

# PAT - Precision Agriculture Tools Plugin for QGIS

**User Manual** 

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## **Enquiries**

Contact PAT@csiro.au for queries about the tools described in this manual.

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## PAT - Precision Agriculture Tools

The Precision Agriculture Tools (PAT) plugin is a suite of open source tools developed by CSIRO for Precision Agriculture (PA) data analysis. The tools run within Quantum Geographic Information System (QGIS) version 2.18 (QGIS Development Team 2017), a free and open-source desktop geographic information system that supports viewing, editing, and analysis of geospatial data. PAT aims to provide an easy-to-use interface for processing PA data through an established workflow developed for constructing maps using on-the-go data (e.g. from yield monitors or EM38 soil surveys) as shown in Figure 1 and Table 1 (Bramley and Williams 2001; Taylor et al. 2007; Bramley et al. 2008; Bramley and Jensen 2014). It also includes tools for analysis of remotely sensed imagery and some on farm experiments and for clustering map layers to create potential management zones. Over time more tools and 'how-to' instructions will be added to expand the functionality and usefulness for both practical and research purposes.

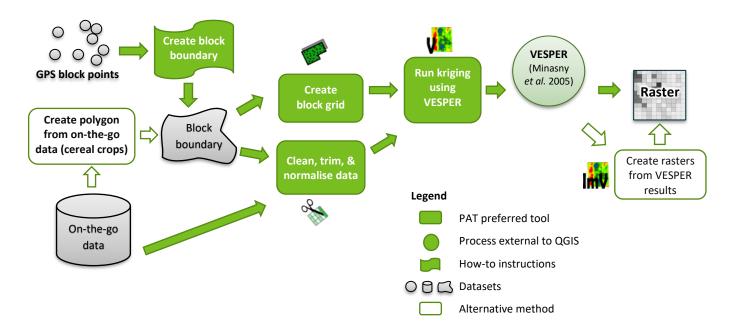


Figure 1 The basic map production workflow used in PAT and some of the existing tools available for processing high density on-the-go data.

Table 1 Brief description of tools available in PAT

	PAT TOOLS	DESCRIPTION
	TAITOOL	Convert polygon features (e.g. a block boundary – see page 57) to
	Create Block Grid	<ul> <li>a TIFF raster; and</li> <li>a VESPER text grid file of X,Y point values used by VESPER.</li> </ul>
	Clean, Trim, Normalise Point Data	Process an on-the-go data file (e.g. from yield monitors) by applying clipping, cleaning and filtering rules and output as a CSV file.
<b>V</b>	Run Kriging Using VESPER	Create a VESPER control file and data files and run VESPER kriging.
ImV	Import VESPER Results	Convert VESPER outputs to raster TIFF format.
3. 0	Create Polygon from	Generate a polygon block boundary from on-the-go data (e.g. from yield monitors) containing GPS points.
	On-The-Go GPS Point Trail Data	The GPS points must cover the entire block (not just be a set of points around the boundary - see page 57 for instructions on creating a boundary from a set of surveyed data points).
	Rescale or Normalise a Raster	<ul> <li>Create rasters by</li> <li>rescaling (standardising) values between a fixed range (i.e. 0-1, or 0-255); or</li> <li>normalising values to a mean of 0 and standard deviation of 1.</li> </ul>
<b>4</b>	Generate Random Pixel Selection	Generate a selection of random pixels from a raster and save to a points Shapefile.
<b>&amp;</b>	Extract Pixel Statistics for Points	Extract pixel statistics using a square neighbourhood footprint from multiple rasters at set locations.
	Calculate Image Indices for Blocks	Resample and smooth imagery to a larger pixel size, as well as calculate indices such as PCD and NDVI.
	Resample Image Band for Blocks	Clip imagery to one or more blocks and resample to a different pixel size.
	Create Zones with k- means Cluster	Create zones with <i>k</i> -means clustering.
		Tool for analysing simple strip-based trials.
	Create Strip Trial Points	It generates pairs of points from a central line within a strip and with 'a pair' offset at a distance outset the strip. These points can be used for the comparison of performance along and adjacent to a strip trial (such as an N-rich strip) using the Lawes and Bramley (2012) moving window t-test tool.
4	Run Strip Trial <i>t</i> -test Analysis	The comparison of performance of a treatment, such as different fertiliser rates, along and adjacent to a strip trial (such as an N-rich strip) using the moving window t test as described in Lawes and Bramley (2012).
	Whole-of-block Analysis	Analyse whole-of-block experiments where 2 or 3 treatments are applied across a block or paddock.
<b>%</b>	Persistor	Summarise the patterns of yield across a block over a number of years relative to a target, as described by Bramley and Hamilton (2005).
<b>P</b> S	Apply Raster Symbology	Apply predefined symbology to a displayed raster.
0	Settings	User settings for PAT. Provides the ability to set data directories and the location of VESPER (if installed).
$\mathcal{L}$	About	About Precision Agriculture Tools (PAT). This provides information on the currently installed release, and relevant open source licences.
?	Help	Display the PAT user manual.

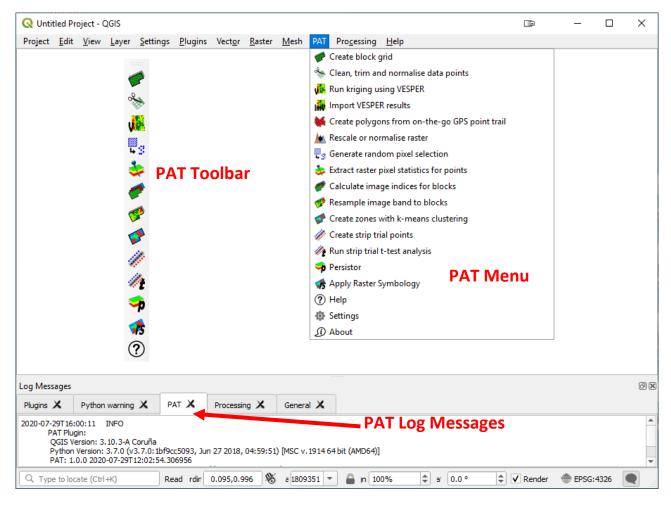


Figure 2 PAT toolbar, menu and log messages panel within QGIS.

# 1 Installing, Upgrading and Uninstalling

#### 1.1 License

The PAT plugin for QGIS is free to download and use under a GPL V3 Open Source license. PAT license details can also be found in the **About** tool under PAT menu. The companion Python library 'pyprecag' (Ratcliff et al. 2019) is also freely available and is released under the CSIRO Open Source Software License Agreement (variation of the BSD / MIT License). All source code and license files are available from https://github.com/CSIRO-Precision-Agriculture.

## 1.2 Requirements

QGIS 3.x: Download and install from the Official QGIS Site.

VESPER: To use the Run Kriging Using VESPER tool requires installation of VESPER (note: only available for Microsoft Windows operating systems). VESPER is a kriging (map interpolation) program (Minasny et al. 2005). It needs to be installed independently if you would like to undertake kriging of maps using VESPER. VESPER is **not** distributed with this plugin but is the recommended map interpolation tool. To download or view more information on VESPER visit https://precisionagriculture.sydney.edu.au/resources/software/download-vesper.

R 3.5.1+: The R statistical language (R Core Team 2018) is required to carry out the analytical processing for the Whole-of-block Analysis (Section 2.14). R is not distributed with this plugin. If Whole-of-block analysis is required, then you must download and install R in addition to QGIS. The minimum R version suitable for this is v3.5.1 Configuring QGIS to use R section on page 10.

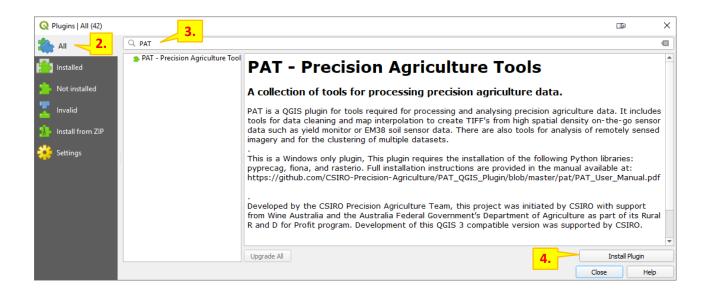
Additional Python Packages: QGIS includes Python and numerous Python packages. However, PAT requires the following additional packages: pyprecag (Ratcliff et al. 2019), Fiona (Gilles et al. 2011), Rasterio (Gilles et al. 2013) and GeoPandas (Jordahl et al. 2020). The Fiona, Rasterio and Geopandas packages have been bundled with the OSGeo4W installer with the PAT plugin and instructions on how to install them are included.

## 1.3 Installing PAT

## 1.3.1 Install the PAT - Precision Agriculture Tools Plugin.

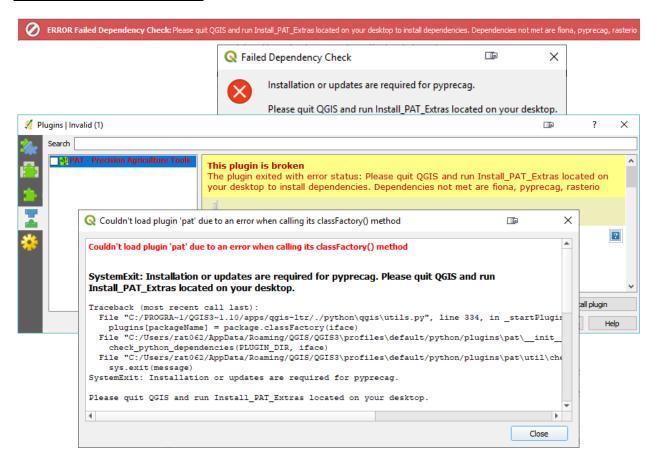
- 1. In QGIS open the plugin manager. Plugins Menu → Manage and Install Plugins
- 2. Select the **All** or **Not installed** side tab.
- 3. Search for and select PAT Precision Agriculture Tools.
- 4. Click *Install plugin*.

You will receive an Failed Dependency Check message box and Error loading plugin messages. This is expected and is resolved by completing the next section titled Installing or Upgrading PAT Python Dependencies.

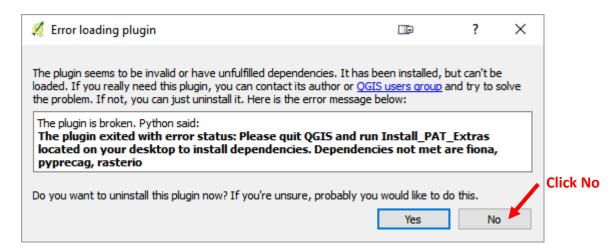


## 1.3.2 Installing or Upgrading PAT Python Dependencies

While installing or upgrading PAT, a check will be completed to ensure the required Python packages are installed on your system. If any of the images below appear, then this check has failed. This is normal and they may appear multiple times.



If the following message appears click No.



#### Steps to install dependencies.

- 1. Close QGIS and go to your windows desktop.
- 2. A shortcut has been created on your desktop named *Install PAT3 Extras*. Double click the shortcut to run it.



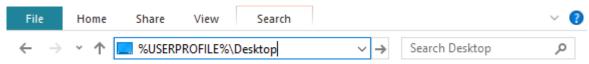
Extras

- 3. Choose YES for the Windows Account Control and any other messages which may appear.
- 4. Restart QGIS. The check will run again to ensure the installation occurred correctly.

If the shortcut does not appear on your desktop then try one of the following.

#### Using the shortcut

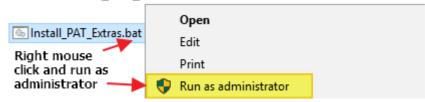
- a) Open windows explorer
- b) Add %USERPROFILE%\Desktop and hit enter.



- c) Double click the *Install PAT Extras* file in this folder.
- d) Continue from step 3 above

#### Running batch file as administrator.

- a) Follow steps a & b above but enter %AppData%\QGIS\QGIS3\profiles\default\python\plugins\pat\python\_packages into the address bar.
- b) Right click the *Install\_PAT\_Extras.bat* file and select *run as administrator*.



c) Continue from step 3 above

If the PAT menu and toolbar are not present, then reinstall and/or check/activate PAT using the plugin manager. QGIS should now contain the PAT menu, toolbar and log panel similar to Figure 2 on page 6. PAT has now been successfully installed.

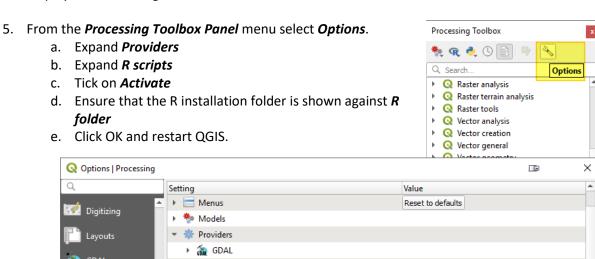
## 1.3.3 Configuring QGIS to use R

If R is not installed and you launch the *Whole-of-block analysis tools* via the menu or toolbar you will a warning message as shown.



To resolve this, complete the following steps to configure R and QGIS

- 1. Download R from Windows version can be downloaded from http://r-project.org/ . The minimum R version suitable for this is v3.5.1.
- 2. Install using all the default options
- 3. Using the QGIS plugin manager
  - a. ensure the **Processing** plugin is checked
  - b. install the *Processing R Provider*.
- 4. Display the *Processing Toolbox Panel*



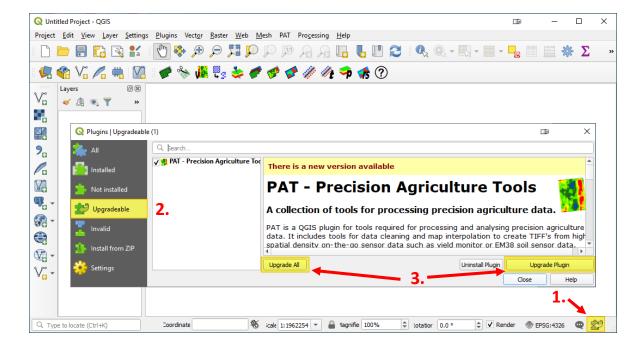
## 1.4 Update PAT

If the Check for Updates option is checked on in the plugin manager, QGIS will advise the user if new plugins are available, or if installed plugins have been updated. This notification is displayed in the QGIS interface's status bar as shown below. It is recommended that *Check for Updates* is checked.

## To Update

- 1. In QGIS open the Plugin Manager by clicking on the link in the status bar or via Plugins. Menu  $\rightarrow$  Manage and Install Plugins.
- 2. Either select the *Upgradeable* left side tab or search for your plugin.
- 3. Upgrade by selecting the PAT plugin and click Upgrade plugin.

A check will be run to ensure the Python packages required by PAT are installed and are of the correct version. If this check fails, follow the instructions in *Installing or Upgrading PAT Python Dependencies* to upgrade the dependencies.

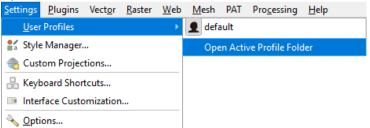


## 1.5 Uninstall PAT

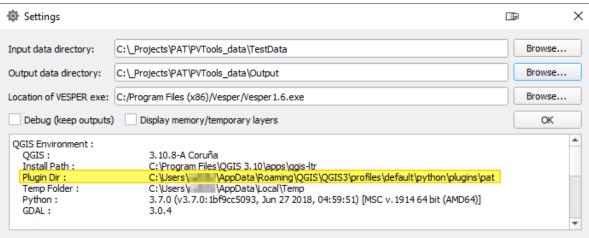
If you wish to uninstall PAT and its Python dependencies you must first uninstall the Python dependencies, followed by the plugin as instructed below.

## 1.5.1 Uninstall PAT Python dependencies for all users.

- 1. Navigate to the current user's PAT installation directory.
  - a) From the Settings -> User Profiles -> Open Active Profile Folder then continue navigating to python\plugins\pat



b) Navigating to the folder as listed in as **Plugin Dir** in the *settings dialog* as shown highlighted below.



- 2. Find the *Uninstall\_PAT\_Extras.bat* file in the python\_packages folder.
- 3. Right click the file and select *Run as Administrator* and choose *Yes* for any messages which appear.

## 1.5.2 Uninstall PAT Plugin.

PAT Python dependencies should be uninstalled <u>PRIOR</u> to uninstalling the PAT plugin. The plugin can be uninstalled in one of two ways.

#### 1. Via the QGIS Plugin Manager

- a. In QGIS, open the plugin manager, and find the PAT plugin.
- b. Click Uninstall plugin.

#### 2. Via Windows Explorer.

- a. Navigate to the current users PAT installation directory as show in 1.5.1.1 above.
- b. Delete the entire pat folder.

## 2 Individual Tools



# Create Block Grid

## **Summary**

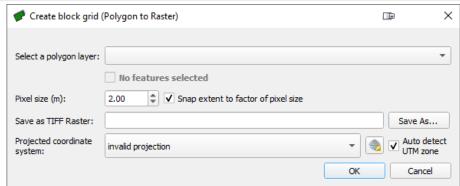
This tool converts polygon features such as a block boundary to a raster grid using a set pixel size. This is a critical part of setting up the analysis environment for creating maps from on-the-go data because it generates the base grid onto which maps will be interpolated, by using the outer most extent of the block boundary as the 'frame' for the grid. Refer to instructions on page 57 on how to generate a block boundary.

The raster outputs created by the Block Grid tool are:

- a TIFF raster; and
- a grid file of X, Y point values used by VESPER for kriging.

Areas inside the polygon boundary will be assigned a value of 1 while areas outside will be assigned a nodata value of -9999.

If your shapefile layer does not appear in the layer list, then it is probably in a geographic coordinate system. Refer to page 59 for information on why using a projected coordinate system is important, and how to reproject it. A future version of the Create Block Grid tool will include this conversion when required.



Block Grid (Polygon to Raster)					
Select a polygon layer		A layer containing the block boundary polygon features to be converted to the raster outputs.			
Use selected features	Default is unchecked	If checked, will only use the selected features, if unchecked, all the features will be used. Useful in creating a grid for a single field from a boundary layer comprising more than one field.			
Pixel size (m)	0.00m to 6 km Default is 2 m	The pixel size to assign to the raster outputs. This is expressed in metres.  Recommended Values:  Viticulture: 2 m  Sugar: 2 m  Broadacre grains:  < 200 ha 5 m  > 200 ha 10 m			
Snap extent to factor of pixel Size	Default is checked	Snap the output raster extent to a factor of the pixel size. This will ensure adjacent rasters use a common origin which is important for future analysis.			
Projected coordinate	e system	The output projected coordinate system of the TIFF raster.			
Auto detect WGS84 UTM zone		<ul> <li>If Auto detect is checked the following rules will apply:         <ul> <li>attempt to match the input if it is a projected coordinate system.</li> <li>If the inputs are in Geographic's (lat/long) it will calculate the relevant WGS84 UTM Zone.</li> </ul> </li> <li>Note: If in Australia then MGA GDA 1994 zones, will be used – see Technical Notes on page 56.</li> </ul>			
Save as TIFF raster  Default derived from input layer		The output raster TIFF file to be created. The VESPER grid file will have a suffix of _v.txt .			



## Clean, Trim, Normalise Point Data

## Summary

This tool processes on-the-go data files (e.g. from a yield monitor or EM38 soil survey) containing GPS coordinates recorded as latitude and longitude in decimal degrees, to output point values in a projected coordinate system. The tool also applies cleaning and filtering rules.

#### This tool:

- retains all columns (except coordinate columns) from the original file.
- converts coordinate columns to a projected coordinate system and renames them to Easting and Northing. An additional column (EN\_EPSG) will be created and assigned the EPSG number for the projected coordinate system used to reproject data. An EPSG number is an international numbering system for coordinate systems (see http://www.epsg.org).
- optionally saves a Shapefile version matching the output CSV file. A second Shapefile will also be saved containing the string '\_removed' in its filename. This Shapefile will contain all the points the filter discards and will be attributed by filter type. The description of the values can be found in Table 2.

## This tool filters data by:

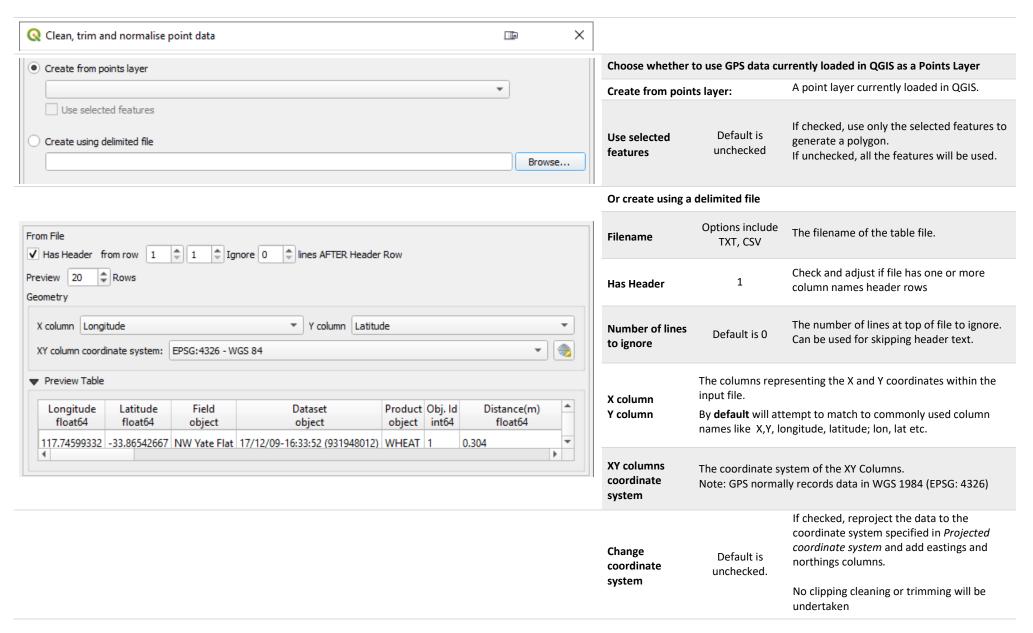
- optionally clipping the data to a Shapefile boundary (i.e. removes points that lie outside the
- optionally removing values from a given column which are null (missing) or are less than or equal to
- creating a normalised column using a prefix of nrm\_ and calculating normalised values for the data such that their mean is zero and standard deviation is 1. The normalised value of column Z is calculated as (Z - mean(col Z))/(s.d.(col Z)).
- optionally trimming normalised outliers based on a set number of standard deviations. This trim can optionally be performed iteratively with the normalised value re-calculated for each iteration. It is recommended that, by default, normalised data are trimmed to +/- 3 standard deviations.
- optionally thinning the data by removing points that are closer together than a set distance. For data that have been georeferenced using a differential GPS, we recommend removing points that are closer than 1 m apart.

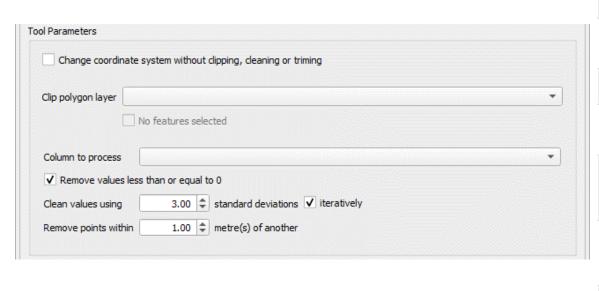
As part of the filtering process, the tool may rename some data columns to adhere to Shapefile column name limitations. The new names are displayed as a PAT log message and written to the log file. For more information on the location of the saved log file and temporary folder refer to the Technical Notes section on page 56.

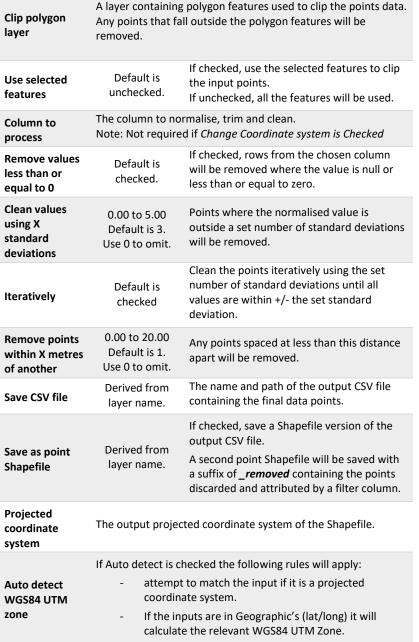
Once processing is complete, the results of filtering are shown as a PAT log message. Only filters which remove points are listed. When iteratively filtering, results of all iterations are shown. An explanation of these is shown in Table 2.

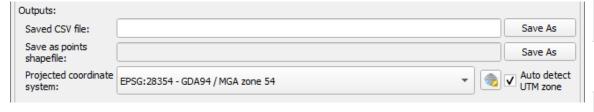
The Clean, Trim, Normalise Point Data Tool can also be used to reproject csv files containing coordinate columns without the cleaning, trim and normalise functionality by ticking the checkbox as highlighted below.











Note: If in Australia then MGA GDA 1994 zones, will be used – see Technical Notes on page 56.

#### **Notes**

Table 2 A description of the filter type used when reporting filtering results.

ORDER	FILTER TYPE	DESCRIPTION
01	clip	Points removed when clipping by a polygon.
02	zero	Nulls and/or Zeros removed.
03	3.0 std iter 1	Points removed during the standard deviation (std) iteration (iter) number 1 and uses 3 standard deviations.
04	3.0 std iter 2	Points removed during the standard deviation (std) iteration (iter) number 2 and uses 3 standard deviations.
05	3.0 std iter 3	Points removed during the standard deviation (std) iteration (iter) number 3 and uses 3 standard deviations.
06	pointXY (1.0 m)	Points removed which are spaced at less than the specified distance apart, in this case 1m.
07	pointX (1.0 m)	Points removed after sorting by the X coordinate, which are spaced at less than the specified distance apart, in this case <b>1 m</b> .
08	pointY (1.0 m)	Points removed after sorting by the Y coordinate, which are spaced at less than the specified distance apart, in this case <b>1 m</b> .
	Pts remaining	The number of points left after filtering occurs.
	Total	The total number of points in the dataset.

The following notification occurs when a delimited text file contains invalid characters in a column name. A copy of the file with corrected column names will be made in the PAT temporary folder (location shown in PAT log message). For more information on the location of the saved log file and temporary folder see the Technical Notes section on page 56.





## Run Kriging Using VESPER

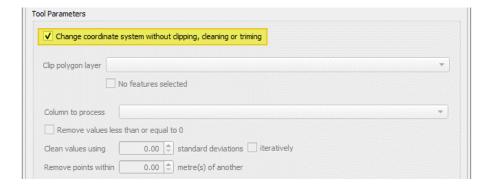
## Summary

This tool is used for mapping on-the-go data at high spatial density, such as are collected by a yield monitor or an EM38 soil survey, following established protocols (Bramley and Williams 2001; Taylor et al. 2007; Bramley et al. 2008; Bramley and Jensen 2014) using local block kriging. It will create a VESPER control file and collate the files required for kriging. The following files will be created in a VESPER sub-folder located in the specified output folder.

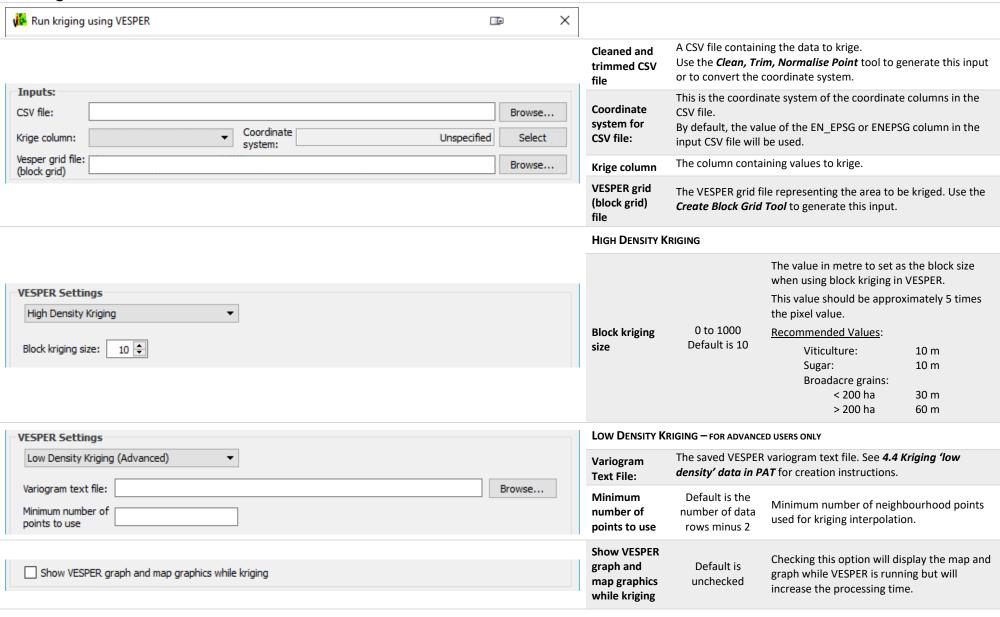
- the VESPER control file. The control filename will be used as a base to derive other VESPER output files like the kriged map result.
- a subset of data to krige. All non-required columns are deleted.
- a Windows batch file (Do VESPER.bat) which can be used to launch VESPER processing for all control files in the VESPER sub-folder. This process can be run outside of QGIS and the Python/pyPrecAg environment.

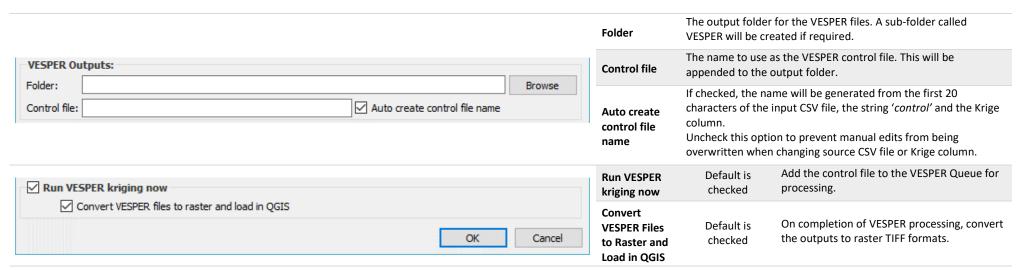
To run VESPER control files from within QGIS, VESPER must already be installed on the PC and configured in the PAT settings (2.17 Settings on page 55).

Data validation checks will be completed before running the tool. This aims at ensuring that the input csv file which contains eastings and northings, overlaps with provided VESPER grid text file. If you receive a warning, please check that these two files use the same coordinate system and indeed overlap. If your csv coordinates are in a different coordinate system to the VESPER grid file, the Clean, Trim, Normalise Point Data Tool can be used to project coordinates without using the clipping, cleaning, trim functionality, by ticking the checkbox as highlighted below.



Kriging of data collected at much lower spatial density, such as through hand sampling (e.g. grape bunch number, mid-season cuts of wheat biomass) is also possible in PAT. However, it should be noted that we DO NOT recommend interpolation of maps from 'low density' data when fewer than 100 data points are available (Webster and Oliver, 2007). Experience also suggests that, in vineyards and sugarcane fields, a sample intensity of around 26 points per ha is required, given the typical dimensions of such fields. A lower sample density may suffice for broadacre situations but the 100 point minimum always applies. Note also that this analysis should only be undertaken by users with an understanding of map interpolation using kriging and the generation and role of the variogram in this process (Webster and Oliver, 2007). Detailed instructions on this process can be found in section 4.4 Kriging 'low density' data in PAT.





## **File Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed with the exception of hyphens (-) and underscores ( ).

DESCRIPTION	FILENAME	<b>EXTENSION</b>	EXAMPLE
Files <u>required</u> by VESPER and copied	d to the VESPER output folder.		
VESPER control file	< 20 characters of CSV file>_ <low density="" high="" or="">_<krige column="">_control</krige></low>	.txt	swblock_HighDensity_YldMassDry_control.txt
VESPER grid file	<control file=""> where control is replaced by vespergrid.</control>	.txt	swblock_HighDensity_YldMassDry_vespergrid.txt
VESPER data file	< <u>control file</u> > where <u>control</u> is replaced by vesperdata.	.csv	swblock_HighDensity_YldMassDry_vesperdata.csv
Files <u>created</u> by VESPER and saved	to the VESPER output folder.		
VESPER Krige file (contains Prediction and SE)	<control file=""> where control is replaced by kriged.</control>	.txt	swblock_HighDensity_YldMassDry_kriged.txt
VESPER parameter file	<control file=""> where control is replaced by parameter.</control>	.txt	swblock_HighDensity_YldMassDry_parameter.txt
VESPER report file	< <u>control file</u> > where <u>control</u> is replaced by report	.txt	swblock_HighDensity_YldMassDry_report.txt
If converting VESPER files to raster	is checked or the <b>Create Rasters from VESPER Results</b> tool is used.		
Kriged Prediction TIFF source: VESPER Krige result above	<control file=""> where kriged is replaced by PRED.</control>	.tif	swblock_HighDensity_YldMassDry_PRED.tif
Standard Error TIFF source: VESPER Krige result above	<control file=""> where kriged is replaced by SE.</control>	.tif	swblock_HighDensity_YldMassDry_SE.tif
Confidence Interval (CI) metadata text file	<control file=""> where control is replaced by CI.</control>	.txt	swblock_HighDensity_YldMassDry_Cl.txt



## Import VESPER Results

## **Summary**

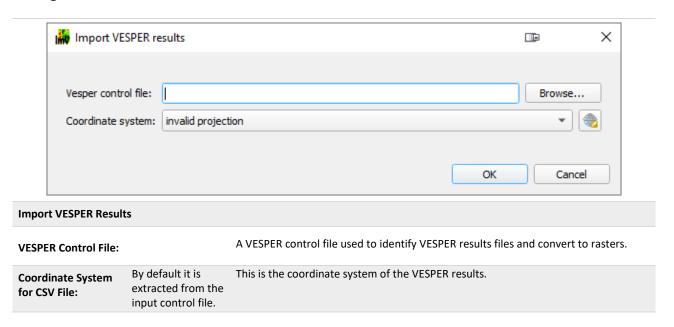
This tool converts the output files generated by VESPER into raster TIFFs.

The output filenames will be created using the VESPER control file name as a base and demonstrated in the Run Kriging Using VESPER Tool - File Naming Conventions section on page 22.

The files created include:

- a TIFF file representing the predicted (kriged) value; these are the values used to generate the resulting map.
- a TIFF file representing the standard error (SE) of the kriging prediction.
- a text file containing the calculated median kriging prediction SE and the 95% confidence interval for the map. This is important information for testing the significance of the difference between different zones when different map layers are clustered to generate potential management zones.

The coordinate system to be assigned to the rasters will be extracted from the value stored within the control file. If the coordinate system cannot be found or is incorrect, it can be selected manually.



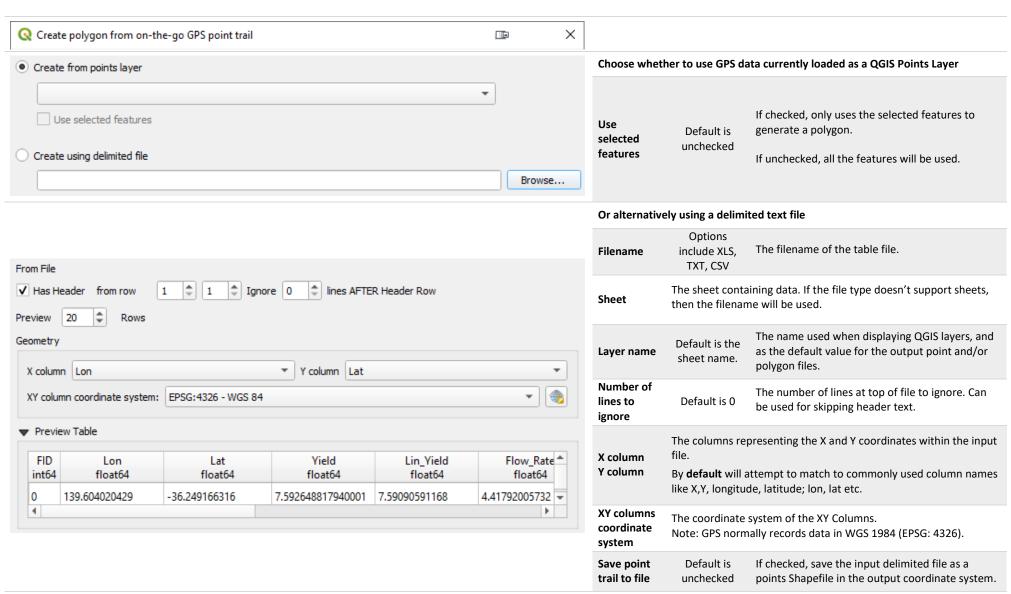


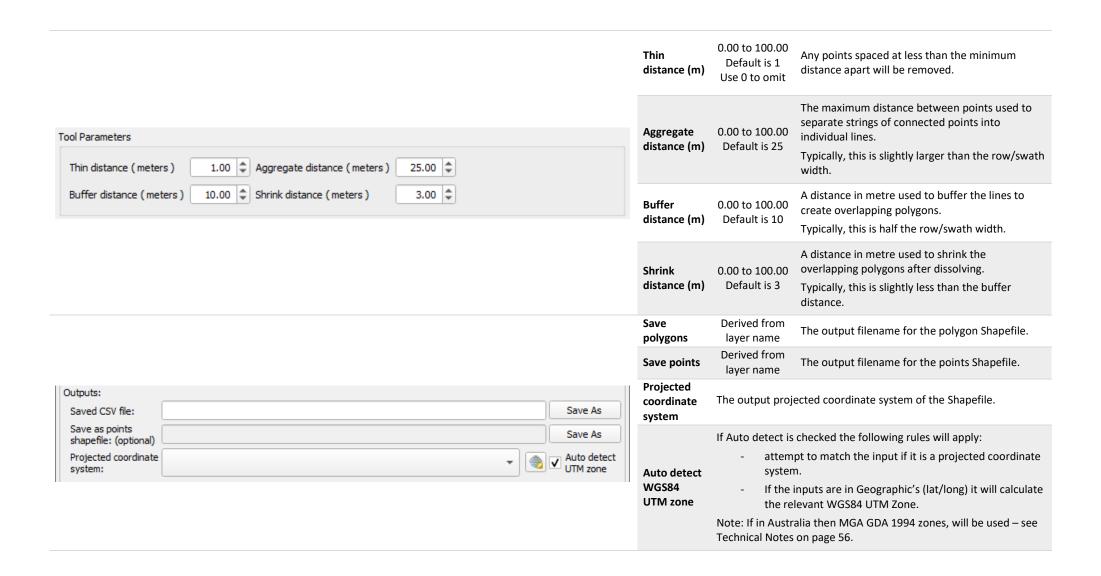
# 2.5 Create Polygon from On-The-Go GPS Point Trail Data

## Summary

Having a block boundary polygon is central to many of the PAT processing steps. Boundary polygons are used to constrain data to a fixed extent. It is preferably that a boundary polygon is created by collecting accurate GPS points around the block and editing them in QGIS to create polygons (refer to How-To -Create a block boundary polygon from a CSV of GPS collected points on page 57 for instructions on this method). However, if accurate GPS data for the boundary is not available, then a less accurate block boundary polygon can be created using this tool, based upon a file of on-the-go GPS points (i.e. from a yield monitor or EM38 survey data). Note that this method is not preferred, but the tool is provided as a 'quick fix' for those wishing to generate maps in the absence of surveyed block or field boundaries.

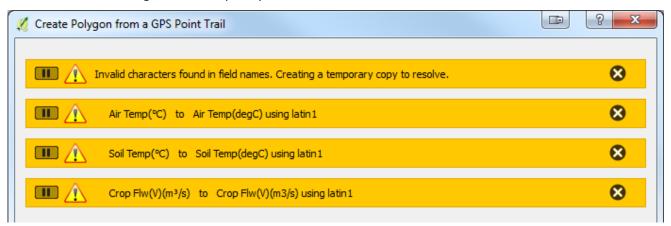
As the process involves a dot-to-dot approach, it is critical that the input file of points are in order (i.e. sorted by an increasing time sequence - as would be normal for data collected by an on-the-go sensor such as yield monitor). For efficiency, points can be thinned by removing points closer than a set distance apart as justified by the accuracy of the GPS. Resulting points will be connected to form lines and then converted to polygons.



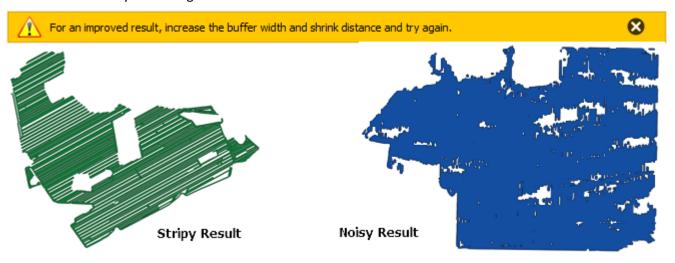


#### **Notes**

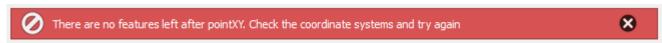
The following notification occurs when a delimited text file contains invalid characters in the column name. A copy of the file will be made in the PAT temporary folder containing corrected column names as specified in the PAT log message and to file. For more information on the location of the saved log file and temporary folder see the Technical Notes section.



A warning is triggered when the resulting polygon is considered stripy or noisy. This can usually be corrected by increasing the buffer and shrink distances.



The following error is caused when the wrong coordinate system is applied to the input source file or layer. Thinning and/or clipping will result all points being removed leaving no points for further processing.





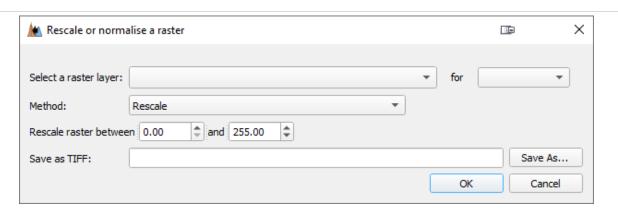
## Rescale or Normalise a Raster

## **Summary**

This tool rescales or normalises a raster and outputs to a new TIFF file. Existing no-data values will be ignored in any calculation.

- Rescale will adjust the raster between the specified values (e.g. output values to a range between 0
- Normalise will adjust the raster to a mean of zero and standard deviation of one

Such re-scaling is useful when you wish to compare several map layers, for example, yield maps for several years.

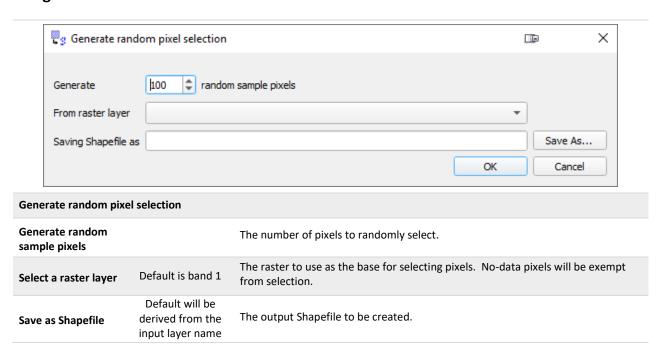


Rescale or normalise a raster					
Select a raster layer for band X	Default is band 1	The raster layer and band to rescale or normalise.			
Method	Default is rescale	Options:  Rescale – adjust values to a fixed range; or Normalise – adjust values to a mean of zero and a standard deviation of one.			
Rescale between (when selected)	Default 0 to 255	The range of values used with rescaling.			
Default will be Save as Shapefile derived from the The output Shapefil input layer name		The output Shapefile to be created.			



## **Summary**

This tool is used to select randomly distributed pixel locations from an existing raster and save them to a point Shapefile. Pixels will be selected from areas inside the raster extent which contain valid data. The resulting points will be located in the centre of the chosen pixel and the output shapefile will contain columns representing the X and Y coordinates. This tool provides the key location input for the Extract Pixel Statistics for Points Tool described on the following page.



## Summary

Extract pixel statistics for points is used to extract pixel statistics from multiple rasters using a pre-defined set of points. These points could be identified using the Generate Random Pixel Selection Tool described on the previous page. Statistics are calculated on pixel values within a square neighbourhood and extracted to a CSV file. This tool is useful when relationships are being sought between different maps layers – for example, a regression or correlation analysis between yield and soil map data.

Applying a neighbourhood filter to rasters is useful for removing (smoothing) small anomalies introduced from instrument inaccuracies or on-the-go movement. The neighbourhood consists of a centre pixel and a number of pixels forming a square around the central pixel, as shown in Figure 3.

9				X				9
	7			х			7	
		5		X		5		
			3	х	3			
				С				

Figure 3. A representation of the 3x3, 5x5, 7x7, 9x9 neighbourhood size around a central pixel (C) as is used when calculating filtered statistics.

For example, a mean statistic for a 3x3 neighbourhood on 2 m pixels, will calculate the mean of the central pixel (red) and the surrounding 3x3 area of 8 pixels (yellow) equating to 36 m2 on ground.

Pixels designated no-data will be excluded from the statistical calculation. However, a central no-data pixel may be assigned a value if at least one pixel in the neighbourhood has a valid value.

Currently, the statistical methods supported by this tool are: mean, standard deviation, co-efficient of variation (CV), minimum, maximum and a count of pixels contributing to the statistical calculation.

The output values are saved to a CSV data file. Column names for each raster and statistic combination are explained below.

💺 Extract pixel stati	stics for points		×			
Select a points layer:	□ No features selected			Select point layer Use selected features	A point layer in  Default is unchecked	QGIS used to extract pixel statistics values.  If <b>checked</b> , extract pixel values only for the selected points. If <b>unchecked</b> , all the points will be used.
Select a raster layer:	Only showing rasters of the same pixel size that Add a raster layer to set the pixel size. Empty List  0 Raster(s)	Add  overlap the points.  = No overlap with points  Move Up  Move Dow  Remove	=	Use raster layer	The pick-list is fill layers with the slayer added. The Add, Delete used to manage. The order of ras	to extract values from.  Iltered to only show overlapping raster same pixel size. This is set by the first raster  If Move Up, Move Down, buttons can be the list of rasters.  Interest is retained in the output CSV file with the first raster and statistic combination.
Include currer	nt pixel value (uses 1x1 neighbourhood by default)  Minimum  3x3	g pixel neighbourhood sizes  7x7		Calculate pixel neighbourhood statistics	the list.  Current pixel ju and/or  Statistics to be	red statistical methods for each raster in st extracts the pixel value, calculated using the set neighbourhood size. mn naming format is explained below.
Standard dev		○ 9x9		Neighbourhood size (pixels)	Default is 3x3	The size of the neighbourhood footprint to use while calculating statistics.
Saved CSV file:		OK Cancel		Save CSV file	Derived from layer name	The name and path of the output CSV file containing the final data points.

## **File and Column Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed with the exception of hyphens (-) and underscores ( \_ ).

## **Output Filenames:**

CSV filename	<qgis layer="" name="" points="">_pixelvals.csv</qgis>	
--------------	--	--

#### **Column Names:**

NAMING RULE	EXAMPLE	EXPLANATION		
<statistic><size>_<raster file=""></raster></size></statistic>	mean3x3_swblock_YldMassDry_PRED	The column containing values from the mean 3x3 neighbourhood filter for the raster swblock_YldMassDry_PRED.tif.		
	pixel_swblock_2009YldMassDry_PRED	The column containing the pixel values for the raster swblock_2009YldMassDry_PRED.		
	cv5x5_seblock_2014YldMassDry_PRED	The column containing values for coefficient of variation (CV) 5x5 neighbourhood filter for the raster seblock_2014YldMassDry_PRED.tif.		
	pixelcount7x7_swblock_2024YldMassDry_PRED	Column containing values representing the count of pixels used for statistical calculations with a 7x7 neighbourhood filter for the raster seblock_2014YldMassDry_PRED.tif.		



## Calculate Image Indices for Blocks

#### Summary

Many providers of remotely sensed imagery deliver data to clients as multi band files. The Calculate Image Indices for Blocks tool is used to calculate indices for a multi band image and aligns the output to the extent and pixel size of a block grid. Indices currently supported are Normalised Difference Vegetation Index (NDVI); the 'simple ratio' or Plant Cell Density index (PCD); Green Normalised Difference Vegetation Index (GNDVI); Chlorophyll Red-Edge index (CHLRE); and Normalised Difference Red-Edge index (NDRE). Table 3 describes how these indices are calculated.

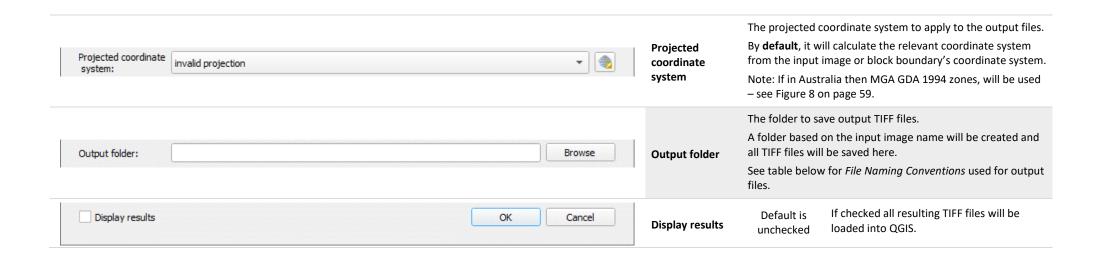
Each band in the image file is mapped to a spectral band e.g. Red, Red-edge, Near Infrared and is then used to calculate relevant image indices. In the case of vineyard data, if a non-vine mask is present in an existing band of the image, it can be used to remove non-vine signals prior to the resampling and alignment to the block grid. This will ensure that spectral signatures relating to ground cover are excluded from the resulting image outputs. This non-vine masking is not relevant for broadacre crops.

An optional block boundary polygon layer, and a column containing the block name or ID, can be used to separate the resulting images into individual blocks. By default, if no column is specified, then all polygons are assumed to be from the one block and will be processed accordingly. If no block boundary layer is specified, then a single polygon outlining the image (excluding no-data) will be used. The output from this tool will be aligned to a block grid that will be identical to the one created separately using the Create Block Grid Tool on page 13.

Table 3 The list of supported indices

Normalized Difference Vegetation Index ( <b>NDVI</b> )	$NDVI = \frac{(NIR - Red)}{(NIR + Red)}$
Green Normalized Difference Vegetation Index (GNDVI)	$\mathbf{GNDVI} = \frac{(NIR - Green)}{(NIR + Green)}$
Normalised Difference Red-Edge Index ( <b>NDRE</b> )	$NDRE = \frac{(NIR - Red Edge)}{(NIR + Red Edge)}$
Plant Cell Density Index (PCD)	$\mathbf{PCD} = \frac{NIR}{Red}$
Chlorophyll Red Edge Index ( <b>ChIRE</b> ) (Gitelson 2004; Gitelson <i>et al.</i> 2005)	$\mathbf{Chl}_{red\ edge} = \left(\frac{NIR}{Red\ Edge}\right) - 1$

Calculate image indices for blocks	×				
Select image: (2 or more bands)	•	Select Image	The image layer containing the appropriate bands required for calculate indices.		
No-data value:		No-data value	Default from Can be used to specify a <u>different</u> no-data image. value.		
		Use a block boundary	A layer in QGIS containing polygon(s) representing blocks.		
Use a block boundary Block ID column: (optional)	•	Use selected features	Default is If <b>checked</b> , only the selected polygons will unchecked be used.		
□ No features selected		Block ID column	A column containing the block id or name. This will be used treat multiple polygons with the same block id or name as one.		
Resample to 2.00 pmetre pixels		Resample pixel size (m)	The pixel size to apply to the raster outputs. This is expressed in metres. Ideally this should match your <b>Block Grid TIFF</b> file.  0.00m to 6 km Default is 2 m  Viticulture: 2 m Sugarcane: 2 m Broadacre grains:  < 200 ha 5 m > 200 ha 10 m		
Specify image bands used for index calculations  Green		Specify image bands used for index calculations	Mapping of band numbers to band types. The mapped bands will enable/disable indices based on individual index requirements.  For most images, Green is band 2, Red is band 3, Near Infrared is band 4.  Non-vine mask is a band where pixels not containing vine signals are set to no-data. Not required for broadacre data.		
Non-vine mask ▼		Select the indices to calculate	The indices to calculate.  If an index is disabled (greyed out) the band it requires hasn't been mapped.  For index acronym and equation, see table above.		



## **File Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed from strings with the exception of hyphens (-) and underscores ( \_ ).

#### Filenames:

NAMING RULE	EXAMPLE	EXPLANATION
Output Folder:		
<output_folder>\<image_name></image_name></output_folder>	C:\data\vineyard\rgbi_jan_50cm_84sutm54_tif	A new folder based on the image name is created in the output folder and all created images are saved here. In this example the image rgbi_jan_50cm_84sutm54.tif is used to create a new folder called rgbi_jan_50cm_84sutm54_tif.
Image Names:		
   	B1_NDVI_2m.tif	The TIFF file resampled to 2m pixels for the Normalised Difference Vegetation Index (NDVI) created for block id/name of B1.
	PCD_250cm.tif	The TIFF file resampled <b>250cm</b> pixels for the Plant Cell Density ( <b>PCD</b> ) Index created without specifying a block id column.



# 2.10 Resample Image Band for Blocks

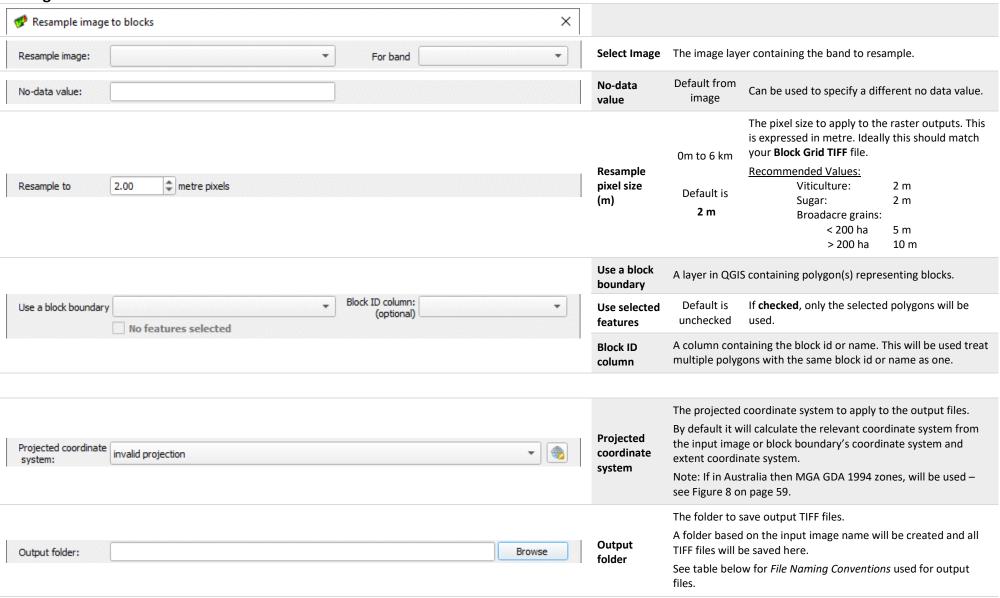
## **Summary**

Resample Image Band for Blocks is used to resample, align and smooth an existing band of an image to match a block grid.

An optional block boundary polygon layer and a column containing the block name or ID, can be used to separate the resulting images into individual blocks. By default, if no column is specified, then all polygons are assumed to be from the one block and will be processed accordingly. If no block boundary layer is specified, then a single polygon outlining the input image will be used.

The output from this tool will be aligned to a block grid that will be identical to the one created separately using the *Create Block Grid Tool* on page 13.

### **Dialog**



Display results	Display results	Default is unchecked	If checked all resulting TIFF files will be loaded into QGIS.
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### **File Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed from strings with the exception of hyphens (-) and underscores ( \_ ).

### Filenames:

NAMING RULE	EXAMPLE	EXPLANATION
Output Folder:		
<output_folder>\<image_name></image_name></output_folder>	C:\data\vineyard\area1_rgbi_jan_50cm_84sutm54_tif	A new folder based on the image name is created in the output folder and all created images are saved here. In this example the image area1_rgbi_jan_50cm_84sutm54.tif is used to create a new folder called area1_rgbi_jan_50cm_84sutm54_tif.
Image Names:		
   	B1_Band6_2m.tif	The TIFF file where <b>Band 6</b> is resampled to <b>2m</b> for block id/name of <b>B1</b> .
	Band7_250cm.tif	The TIFF file where <b>Band 7</b> is resampled <b>250cm</b> pixels <b>without</b> specifying a block id column.



## 2.11 Create Zones with k-means Clusters

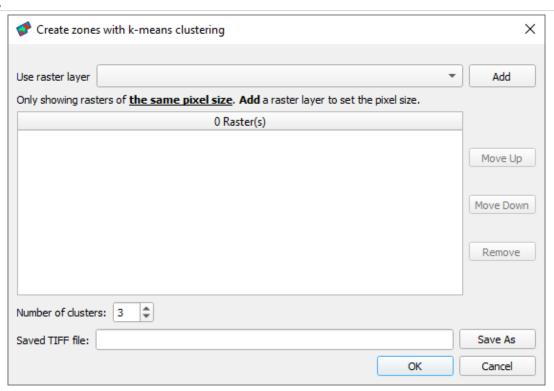
### **Summary**

This tool allows zones to be created by performing k-means clustering on multiple rasters (i.e. map layers) to create clusters of similarity by minimising variability within clusters while maximising variability between clusters. If significant differences between clusters are observed, then the clustered results can be used as potential management zones.

Raster files with the same single pixel size, and coordinate systems are used as inputs and the common area of overlap will be used to generate an output TIFF containing the clustered result.

On completion of k-means clustering, the mean and standard deviation for each zone/cluster and source raster combination will be calculated and written to a CSV File alongside the output TIFF file as well as being displayed in PAT's log messages panel.

### **Dialog**



### Create zones with k-means clustering

The raster layer to extract values from.

Only raster layers of the same pixel size can be analysed at a time. The first raster layer added sets both the pixel size and the output coordinate system.

Use raster layer:

The Add, Delete, Move Up, Move Down, buttons can be used to manage and order the list of rasters.

The order of rasters is retained in the output CSV file with separate columns for each raster and statistic combination.

Number of clusters:

Default is 3

The number of clusters/zones to create. Aside from trying 3, it is recommended that users also try 2 clusters. 4, 5 or an even larger number of clusters may also be

appropriate.

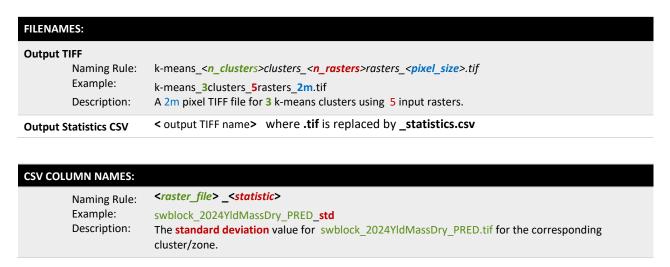
The name and path of the output TIFF file representing the zones.

Save TIFF file

In addition, a statistics CSV file will be written to disk along with the TIFF, and results printed to PAT's log messages panel.

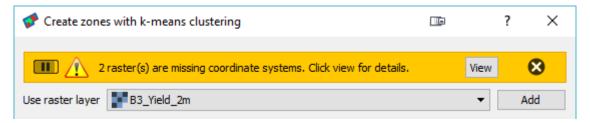
### **File and Column Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed from strings with the exception of hyphens (-) and underscores (\_).

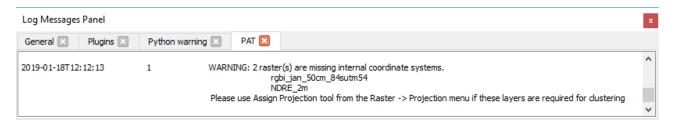


#### **Notes**

To successfully use this tool, all input raster files must contain the coordinate system internal to the file. When a file without a coordinate system is loaded into QGIS, an external coordinate system will be applied based on your QGIS settings. These files cannot be used by this tool. The user will be notified of this when launching the tool as follows.



Clicking on view will open the PAT Log Messages Panel and provide a list of those images. Users can then use the Assign Projection Tool from the Raster -> Projection menu to assign the appropriate internal coordinate system to the files. If the Assign Projection tool is unavailable activate the GDALTools plugin.



This tool is used to generate the data needed to conduct the moving window t-test method of Lawes and Bramley (2012) and is available in PAT's Run Strip Trial t-test Analysis Tool. It may be useful to users who, for example, have implemented a treatment like an N-rich strip in their cereal paddock, or who want to compare a row of vines that have been pruned differently to the rest of a vineyard.

The tool uses a line representing the centre of a strip trial (or treatment) and creates points along the line spaced using the nominated distance. In addition, two sets of parallel 'control' points are created at an offset to the strip trial line as shown in Figure 4. This allows the treatment strip data, to be compared to data from the neighbouring (control) area.

The tool creates a number of identifiers to facilitate the analysis: each point has values assigned to new columns headers Trial<u>ID</u>, <u>Strip Name</u>, <u>PointID</u> and <u>DistOnLine</u>. The treatment strip is assigned a unique identifier in the column TrialID and the value of 'Strip' in the Strip Name column. Points are created along the line and assigned a unique point identifier in the column PointID. In addition, two sets of control points are placed at the offset distance either side of the strip. The PointID of the control points are paired with the corresponding treatment point. The Strip Name values of the control points are assigned based on the direction of offset from the original treatment line as shown in Figure 4. Each point is also assigned a value in the column <u>DistOnLine</u> representing the distance in metres of the point from the start of respective line.

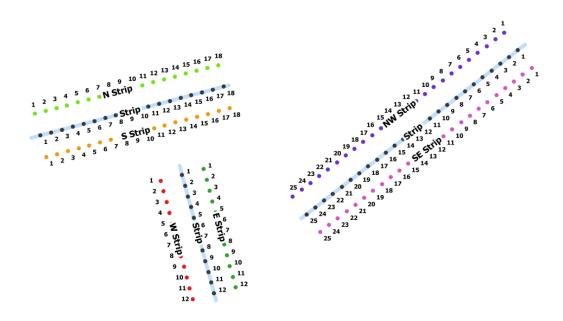
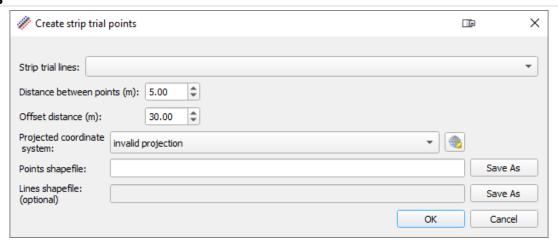


Figure 4 Output from the Create Strip Trial Points tool. The blue line represents the treatment strip and the dots indicate the equally spaced treatment and control (i.e. offset) points.

### **Dialog**



#### Create strip trial points The layer containing the strip trial lines. Strip trial lines: The distance used to space points along the centre line. In this instance, users are advised to select a value appropriate to their trial rather than accepting the default. 1 - 500 mDistance between If used in conjunction with a strip trial t-test then this distance should be the pixel size of points (m): Default is 5m the input rasters used for the t-test. This will result in every pixel in the strip being used for the analysis. The distance used to offset the points <u>from</u> the line. If used in conjunction with a strip trial t-test then this should be far enough away from 1 - 500mOffset distance (m) the strip trial to remove the trial's treatment influence. In the case of an N rich strip trial Default is 30m laid down using a boom spray, the offset distance ought to be at least equivalent to the length of the boom. The projected coordinate system to apply to the output files. By default it will calculate the relevant coordinate system from input layer's coordinate system. **Projected** coordinate system: Note: If in Australia then MGA GDA 1994 zones, will be used – see Figure 8 on page 59. Derived from The name and path of the output points shapefile. Points shapefile: layer name The name and path of the output line shapefile. Save line shapefile: File name is derived from the output points shapefile

### **Column Naming Conventions**

CSV COLUMN NAM	IES:
FID	Unique feature identifier for each point.
TrialID	The unique strip trial identifier.
Strip_Name	A name assigned to each line. The treatment will be assigned a value of 'Strip' while the offset strips will be assigned the cardinal direction of the offset to the treatment as shown in Figure 4.
PointID	The identifier of each point along the centreline, and its paired control points.
DistOnLine	The distance in metres of the point from the start of each line.

This tool enables a user to analyse strip trials to determine the effectiveness of treatments. For example, it could be used analyse the difference between an N-Rich strip treatment and the adjacent normally fertilized paddock, or to compare different pruning or spray treatments in a vineyard.

Using points created from the Create Strip Trial Points Tool, values will be extracted from input rasters and a pairwise moving window t-test conducted based on a calculated Response Index (RI) and treatment differences. These are then graphed as described in Lawes and Bramley (2012).

The input rasters include:

- Strip values raster: A map produced through following the high density on-the-go sensor mapping workflow (Figure 1) which includes data sourced from the trial strip.
- Control values raster (optional): A map produced through following the high density on-the-go sensor mapping workflow after data sourced from the trial strip have been removed. Often this is desirable because, when interpolating a map containing all the data for a 'nitrogen rich' strip, the higher values in the strip may impact the interpolation of adjacent areas. In general, our recommendation would be to make use of a control values raster by re-kriging the original data excluding the data points under the trial strip.
- Zone raster (optional): A zone raster can be created using the Create Zones with k-means Clusters tool. You may want to use this to identify which zone different parts of the experimental strips are located in. Desirably in a field with defined zones the strip would run across these zones.

For each input strip a map image will be produced along with a graph and CSV file, for each strip combination a) Strip vs Control 1, b) Strip vs Control 2, c) Strip vs Average of Control 1 & 2 (Figure 5).

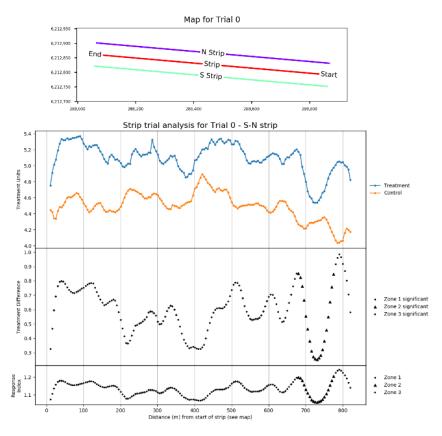
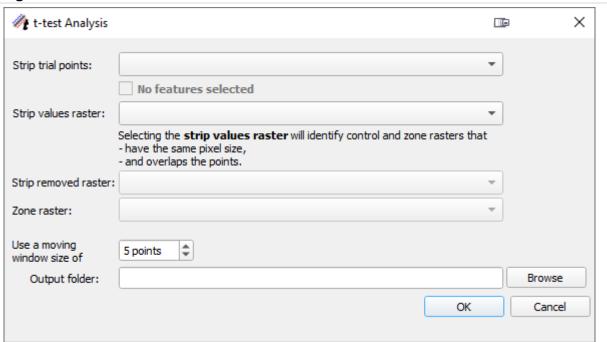


Figure 5 Output graph and map from the Run Strip Trial t-test analysis tool.

### **Dialog**



Run Strip Trial t-test Analysis			
Strip trial points:	The layer containing the strip trial points. This must be the output from the <i>Create Strip Trial Points Tool</i> .		
Strip values raster:	The raster containing values to extract for analysis.		
Strip removed raster:	Optional The raster layer created after treatment points have been removed.		
Zone raster	Optional  The raster containing the zone values. If not provided, a zone of 1 will be applied to the entire strip.		
Use a moving window size	5 points The number of points to use for the moving window t-test calculation.		
Output folder:	The folder to save output map, graph and CSV files.  See table below for used for output files.		

### **File and Column Naming Conventions**

- <> denotes an existing element or input
- non-alphanumeric characters are removed from strings with the exception of hyphens (-) and underscores ( \_ ).

*TrialID* represents the unique strip trial identifier from input points *TrialID* attribute.

strip represents the value(s) from input points Strip\_Name attribute. When the analysis uses both control strip points their values will be hyphenated e.g. N-S

FILENAMES:	
Map file:	A map showing the Trial points.
Naming Rule:	Trial-< <mark>TrialID&gt;</mark> _map.png
Example:	Trial-3_map.png
Description:	The map for the strip with a TrialID of 3.
Graph files:	The results displayed as graphs.
Naming Rule:	Trial-< <i>TrialID&gt;_</i> < <i>strip</i> >_graph.png
Example:	Trial-3_N-strip_graph.tif
Description:	The graph result for Trial 3 which compares the N control strip to the treatment strip.
CSV files:	The csv file used to create the graphs.
Naming Rule:	Trial-< <i>TrialID&gt;_</i> < <i>strip</i> >.csv
Example:	Trial-3_N-S-strip.csv
Description:	The csv file containing the Trial 3 results for the comparison of N-S control strip to the treatment strip.

CSV COLUMN NAMES:	
TrialID	The unique strip trial identifier from the input Strip Trial Points.
PointID	The identifier of each strip trial point from the input Strip Trial Points, and the paired control points.
DistOnLine	The distance in metres of the point from the start of the centreline of the treatment strip.
Strip Value	The number extracted from the <b>Strip value raster</b> .  The units of value will be inherited from the source data (eg NDVI, Yield t/ha).
<strip> Control (if provided)</strip>	The number extracted from the <b>Control values raster</b> .  Applies only when a <b>Control values raster</b> is used.  The units of value will be inherited from the source data (eg NDVI, Yield t/ha).
Strip Zone (if provided)	The zone identifier extracted from the <b>Zone raster</b> .  Creates a Zone value of 1 if not specified.
< <b>strip</b> >_mean	The mean of the input control strips for each point if both control strip are used.
treat_diff	The difference between the treatment and controls points columns.
av_treat_diff	The average of the column <u>treat_diff</u> using a moving window.
p_value	The t-test p-value calculated using a moving window.
RI	The RI calculated using a moving window:  RI = mean of treatment / mean of control

The column names for the output file are dependent on the input scenarios. In the following example, a subset of the column names for the East-West strip trial shown in Figure 4 where a strip values raster, control raster and zone raster were provided as inputs.

CSV COLUMN NAMES:			
Strip Value	The pixel value from the <u>strip value raster</u> for the strip trial points.		
N Strip Control S Strip Control - or -	The pixel value from the control raster for control line to the north or south of the treatment strip.		
N Strip Value S Strip Value	If a <u>control raster</u> is not provided, then pixel value will come from the strip value raster and the column name will be <b>N Strip Value</b> or <b>S Strip Value</b> .		
Strip Zone	The pixel value from the <u>zone raster</u> for the strip trial.  If a zones raster is not provided, then a value of 1 will be assigned.		
N Strip Zone S Strip Zone	The pixel value from the zone <u>raster</u> for control points to the north or south of the strip trial.  If a zones raster is not provided, then a value of 1 will be assigned.		
N-S_mean	The average of the control strip points when both of sets of control strips are used in the analysis.		

Whole-of-block experimentation, also known as spatially distributed experimentation, involves on-farm experiments where treatments are applied in a highly replicated design over an entire block or land management unit, or a large section of it. Given their scale, they are typically laid down using the farmer's normal equipment. These experiments can be thought of as a three-dimensional version of the strip trial, discussed in the previous section *Create Strip Trial Points*. Users interested in this experimental approach may wish to read the book chapter by Bramley et al. (2013) and perhaps the references listed in that chapter. In brief, the objective of whole-of-block experimentation is to recognise the possible effect of underlying spatial variability in the field in which the experiment is located, on the crop response to experimental treatments, and so use this variability as an experimental tool to guide decisions about variable rate or targeted management. Whole-of-block experimentation therefore moves away from the classical approach to agronomic experimentation which seeks to identify whether treatment A is better than treatment B and typically uses small plots. Instead, it is recognised that treatments A and B may both deliver benefit, albeit in different parts of the block or field; the whole-of-block approach identifies these different areas.

An example is given in Figure 6. Here, some different vineyard floor management treatments have been evaluated in an experiment aimed at improving the availability of water to the vines. The cereal/legume treatment delivers a significant benefit over the other treatments, but this is expensive and so the vineyard manager would only want to use this in those areas where it is beneficial. The whole-of-block analysis tool enables users to generate maps such as those shown in Figure 6 – maps showing the response to the different treatments over the entire block, and maps indicating the significance of differences between these treatments.

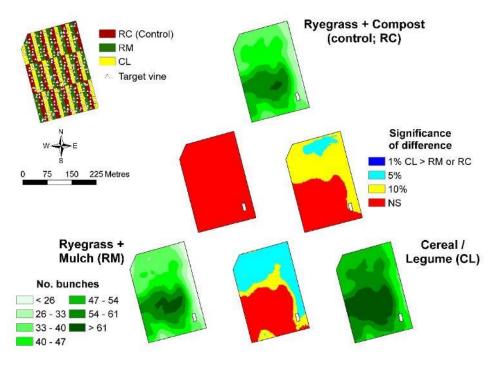


Figure 6 An example of a whole-of-block experiment conducted in a 4.8 ha Clare Valley vineyard. Panten and Bramley (2011) provide further information

The purpose of the Whole-of-Block analysis tool is to predict treatment responses and their differences over the entire experimentation area as illustrated by Figure 6. Similar to the use of VESPER for kriging, this tool interpolates point data (low or high density) to the area of the entire block. Low density data would typically include up to a few hundred points, perhaps collected through hand sampling, dispersed across a block, whereas high density data includes a few thousand or more points, typically collected using a sensor such as a yield monitor. The tool will handle either two or three treatments and will identify areas of the block where the treatments are significantly different from each other. To address the technical and computational challenges in this analysis, the tool uses fast and efficient global and local cokriging techniques.

#### Use of R

This tool is somewhat different from the other PAT tools in that it uses the R statistical language (http://rproject.org/) to carry out the analytical processing. This means that R must be installed on your PC for this to work – see *Configuring QGIS to use R* on page 10. The minimum R version suitable for this is v3.5.1.

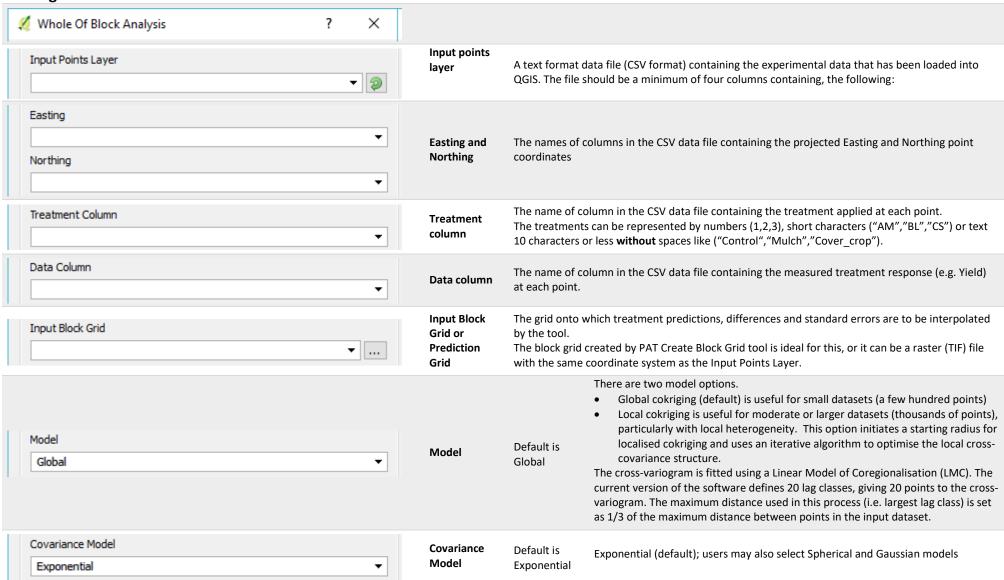
### Algorithm for the Local Cokriging System – a brief commentary on the data radius used

A spatially-varying local co-kriging method is used to balance the trade-off between the computation efficiency and prediction accuracy. We use a neighbourhood radius r(s) to identify neighbours for location

For subregions with lower density observations, such as at the corners, or edges of a farm field, the neighbourhood radius required by local cokriging will be relatively bigger to ensure the stability of the estimated cross-variograms and balance the number of observations from different treatments within the neighbourhood.

For high density datasets we will use a relatively small neighbourhood radius, and ensure it will change relatively smoothly across the field. This adaption is achieved by two specific algorithm choices. Firstly, an initial neighbourhood radius is selected automatically based on the observation density and achieving a reasonable balance in the number of observations from different treatments. This neighbourhood radius increases iteratively to include more observations if the current radius is not large enough to meet prediction quality criteria. Secondly, a spatial-varying neighbourhood radius surface is estimated based on neighbourhood radius values from a subset of locations. This exploits the spatial-smoothness of the neighbourhood radius and provides a better default neighbourhood radius from which to optimise.

### **Dialog**



User Defined Neighbourhood for Local CoKriging Input Neighbourhood Size in Metre  30	User Defined Neighbourhood radius for Local Cokriging	Only applicable to the Local Cokriging method, and allows a user to define an initial neighbourhood radius. If this option is ticked then user should define the "Input Neighbourhood Size in Metres". The default is 30 m. A suggested starting value can be calculated as 15 times the pixel size of the block grid. However, if "User Defined Neighbourhood for Local Cokriging" is not ticked then the initial neighbourhood radius will be chosen as a function of total number of data points and corresponding number of treatments. This is the recommended starting point for using this tool.	
Save Output [optional]	Save output	Folder in which to save the output files.  See File Naming Conventions (below) for an explanation of the different output files that will be created in this folder.	

### **File Naming Conventions**

- <> denotes an existing element or input. For example <Data Col> refers to the name of the Data Column selected in the input parameters (above).
- non-alphanumeric characters are removed from strings with the exception of hyphens (-) and underscores ( \_ ).

### Filenames:

NAMING RULE	EXAMPLE	EXPLANATION
Output file names:		
<data col="">_dump_file_<model><covariance model="">.tif</covariance></model></data>	Bunch_wt_dump_file_GlobalExplog	Logged information mainly useful for debugging purposes
<pre><data col="">_model_parameters_<model><covariance model="">.txt</covariance></model></data></pre>	Bunch_wt_model_parameters_GlobalExptxt	Summary of parameters used during processing
<data col="">_list_of_tif_files_<model><covariance model="">.txt</covariance></model></data>	Bunch_wt_list_of_tif_files_GlobalExptxt	List of TIF files generated as output
<data col="">_tr_<tmt1>_<model><covariance model="">tif</covariance></model></tmt1></data>	Bunch_wt_tr_KC_GlobalExptif	Treatment inference in "Input Block Grid or Prediction Grid".
<data col="">_tr_<tmt1>_var_<model><covariance model="">tif</covariance></model></tmt1></data>	Bunch_wt_tr_KC_GlobalExptif	Corresponding variances of the treatments.
<data col="">_ tr_diff_<tmt1>_<tmt2>_<model><covariance model="">tif</covariance></model></tmt2></tmt1></data>	Bunch_wt_p_val_KC_SP_GlobalExptif	Predictions of treatment response differences between treatments
<pre><data col="">_tr_<tmt1>_var_<model><covariance model="">tif</covariance></model></tmt1></data></pre>	Bunch_wt_tr_KC_var_GlobalExptif	Corresponding covariances of the treatment differences.
<data col="">_p_val_<tmt1>_<tmt2>_<model><covariance model="">tif</covariance></model></tmt2></tmt1></data>	Bunch_wt_p_val_KC_SP_GlobalExptif	p-values of the treatment differences between pairs of treatments.
<data col="">_z_<tmt1>_<tmt2>_<model><covariance model="">tif</covariance></model></tmt2></tmt1></data>	Bunch_wt_z_KC_SP_GlobalExptif	Z-statistics of the treatment differences.



Persistor is used to summarise the similarities in patterns of variation across a block over a number of years (Bramley and Hamilton 2005). It might be used as an alternative to k-means clustering for identifying clusters based on yield, but also enables zones to be assessed in terms of a target level of performance (e.g. mean yield + 10%) and so provides information about how frequently such a target is met.

The "Target over all years" method assigns a value to each pixel to indicate the number of instances (in the raster list) in which that pixel was either less than or greater than the mean (+/- a nominated percentage) of that raster.

The "Target probability" method builds on the target over all years method, in that it includes an upper range (i.e. cells with a given frequency of values that are above the mean +/- a given percentage) and a lower range (i.e. cells with a given frequency of values that are below the mean +/- a given percentage).

The tool assigns a value to each pixel which indicates whether the performance in that pixel over a given proportion of years is:

- a) Greater than the mean plus or minus the nominated percentage (value = 1)
- b) Less than the mean plus or minus the nominated percentage (value = -1)

The remaining pixels which do not fall into category a) or b) are given a value of 0.

All input grids must overlap and have the same coordinate system and pixel size.

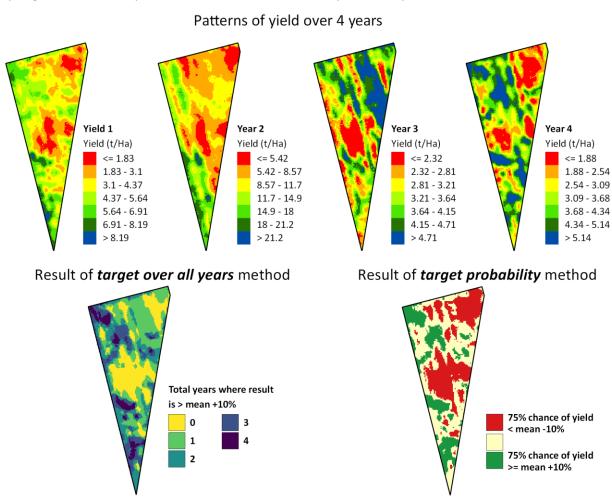
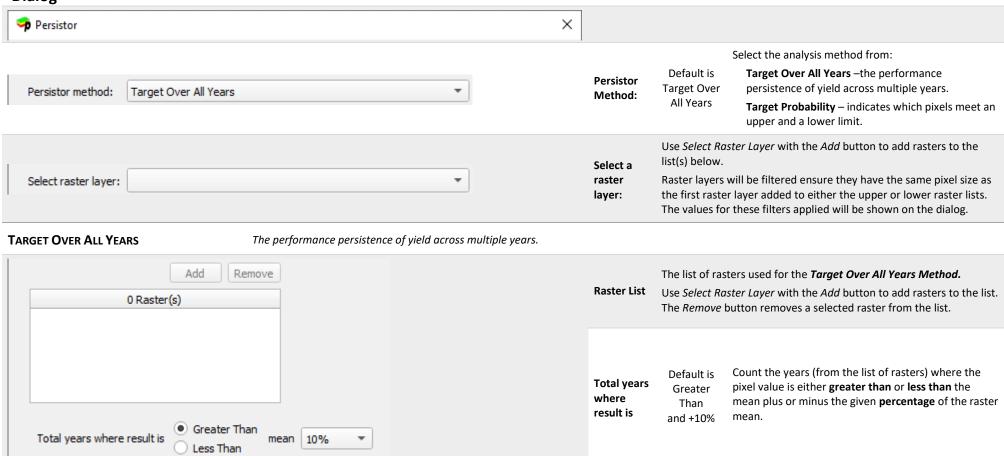
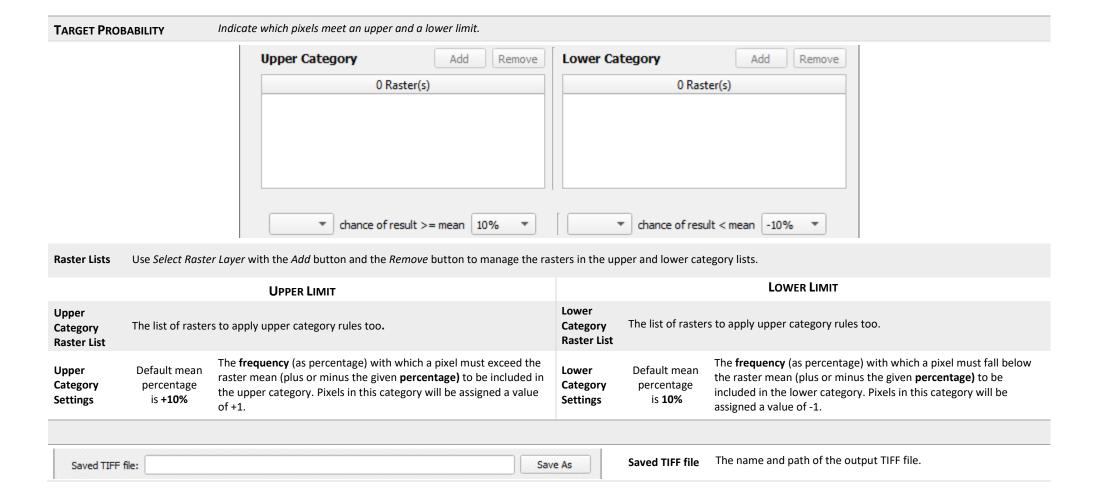


Figure 7 An illustration of the *Persistor Tool* results.

### **Dialog**



continued on next page......

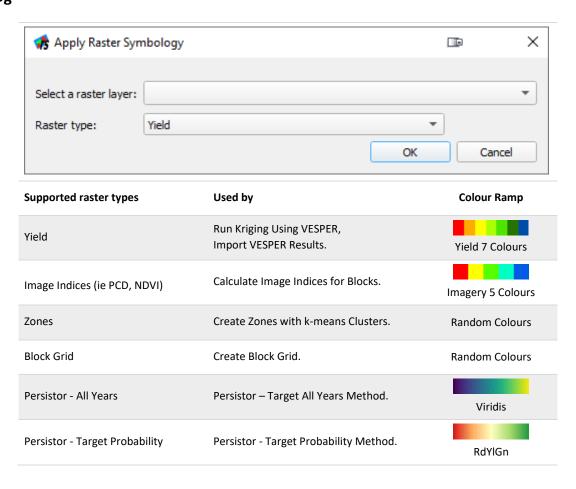




The *Apply Raster Symbology Tool* allows for the quick application of the symbology for the different PAT tool output rasters.

Instructions for installing the PAT symbology can be found in 4.3 Loading PAT Symbols into QGIS.

### **Dialog**





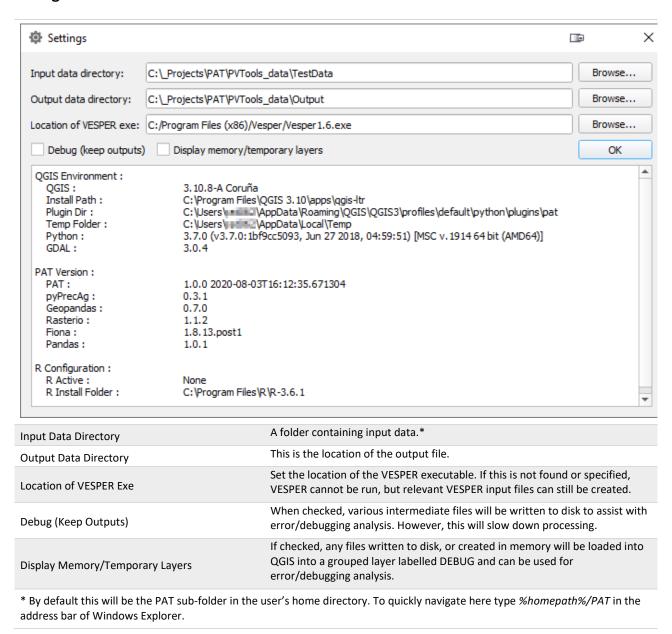
This tool is used to display and edit PAT settings.

The input and output data directories set here will be used to set the default paths by the browse for file/folder functionality for all tools. Each tool will store and access its own values after first time use.

Checking the Debug (keep outputs) box will save intermediate files created while processing data to file. It should be noted that this will slow down the time taken to run tools, but can be a useful diagnostic tool.

The Display Memory/Temporary Layers checkbox can be used to add the intermediate files to QGIS along with any in-memory or virtual layers which are used but not saved to disk.

### Dialog



## 3 Technical Notes

### 3.1 General Notes

- PAT makes use of the CSIRO-developed pyPrecAg Python module which is an open source Python package containing a range of specialised analysis functions.
- All intermediate files created while processing is located in the PrecisionAg folder of the user's temporary folder. To quickly navigate to the temporary folder, type %temp%/PrecisionAg in the address bar of windows explorer. This folder is deleted when QGIS exits.
- All progress, messages and errors are displayed in the PAT tab of the Log Panel as shown in Figure 2 and are saved to a log file located in the PrecisionAg folder of the user's temporary folder. A list of important paths including the location of temp and the user's plugin folder.
- A Users QGIS Plugin folder can be found by typing %homepath%/.qgis2/python/plugins into the address bar of Windows Explorer.

## 4 How-To's

## 4.1 Create a block boundary polygon from a CSV of GPS collected points

1. Using the *New Shapefile Layer* tool from the *manage layers toolbar* (or Layer menu -> Create Layer -> New Shapefile Layer) create a new polygon shapefile adding the relevant coordinate system\* and attribute fields you require. Clicking OK will prompt you for the location to save the shapefile.

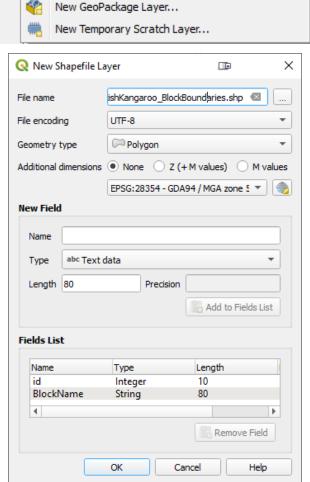
More information on coordinate systems can be found at

- 2. Set a style and labelling to the polygon layer. A hatching polygon fill works well for editing.
- 3. Launch the *Add Delimited Text Layer* tool from the *manage layers toolbar* (or Layer menu-> Add Layer -> Add Delimited Text Layer) and load your GPS CSV file as a layer into QGIS.

  Hint: Your coordinate system is probably WGS 84.
- 4. If required, load other vector or raster data, like imagery, which can be used as reference.
- 5. Setup your snapping environment.
  - a. Open *snapping* options (*Project menu ->* Snapping Options)
  - b. Enable snapping by clicking the horseshoe icon.



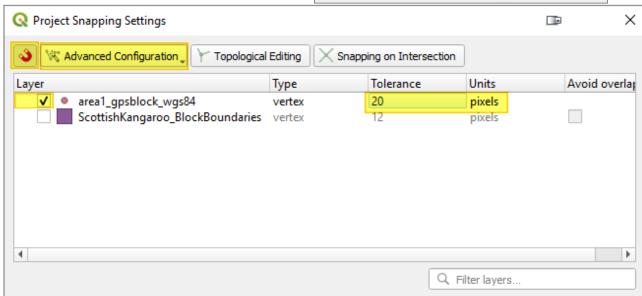
- c. Change *layer selection* to *advanced configuration*.
- d. **Tick** the layer containing the **point layer** loaded in step 3.
- e. Change the tolerance to 20 and set units to pixels



New Shapefile Layer...

New SpatiaLite Layer...

Ctrl+Shift+N



<sup>\*</sup>More information on selecting a coordinate system can be found on page 59.



- 7. Add new features by using the *add features* tool from the *digitizing toolbar*.
- 8. As you move the mouse close to a point the point will change to show a magenta cross hairs (+) or box ( ), this means the mouse has snapped to this point. Clicking the mouse will add this point as a vertex in the polygon. Continue following around the points to form a polygon. Right-mouse-click to finish a polygon.
- 9. When you finish a polygon a dialog will open to allow you to enter attributes. Click **OK** to add attributes and finalise polygon.

on *digitizing toolbar* and *toggle editing* off Save your edits using the save layer edits icon when complete.

**To add, move or delete a vertex**, toggle to node mode using the *node tool* . Click the polygon to edit. Nodes/Vertex will appear as red squares.

- **Double click** to **add** new vertex.
- Single click to select existing vertex. The square will turn blue. Use the DEL key to delete
- Click and drag a vertex to move.

To add a hole (donut) to a polygon use the add ring tool from the *advanced digitizing toolbar* and sketch your polygon as described in step 8.

To delete a hole (donut) in a polygon use the delete ring tool [2] from the advanced digitizing toolbar and click in the hole.

from the advanced digitizing toolbar and sketch the To split a polygon, use the *split features* tool path to split. Multiple polygons will be created having the same attribution.

## Useful editing shortcut keys.

Add new feature	Ctrl+.	Zoom in	Scroll wheel or	Ctrl ++
Delete last vertex	Del	Zoom out	Scroll wheel or	Ctrl +-
Undo	Ctrl+Z	Zoom full	Ctrl+Shift+F	
Cancel edit	Escape	Pan	Middle mouse or Spacebar (while adding a feature only). Note: spacebar also turns active layer visibility on/off.	

### 4.2 How to reproject a shapefile and why you need to do this

A projected coordinate system is used to depict the curved earth surface as a flattened surface – as it is when printed on a piece of paper. Currently the block grid tool requires the polygon shapefile to be in a projected coordinate system. However, raw GPS data such as you might have in your yield monitor data file uses a geographic coordinate system most likely with position expressed as decimal degrees or degrees minutes second (such as in Google Maps). These will typically be labelled as latitudes and longitudes. If your shapefile is in a geographic coordinate system then you will need to reproject it to a projected coordinate system.

Geographic coordinate systems include:

Global: EPSG: 4326, WGS 84 Australian: EPSG: 4283, GDA94

In projected coordinate systems, the Globe/Australia is divided up into zones (see UTM zone map). Map grid of Australia (MGA) zones align closely with the UTM zones, but use the 1994 Geocentric Datum of Australia. Figure 8 shows the position of the MGA zones.

Projected coordinate systems include:

Global: WGS84 / UTM Zone divided in to North and South of the equator

Australian: GDA94 / MGA zones

You need to know the projected coordinate system for your data and which zone it lies in. This projected coordinate system will get used throughout PAT. This is made simpler through the use of EPSG numbers. An EPSG number is an international numbering system for coordinate systems (see http://www.epsg.org)

#### 1. Determine your projected coordinate system.

- Within Australia use GDA 1994 Map grid of Australia (MGA) EPSG: 283xx
- Outside Australia use the WGS84 UTM Zone system.

North of the Equator - EPSG: 326xx South of the Equator - EPSG: 327xx

Using your EPSG prefix listed replace the xx with your zone. For example Adelaide is an Australian city in zone 54 so projected coordinate system then becomes EPSG: 28354 GDA94 / MGA zone 54.

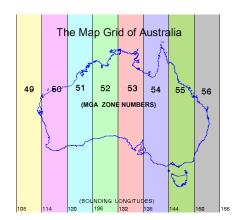


Figure 8 The Map Grid of Australia. (www.environment.gov.au/erin/tools/mga 2geo-gda.html).

### 2. Reproject your shapefile

- a. In QGIS, select your shapefile and select **Save as** from the **Layer menu.**
- In the dialog, browse for a new shapefile and assign a name.
- Click the icon to the far right of the CRS options.

Selected CRS (EPSG:4326, WGS 84) -

- d. Search for your projected coordinate system. The easiest way is to use the EPSG number from step
- e. Leave all other options as is and click OK

The shapefile will be reprojected and loaded into QGIS and is now ready for use.

For further information on coordinate systems see

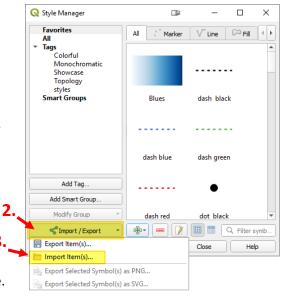
https://docs.qgis.org/3.10/en/docs/gentle\_gis\_introduction/coordinate\_reference\_systems.html

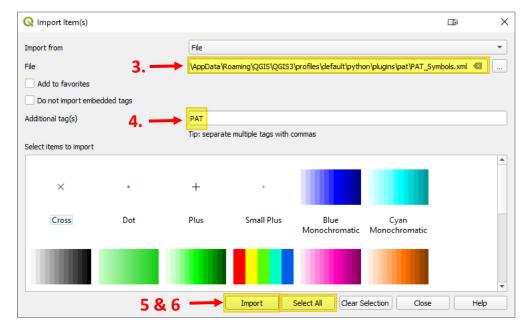
### 4.3 Loading PAT Symbols into QGIS

The PAT plugin includes a pre-defined set of symbols and colour ramps for use with datasets derived while using the plugin. They should be automatically loaded when PAT is installed.

To manually load the styles:

- 1. In QGIS, launch the **Style Manager** from the settings menu.
- 2. From the lower left corner of the dialog, select *Import* dialog.
- 3. Set Import from to file specified below. Browse to the users QGIS plugin folder and find the PAT symbols.xml file in the pat folder (see 1.5 Uninstall PAT for help finding this folder).
- 4. Enter PAT as an additional tag(s).
- 5. Select symbols to import or click Select all.
- 6. Click *Import*. If symbols with the same name are already loaded, you will be notified and given the option to overwrite.





### 4.4 Kriging 'low density' data in PAT

The mapping protocol which forms the 'backbone' of the PAT workflow is designed to handle data collected at a high spatial density using an on-the-go sensor – such as a yield monitor, proximal canopy sensor or electromagnetic soil sensor. Sometimes, there will be a need to collect data using hand sampling – for example using biomass cuts in a wheat field, counts of bunch numbers in a vineyard or assessments of disease – and for mapping these. Whereas the interpolation of high density, on-the-go data uses local block kriging, for hand sampled data collected at much lower density, global point kriging is more appropriate. This can be done in VESPER, but requires more user input, not least because the 'variogram' which underpins interpolation using kriging, needs to be generated manually and carefully. Additional information on this can be found in the VESPER manual available here with additional information on geostatistics available in books such as Webster and Oliver (2007). These notes are intended to guide the generation of the variogram and other input that PAT needs to run VESPER for low density data sets. Note for this current release of PAT, the procedure requires some use of VESPER independently of PAT.

#### Important notes

- It is not the intention in these notes to provide training in geostatistics. Users wishing to gain a better understanding of this topic should consult appropriate texts such as Webster and Oliver (2007). However, in general, we do not recommend that PAT users engage in low data density map interpolation unless they have acquired some prior understanding of map interpolation using kriging.
- It should also be noted that we DO NOT recommend interpolation of maps from 'low density' data when fewer than 100 data points are available (Webster and Oliver, 2007).
- Experience with work in vineyards (3-10 ha) suggests that a sample density of 26 samples or data points per hectare leads to satisfactory maps but a general rule of thumb is that higher data densities give better maps.

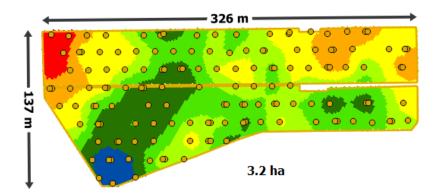


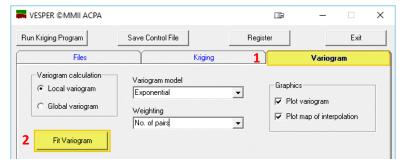
Figure 9 A map showing the data points used for the low data density kriging of 140 measures of grape berry weight in a 3.2 ha vineyard along with the final kriged result. This is the example used for the variogram fitting below.

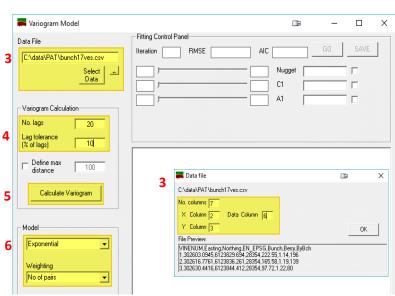
### Requirements

- A CSV file containing separate columns for each of X coordinates (e.g. Easting), Y coordinates (e.g. Northing) and the data value(s) to be mapped.
- A VESPER grid file created using the Create Block Grid Tool
- The coordinate system for both the CSV file and VESPER grid file. Both these files should have the same coordinate. The Clean, Trim, Normalise Point Data Tool can be used change the coordinate system of the CSV file to match the VESPER grid file described in 2.2.
- **VESPER:** Run Kriging Using VESPER tool requires installation of VESPER (see section 1.3.2)

### **Setup the Variogram**

- 1. Open VESPER independently of PAT and QGIS (i.e. as a stand-alone application).
- 2. In the Variogram tab, click Fit Variogram.
- 3. Browse to your csv file and in the Data files dialog that will appear, select the columns representing your X, Y and data value. For this example the X (Easting) is in column 2, Y (Northing) column 3 and data value (Berry) column 6.
- 4. Change the Variogram computation to
  - Number of Lags to 20 a.
  - Lag tolerance (%) to 10
- 5. Ensure the Variogram Model is Exponential and Weighting is No. of pairs.
- 6. Click Calculate Variogram to calculate and display the variogram.





### The Variogram

A variogram is a function which describes the variation in a dataset as a function of the distance between the points which comprise that dataset. It is key to the kriging process because the weights used to interpolate a map surface from sample data is a function of the distance between nearby data points and the location of the points for which map values are being interpolated. These weights derive from the variogram. Both Webster and Oliver (2007) and the VESPER Manual (Minasny et al., 2005) provide further information

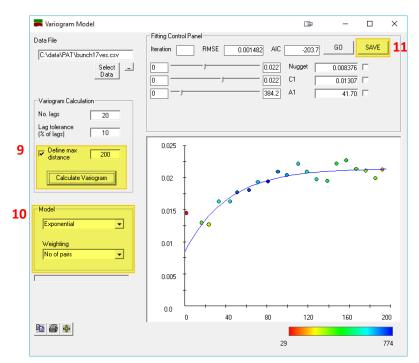
- 7. The colour coding in the variogram plot relates to the number of pairs of samples that have been used (or are available) to calculate the 'semivariance' of samples separated by each lag (i.e. distance) class.
- 8. In this example, it can be seen that beyond a distance of about 200 m, the number of pairs of points drops off. Because these points are at large lags, and the important part of the variogram is the sloping part at shorter lags, it can often be useful to set the maximum distance. The distance will vary between datasets.

When a maximum distance is set, it should reflect the drop-in pairs of points and should also be larger than the apparent range of spatial dependence (see Webster

Variogram Model Fitting Control Panel Data File SAVE RMSE [ -197.0 0.001739 AIC Iteration C:\data\PAT\bunch17ves.csv Select \_\_\_ 0.025 Nugget 0.008448 F C1 0.025 0.01403 0 491.4 49.49 Variogram Calculation No. lags 20 0.03 8. decreasing pair count Define max Calculate Variogram 0.024 Model 0.018 Exponential -Weighting 0.012 • No of pairs 0.006 ₽ **4** <sup>33</sup>7 coloured by pair

and Oliver, 2007 for further discussion of the form of the variogram), and also needs to be sensible in relation to the size of the field being considered. For example, if your vineyard being mapped is a 1 ha square of 100 m side, one would generally expect the maximum distance to be less than 100 m.

- For this dataset, tick the Define max distance and enter 200 m, then click Calculate Variogram to update the variogram.
- 10. To this point, the default settings of an Exponential variogram model have been used with a weighting for the fitting of the curve dependent on the No of pairs. Often, a curve which describes the data a little better might be obtained by selecting a different model and/or using a different weighting method. In general, and with the exception of advanced users, we would recommend that the variogram model selection is confined to either Exponential (the default) or Spherical and that the Weighting used



is either No of pairs or No\_pairs/st\_dev. Further information on settings is available in the VESPER Manual (Minasny et al., 2005).

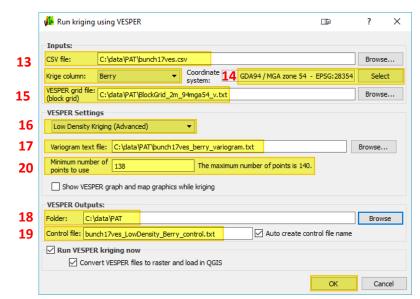
11. Once satisfied with the fitted variogram, click *Save* to save variogram parameters to file. It is useful if the output has the same prefix as the CSV file with the name of the data column you used in step 3 and variogram is added to the end (e.g bunch17ves\_berry\_variogram.txt). This is the input variogram file for the low density kriging option in *Run Kriging Using VESPER Tool*.

### **Run Kriging using PAT**

- 12. Close VESPER, launch QGIS and the open then dialog for Run Kriging Using VESPER Tool from the PAT toolbar or menu. Further details on this dialog can be found section 2.3. The Low Density Kriging implementation is as described 🗞 🚜 🖫 😓 🏉 🤣 🔗 🥢 🚜 🚓 🦈 🂰 🎱 below.
- 13. Select the same *CSV file* and *Krige* (data) column used in step 3
- 14. The coordinate system will be automatically determined if a column (EPSG or ENEPSG) exists in the CSV file. If this does not exist then you will be required to set it prior to continuing.

If your CSV file needs reprojecting, the you can use the Clean, Trim, Normalise Point Data Tool using the settings described in 2.2.

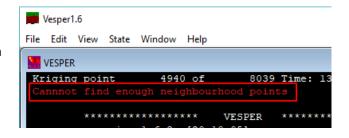
15. Browse to the VESPER grid file.



At this point a dataset overlap check will be completed if you have provided the CSV and VESPER grid file where it is assumed that these two files have the same coordinates system. If this fails you will be warned as shown. No overlap will cause VESPER kriging to fail.

- 16. Choose the Low Density Kriging (Advanced) method.
- 17. Browse for the *variogram text* file saved in step 11.
- 18. Specify the *output folder* to store the VESPER files and results in. A sub-folder called VESPER will be automatically created if it does not already exist.
- 19. If required change the default control file name. This is saved into the folder from step 18.
- 20. Use the default (total point count minus 2) minimum number of points to use during VESPER kriging.

In some cases, this will cause a Cannot find enough neighbourhood points error in VESPER and create an empty output. If this occurs re-run the kriging by repeating steps 13-21 and decrease the *minimum number of points to use* by one and try again. You may have to re-try multiple times to find the right value.



21. Click OK and the control file will be created and added to the VESPER queue. When the VESPER processing is completed, the resulting kriged text file will be converted to a TIF and loaded into QGIS.

## 5 Contributors

The following individuals have contributed the production of PAT and the underpinning pyprecag Python package, through coding, configuration and expertise in PA, and in the case of the Whole-of-block analysis tool, the development of statistical methodology.

Core Project Team: Christina Ratcliff (lead developer), Rob Bramley (project leader) and David Gobbett (technical lead)

Whole-of-block analysis tool: Shuvo Bakar, Warren Jin and Brent Henderson pyprecag packaging and publishing: Andrew Spiers, Adrian D'Alessandro, David Benn and Daniel Collins

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