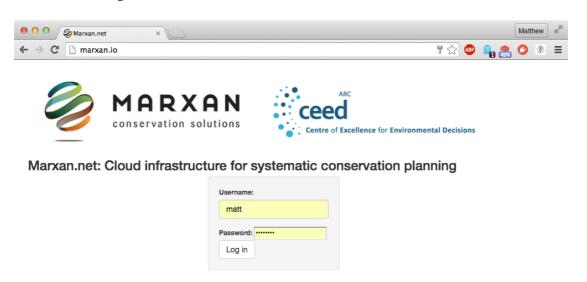
Put all the files in a directory with the name you want for your Marxan database then zip up the directory. This zip file is your Marxan dataset.

Sample datasets

A sample dataset "Tasmania" is included in your account by default. Several additional sample datasets are provided for download from http://marxan.net

Login to marxan.io

We recommend Google Chrome web browser if other browsers don't work as expected with the apps. Using your browser, visit http://marxan.net and click "marxan.io – login"



Enter the username and password provided. Then click the "Log in" button to be authenticated on the system. Once you're authenticated, you'll see this screen:

Login from a82-83.nat.uq.edu.au
Last login Mon Apr 4 08:30:37 2016 from a8283.nat.uq.edu.au
marxan.io user guide
Run Marxan
Parameter testing
Upload a dataset

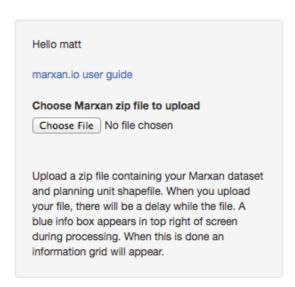
Click the blue "Run Marxan" link to run Marxan, and visualise output maps and figures. A new tab will open in your browser with this app leaving the authentication tab open where you can launch other apps.

Click the blue "Parameter testing" link to do parameter testing, and visualise output maps, figures, and tables. A new tab will open in your browser with this app leaving the authentication tab open where you can launch other apps.

Similarily, click the blue "Upload a dataset" link to upload your own Marxan datasets. Again, a new tab will open in your browser with this app leaving the authentication tab open where you can launch other apps.

Upload a dataset

The "Upload a dataset" screen looks like this:



Choose File

Click the "Choose File" button to select and upload the zip file containing your dataset. When you upload your file, there will be a delay while the file is processed and your data is extracted and ingested.

While processing occurs, a blue progress messages are displayed in the top right corner of the screen like this one:

```
Please wait: processing dataset

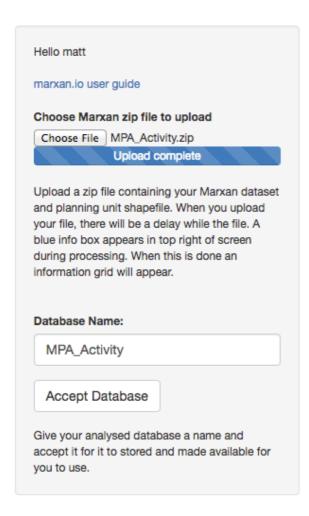
Planning units simplifying shapefile
```

For large datasets, the delay might be a minute or even several minutes while the files are analysed. Large and complex datasets can take a while to process.

The progress messages displayed are:

```
Please wait: processing dataset
Reading zip
      Copying
       Unzipping
Input files
       converting Marxan files
       converting MarZone files
       reading Marxan files
       reading MarZone files
       writing Marxan files
       converting matrix file
Planning units
      reading shapefile
      writing shapefile
       reading dbf table
       querying dbf table
       writing dbf table
       dissolving shapefile
       creating shapefile outlines
       converting shapefile
       saving pulayer.Rdata
```

When this is done, an information grid is presented to you like the screen below:



	V1	V2
1	name	MPA_Activity.zip
2	size	19749 bytes
3	elapsed	2.3480000000001 seconds
4	type	application/zip
5	Warnings	0
6	Errors	0

If the the blue progress messages disappear and the information grid does not appear, an error might have occurred in the processing of your dataset. You can email m.watts@uq.edu.au to check your data that is logged in the system.

If an error is listed in the "Errors" box, it will contain diagnostic information that might help you correct the error and upload a corrected zip file. If you get stuck you can email m.watts@uq.edu.au to check your data that is logged in the system.

Diagnosis of errors will allow us to improve the service over time for all users, so your patience and help is appreciated.

Database Name

If the information grid is displayed with summary information on the examination of your dataset, and no errors are displayed, then you can specify a name for your dataset.

Database Name:	
MPA_Activity	

Please only use alphabetic and numeric characters with underscores and no spaces, as this becomes the directory name for your dataset.

Accept Database

To accept a dataset that has been successfully analysed, click the "Accept Database" button.

Accept Database

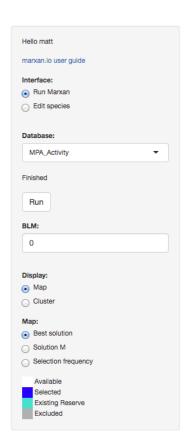
Your dataset will then be imported into your account for use. A blue message will briefly appear on the top right corner of the screen screen saying "Dataset accepted".

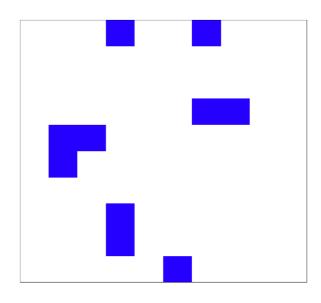
If you already have a dataset with the same name, a blue message will briefly appear on the top right corner of the screen saying "Duplicate dataset name". In this case, enter a new unique name before pressing "Accept Database".

Run Marxan

The "Run Marxan" web app is a graphical user interface that allows users to run Marxan, display maps, tables and graphs, and edit key parameters. It has a control panel on the left, and an output panel on the right.

The "Run Marxan" screen looks like this:





User Interface

The app has an "Interface" control:



Select "Run Marxan" to set the BLM (boundary length modifier), display output maps and cluster analysis graphs, and run Marxan.

Select "Edit Species" to set the feature prop (proportional target) and SPF (species penalty factor) for each of your species.

Database

The app has a "Database" control:



Click the drop down arrow on the right of the control to select the dataset you want to work with.

The first time you select a dataset you have uploaded, there will be a delay while the Marxan outputs are rendered. Blue progress messages appear in the top right corner of the screen while processing occurs.

This delay is a minute or so for small datasets and several minutes for large and complex datasets.

Please be patient after selecting a dataset while the output renders. The blue progress messages disappear and the display refreshes with output from your dataset once it's rendered.

BLM

The app has a "BLM" control:



To edit the boundary length modifier (BLM), enter a new value in this control.

Run

The app has a run button:

Run

When you've changed BLM, SPF, or targets, click this button to run Marxan with the new parameters.

A delay of a few seconds for small datasets and longer for larger datasets occurs. During processing, progress messages appear in the top right corner of the screen and the "Finished" label greys out. After the results are rendered, the blue progress messages disappear, the "Finished" label appears black again and the display refreshes with new output.

Finished

Display output: Map

The app has a "Display output" control:



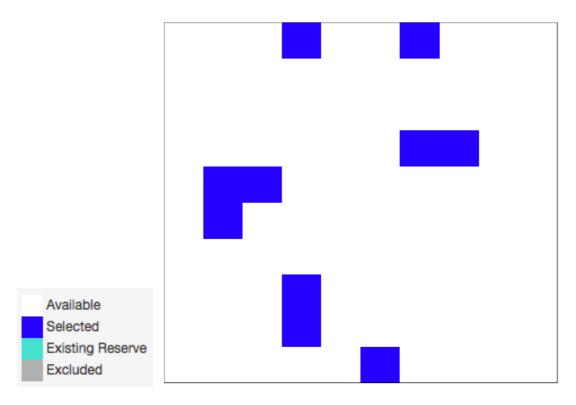
Select "Map" to display output maps.

When "Map" is selected, a "Map to display" control appears:



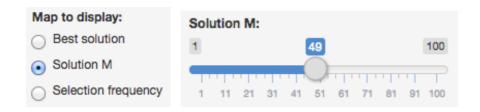
Best solution

Selecting "Best solution" displays the Marxan solution with the lowest objective function score. A map and map legend appears:



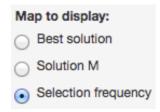
Solution M

Selecting "Solution M" displays a slider to select any of the 100 solutions:

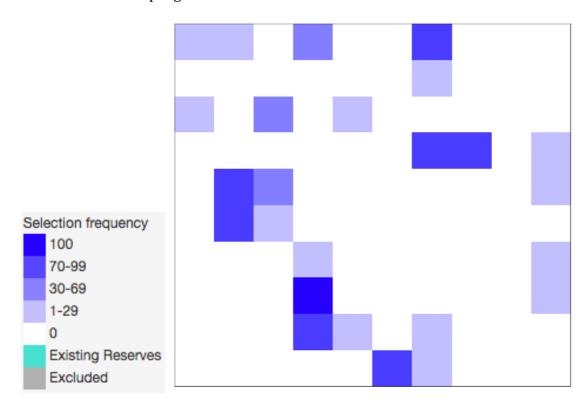


When you drag the slider, the map display is refreshed with the solution selected in the slider.

Selection frequency



Selecting "Selection frequency" displays a selection frequency map of 100 solutions and a map legend:



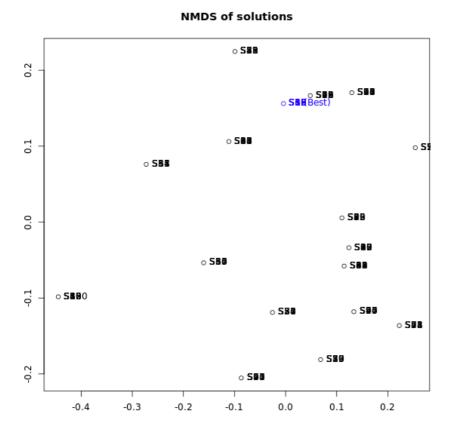
Display output: Cluster

Selecting "Cluster" with the "Display output" control causes the "Cluster to display" control to appear:



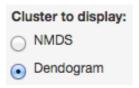
NMDS of solutions

Selecting "NMDS" with the "Cluster to display" control displays the 2 dimensional scaling of the dissimilarity of the 100 solutions:



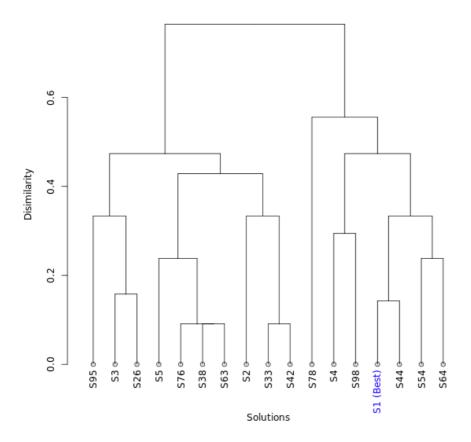
The 100 solutions are labeled on the graph, and the solution with the lowest objective function score is labeled in blue.

Bray-Curtis dissimilarity of solutions



Selecting "Dendogram" with the "Cluster to display" control displays a dendogram of the 100 solutions:

Bray-Curtis dissimilarity of solutions



Only unique solutions are displayed, and again each solution is labeled and the solution with the lowest objective function score is labeled in blue.

Edit Species

Selecting "Edit Species" in the "Interface" control causes the species table editor to appear:



Edit "prop" and "spf"

Here you can edit prop (proportional target) and SPF (species penalty factor) for each species.

	id	prop	spf
1	1	0.1	10
2	2	0.1	10
3	3	0.1	10
4	4	0.1	10
5	5	0.1	10
6	6	0.1	10
7	7	0.1	10
8	8	0.1	10
9	9	0.1	10
10	10	0.1	10

To edit prop or SPF for a species, click the cell you want to edit and type in a new value. Here we have changed the prop for species 2 from 0.1 to 0.4:

	id	prop	spf
1	1	0.1	10
2	2	0.4	10
3	3	0.1	10
4	4	0.1	10
5	5	0.1	10
6	6	0.1	10
7	7	0.1	10
8	8	0.1	10
9	9	0.1	10
10	10	0.1	10

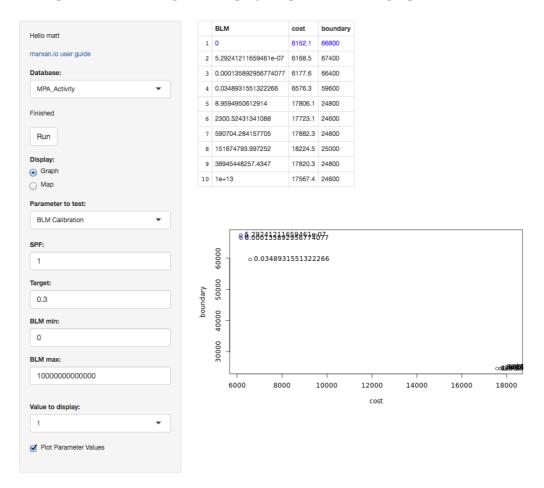
Save

When you have entered the prop and SPF values for the species you want to change, click the "Save" button to save the changes to your dataset:



Parameter Testing

The "Parameter testing" web app is a graphical user interface that allows users to do parameter testing, and display maps, tables and graphs:



Here you can do calibration and sensitivity analysis of key parameters. You can iteratively run tests for parameters of interest, changing parameters each time, to interactively explore the response of variables of interest. This allows the dynamics of a problem to be systematically explored in a structured way.

Appropriate values for the key parameters can thus be rapidly and efficiently determined.

Database

The app has a "Database" control:



Click the drop down arrow on the right of the control to select the dataset you want to work with.

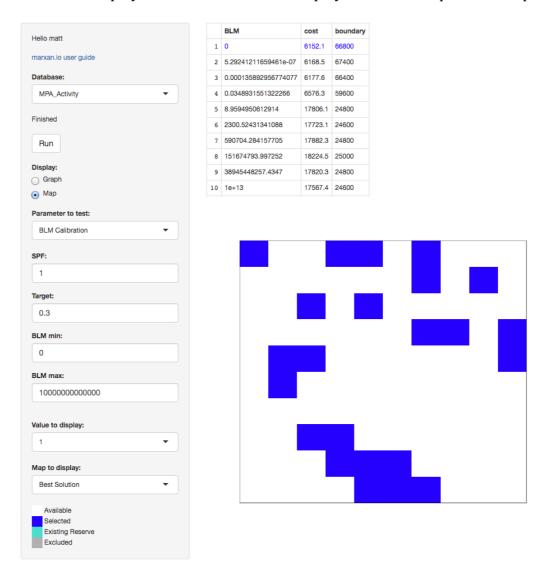
The first time you select a dataset you have uploaded, there will be a delay while the parameter testing outputs for the selected test are rendered. Blue progress messages appear in the top right corner of the screen while processing occurs.

This delay is a minute or so for small datasets and several minutes for large and complex datasets.

Please be patient after selecting a dataset while the output renders. The blue progress messages disappear and the display refreshes with output from your dataset once it's rendered.

Display

Click the "Display" control to switch the display between "Graph" and "Map":



Parameter to test

The "Parameter to test" control selects the test you want to run: "BLM Calibration", "SPF Calibration", or "Target Sensitivity":



The first time you select a parameter to test for a dataset you have uploaded, there will be a delay while the parameter testing outputs for the selected test are rendered. Blue progress messages appear in the top right corner of the screen while processing occurs.

This delay is a minute or so for small datasets and several minutes for large and complex datasets.

Please be patient after selecting a parameter while the output renders. The blue progress messages disappear and the display refreshes with output from your dataset once it's rendered.

BLM Calibration

Click the drop down arrow on the right of the "Parameter to test" control and select "BLM Calibration" to calibrate the boundary length modifier (BLM):



These controls appear for you to enter parameter values for the calibration:



In the "SPF" control, enter the SPF value you want to use.

In the "Target" control, enter the proportional target value to use.

In the "BLM min" control, enter the minimum BLM value to use.

In the "BLM max" control, enter the maximum BLM value to use.

SPF Calibration

Click the drop down arrow on the right of the "Parameter to test" control and select "SPF Calibration" to calibrate the species penalty factor (SPF):



These controls appear for you to enter parameter values for the calibration:

BLM:	
0	
Target:	
0.3	
SPF min:	
0.0001	
SPF max:	
10000000000	000

In the "BLM" control, enter the BLM value you want to use.

In the "Target" control, enter the proportional target value to use.

In the "SPF min" control, enter the minimum SPF value to use.

In the "SPF max" control, enter the maximum SPF value to use.

Target Sensitivity

Click the drop down arrow on the right of the "Parameter to test" control and select "Target Sensitivity" to analyse the sensitivity of the species targets:



These controls appear for you to enter parameter values for the calibration:

BLM:	
0	
SPF:	
1	
Target min:	
iaiget iiiii.	
0	

In the "BLM" control, enter the BLM value you want to use.

In the "SPF" control, enter the proportional SPF value to use.

In the "Target min" control, enter the minimum Target value to use.

In the "Target max" control, enter the maximum Target value to use.

Run

Click the run button to execute the selected parameter test with the parameters you have entered:



A delay of a few seconds for small datasets and several minutes for larger datasets occurs during which the "Finished" label greys out. Blue progress messages appear in the top right corner of the screen while processing occurs.

When processing is finished, the blue progress messages disappear, the "Finished" label appears black again, and the display refreshes with new output.

Finished

A set of 10 values is tried for the parameter you are testing.

A set of 10 solutions are generated for each value you are testing for a total of 100 solutions.

For BLM and SPF calibration, the values are ramped exponentially between the min and max entered.

For Target sensitivity analysis, the values are ramped linearly between the min and max entered.

Output table

	SPF	cost	shortfall
1	1e-04	0	4164.9
2	0.00774263682681128	0	4164.9
3	0.599484250318941	5063.8	313.7
4	46.4158883361278	7420.4	0.1
5	3593.81366380463	7307	0
6	278255.940220712	7526.7	0
7	21544346.9003189	7391.6	0
8	1668100537.20006	7417.2	0
9	129154966501.489	7615.6	0
10	1e+13	7792.6	0

The output table displays the parameter values tried, the average cost of solutions, and the average value for the response variable of the solutions.

For SPF, the response variable is "shortfall" (the total amount by which our features miss meeting their targets).

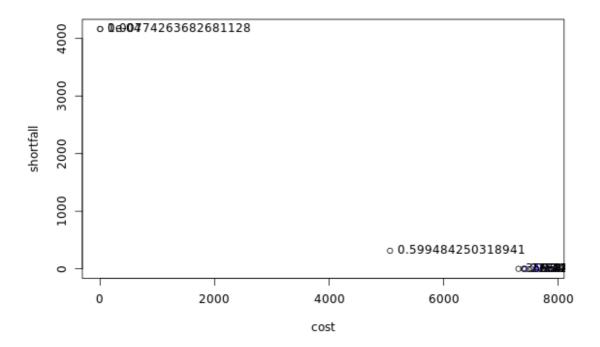
For BLM, the response variable is "boundary" (the total boundary length of the reserve network configuration).

For target, the response variable is simply "target" (the target level we are testing).

Our aim with SPF and BLM calibration is to discover a parameter value that gives an appropriate response in the response variable.

Our aim with target sensitivity analysis is to discover how much it costs to protect different proportions of our species.

Output graph



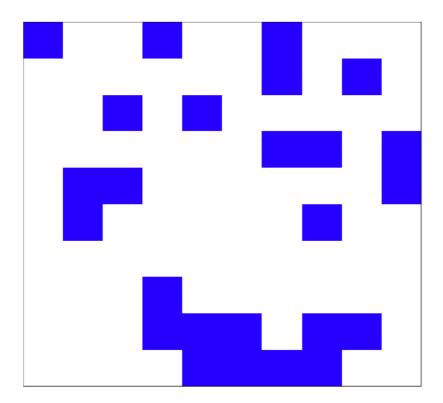
The output graph for BLM and SPF displays the response variable on the Y axis, and the cost on the X axis.

The output graph for target displays target on the Y axis and cost on the X axis.

As a test is rerun with different values of "min" and "max", results are cumulatively added to the graph so the problem dynamics can gradually be revealed.

This iterative process allows rapid and efficient determination of appropriate values for the key parameters.

Output map



The output map displays selected areas or selection frequency for the value selected in the "Values to display" control (see below).

The "Map to display" control also determines which map is displayed (see below).

Value to display

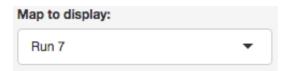


The "values to display" control allows you to select which input parameter value to interrogate.

Choosing "1" selects the "min" value specified. Choosing "10" selects the "max" value specified. Choosing a value between 1 and 10 selects one of the values ramped between the "min" and "max" values.

The selected value is highlighted in blue on the output table and graph, and results from testing the selected value are displayed on the output map.

Map to display



The "Map to display" control allows you to select which map to display.

10 solutions are generated for each input parameter value tried, and selecting "Run 1" to "Run 10" displays that solution on the map.

Selecting "Best Solution" displays the solution with the lowest objective function score on the map.

Selecting "Selection Frequency" displays selection frequency on the map.

Plot Parameter Values



The "Plot Parameter Values" control selects if parameter values will or will not be displayed next to their relevant points on the graph.

Acknowledgements

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