

PAT - Precision Agriculture Tools Plugin for QGIS

User Manual

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Enquiries

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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.

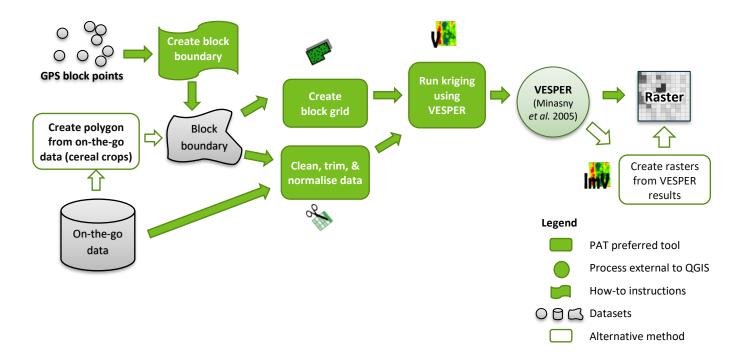


Figure 1 The basic map production workflow used in PAT and some of the existing tools available for processing high density on-the-go data.

Table 1 Brief description of tools available in PAT

	PAT TOOLS	DESCRIPTION
	Create Block Grid	 Convert polygon features (e.g. a block boundary – see page 52) to a TIFF raster and a VESPER text grid file of X,Y point values used by VESPER.
	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.
V	Run Kriging Using VESPER	Create a VESPER control file and data files and run VESPER kriging.
lmV	Import VESPER Results	Convert VESPER outputs to raster TIFF format.
W	Create Polygon from On-The-Go GPS Point Trail Data	Generate a polygon block boundary from on-the-go data (e.g. from yield monitors) containing GPS points. The GPS points must cover the entire block (not just be a set of points around the boundary - see page 52 for instructions on creating a boundary from a set of surveyed data points)
	Rescale or Normalise a Raster	 Create rasters by rescaling (standardising) values between a fixed range (i.e. 0-1, or 0-255) normalising values to a mean of 0 and standard deviation of 1
4 :	Generate Random Pixel Selection	Generate a selection of random pixels from a raster and save to a points Shapefile.
*	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 k-means clustering
		Tool for analysing simple strip-based experiments.
	Create Strip Trial Points	It generates pairs of points from a central line within a strip and 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 method.
A t	Run Strip Trial t-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.
	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.

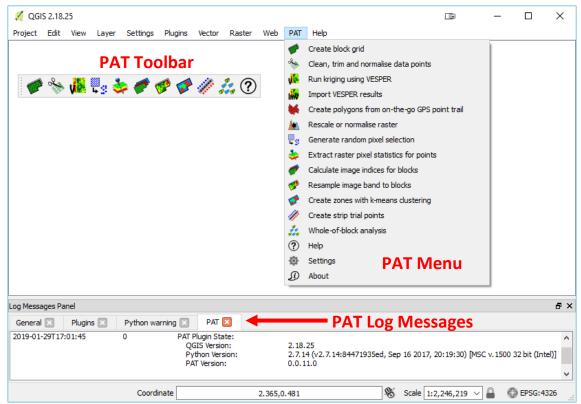


Figure 2 PAT toolbar, menu and log messages panel within QGIS

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

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://sydney.edu.au/agriculture/pal/software/vesper.shtml. **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.

QGIS LTR 2.18.21-26: Download and install the QGIS standalone long term release (versions 2.18.21 to 2.18.26). Note that the PAT tools do not currently work with QGIS version 3 or later.

QGIS 2.18.26 can be downloaded directly using these links.

32bit: http://download.osgeo.org/qgis/win64/QGIS-OSGeo4W-2.18.26-1-Setup-x86.exe 64bit: http://download.osgeo.org/qgis/win64/QGIS-OSGeo4W-2.18.26-1-Setup-x86_64.exe

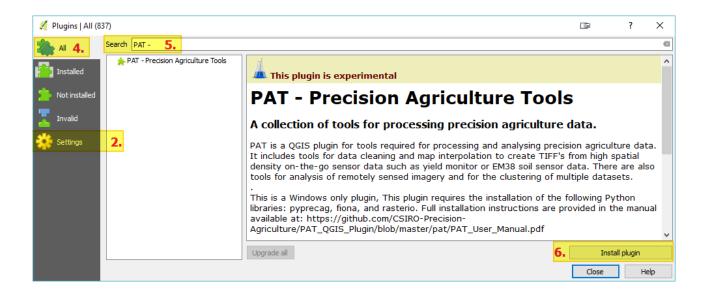
Additional Python Packages: QGIS includes Python and numerous Python packages. However, PAT requires the following additional packages: *pyprecag* (Ratcliff *et al.* 2019), *Fiona* (Gilles *et al.* 2011) *and Rasterio* (Gilles *et al.* 2013). The *Fiona* and *Rasterio* packages have been bundled with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

1.3 Installing PAT

Install the PAT - Precision Agriculture Tools Plugin.

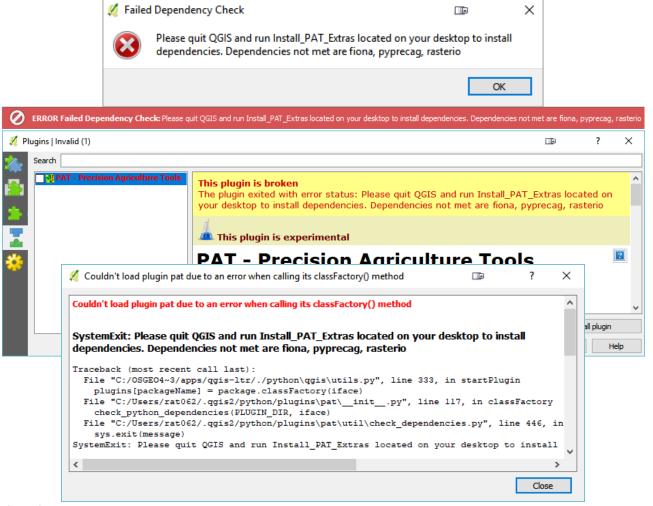
- In QGIS open the plugin manager.
 Plugins Menu → Manage and Install Plugins
- 2. Select the **Settings** side tab.
- 3. Tick *Check for Updates* and *Show Experimental Plugins* options.
- 4. Select the **All** or **Not installed** side tab.
- 5. Search for and select **PAT Precision Agriculture Tools.**
- 6. 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*.

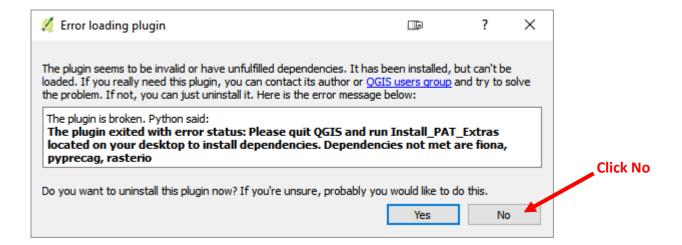


Installing or Upgrading PAT Python Dependencies

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.



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_PAT_Extras*. Double click the shortcut to run it.



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

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.

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.

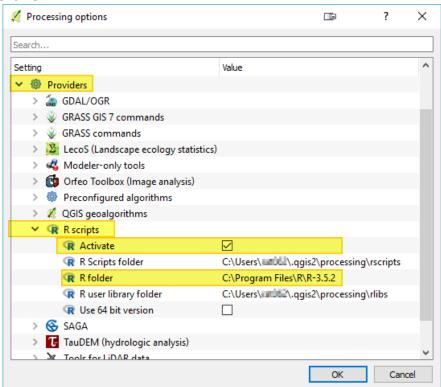


To resolve this, complete the following steps to configure R and QGIS

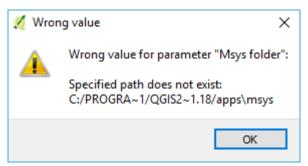
- 1. Download R from Windows version can be downloaded from http://r-project.org/. The minimum R version suitable for this is v3.5.1.
- 2. Install using all the default options
- 3. If required, activate the **Processing** plugin in QGIS plugin manager.
 - a. In QGIS open the plugin manager. Plugins Menu → Manage and Install Plugins
 - b. Select the *Installed* section
 - c. Find **Processing** and ensure the adjacent checkbox is ticked.
 - d. Close the Plugins dialog.

This will give you a processing menu and the processing side panel.

- 4. From the *Processing* menu select *Options*.
 - a. Expand Providers
 - b. Expand *R scripts*
 - c. Tick on Activate
 - d. Ensure that the R installation folder is shown against *R folder*
 - e. Click OK.

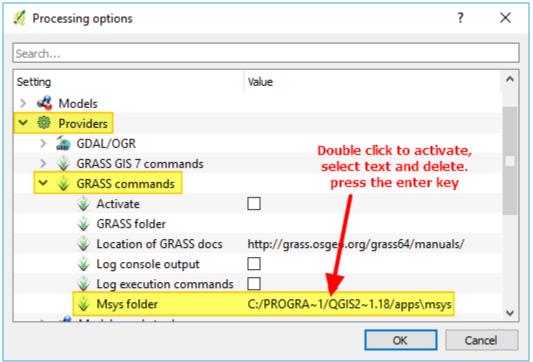


When clicking OK you may get a **Wrong value** error message like this



and your mouse cursor will remain a circle. To resolve treat the mouse circle as an arrow and

- 1. Under **Providers**, expand **GRASS commands**
- 2. Double click the text to the right of Msys folder, select and delete it and press the enter key.



- 3. Click OK
- 4. Close and re-open QGIS to reset your mouse cursor. It should now all work.

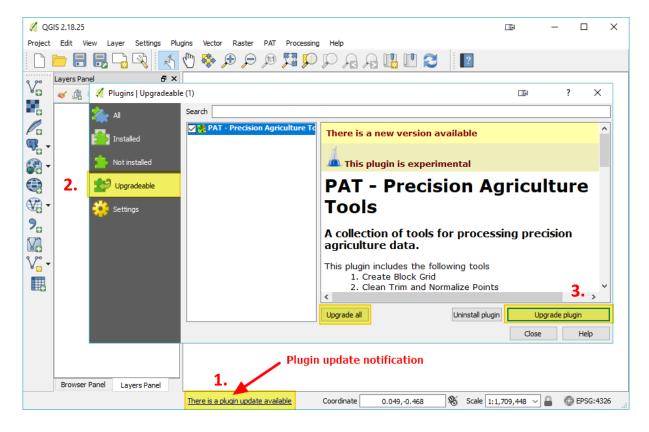
1.4 Update PAT

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 → 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.

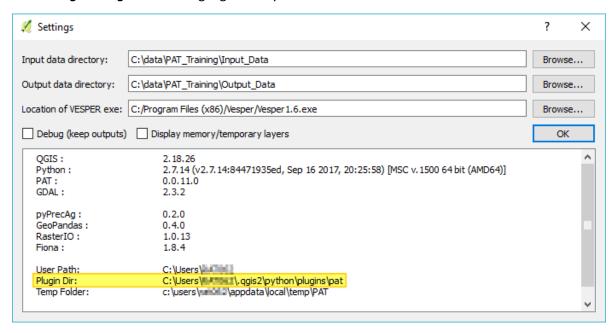


1.5 Uninstall PAT

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. This folder is listed in as Plugin Dir in the settings dialog as shown highlighted in yellow 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 PRIOR 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

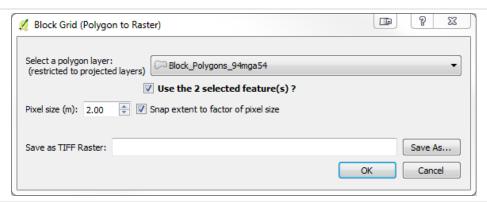
This tool converts polygon features such as a block boundary to a raster grid using a set pixel size. This is a critical part of setting up the analysis environment for creating maps from on-the-go data because it generates the base grid onto which maps will be interpolated, by using the outer most extent of the block boundary as the 'frame' for the grid. Refer to instructions on page 52 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 54 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 in a projected coordinate system 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 Sugarcane: 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.				
Save as TIFF raster	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:

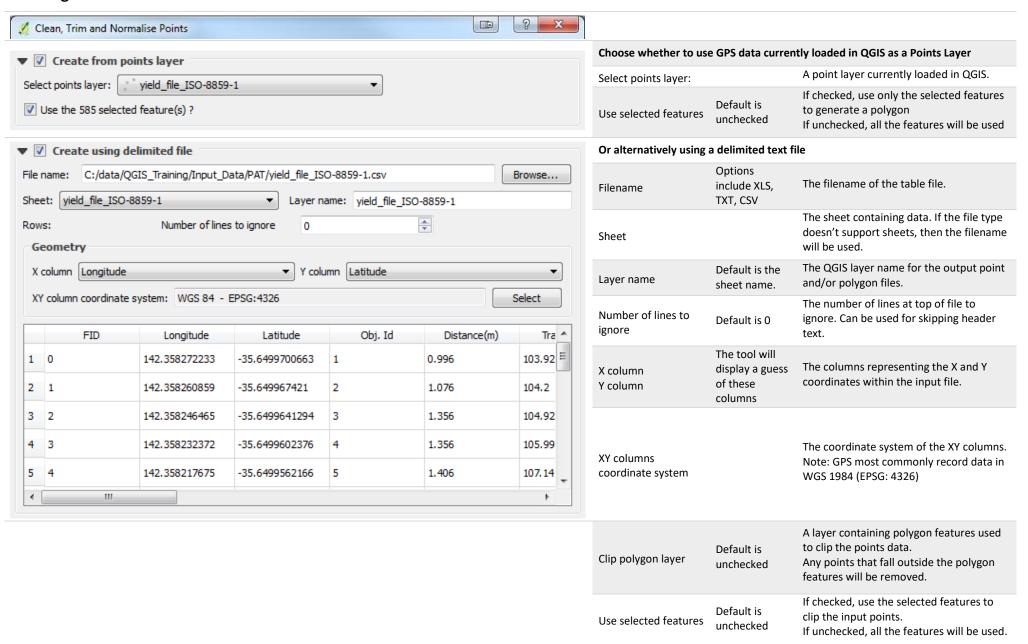
- retains all columns (except coordinate columns) from the original file.
- converts coordinate columns to a projected coordinate system and renames them to Easting and Northing. An additional column (EN EPSG) will be created and assigned the EPSG number for the projected coordinate system used to reproject 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

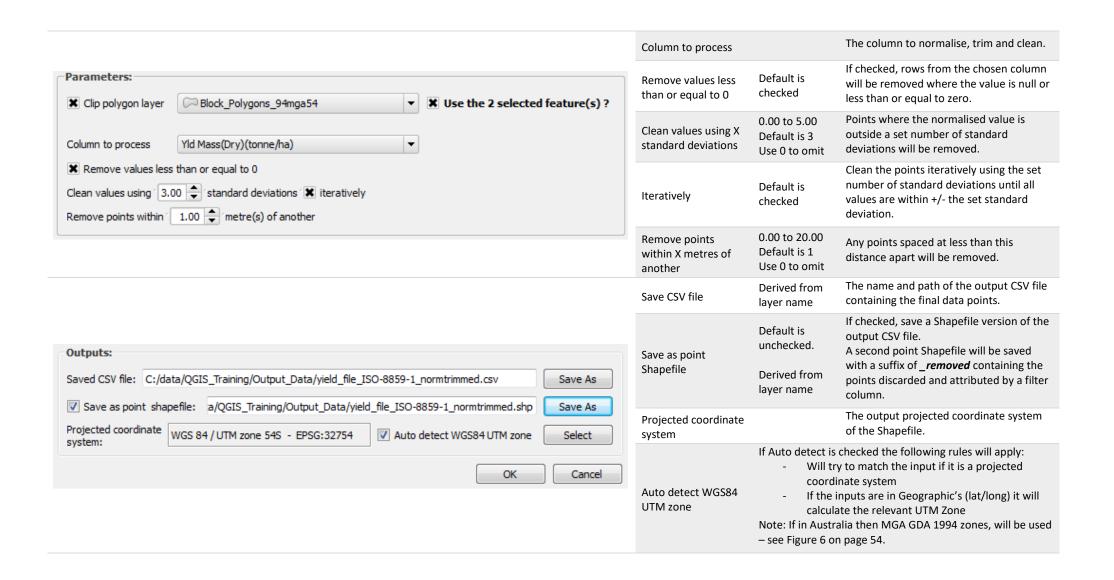
This tool filters data by:

- 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 51.

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.



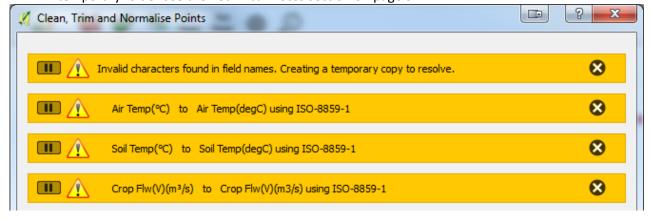


Notes

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 1 m
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 1 m
	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 51.





Run Kriging Using VESPER

Summary

This tool is used for mapping on-the-go data, 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.

Any files matching the control file name will be deleted from the VESPER sub-folder including VESPER kriged results, to ensure that the files remaining belong to the newly create control file. If you want to retain old versions, use a different name for the control file.

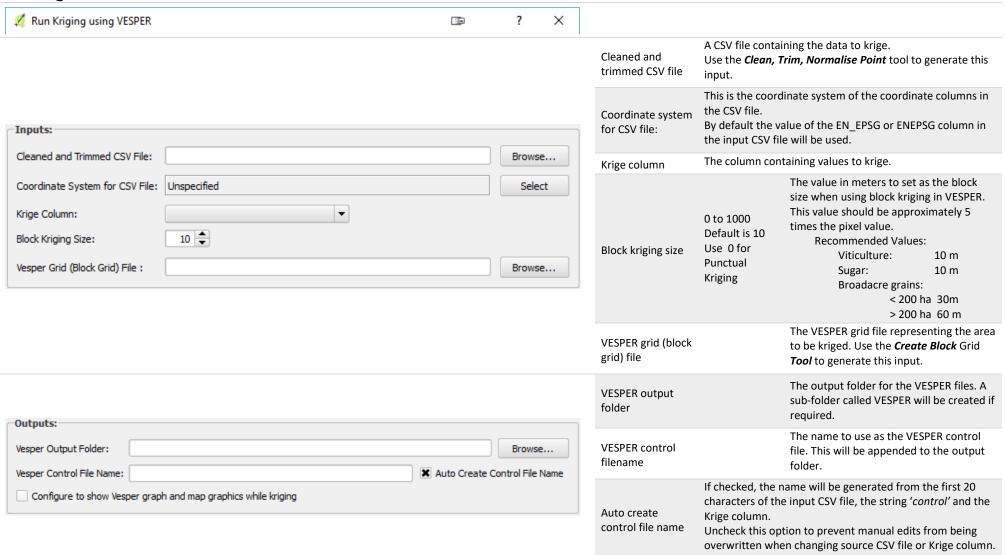
The coordinate system of the CSV file will be determined by interrogating the EN EPSG or ENEPSG columns generated as a by-product of running the Clean, Trim, Normalise Point tool. If the resulting value cannot be found or is incorrect, the coordinate system can be set manually. The resulting EPSG number will be stored within the control file and later used when converting VESPER outputs into raster TIFF format.

To krige (run) VESPER control files from within QGIS, VESPER must already be installed on the PC and configured in the PAT settings (see Settings Tool on page 50).

If the Run VESPER Kriging Now option is checked the output control file will be launched in VESPER and kriged immediately. Running this tool multiple times will add each output control file to the VESPER queue and run consecutively. The QGIS status bar is used to manage the queue; currently the queue can only be displayed and cleared. On the completion of each VESPER run, if the Convert VESPER Files to Raster and **Load in QGIS** option is <u>checked</u> the kriged results will be import to TIFF files and loaded into QGIS.

If the Run VESPER Kriging Now option is unchecked the tool will create the VESPER control, data and batch files in the selected folder allowing the user to run VESPER in their own time.

When Run VESPER Kriging Now or Convert VESPER Files to Raster and Load in QGIS options are unchecked the Import VESPER Results (post VESPER) tool can be used to import VESPER kriged results to TIFFs then loaded into QGIS.





File Naming Conventions

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

DESCRIPTION	FILENAME	EXTEN SION	EXAMPLE				
Files <u>required</u> by VESPER and copied to the VESPER output folder							
VESPER control file	<pre><first 20="" characters="" csv="" file="">_< first 10 characters of krige column>_control</first></pre>	.txt	swblock_YldMassDry_control.txt				
VESPER grid file	<control file="" name=""> where control is replaced by vespergrid</control>	.txt	$swblock_YldMassDry_vespergrid.txt$				
VESPER data file	<control file="" name=""> where control is replaced by vesperdata</control>	.csv	swblock_YldMassDry_vesperdata.csv				
Files <u>created</u> by VESPER and saved to the VESPER ou	tput folder						
VESPER Krige file (contains Prediction and SE)	<control file="" name=""> where control is replaced by kriged</control>	.txt	swblock_YldMassDry_kriged.txt				
VESPER parameter file	<control file="" name=""> where control is replaced by parameter</control>	.txt	$swblock_YldMassDry_parameter.txt$				
VESPER report file	<control file="" name=""> where control is replaced by report</control>	.txt	swblock_YldMassDry_report.txt				
If converting VESPER files to raster is checked or the	e Create Rasters from VESPER Results tool is used						
Kriged Prediction TIFF source: VESPER Krige result above	<control file="" name=""> where kriged is replaced by PRED</control>	.tif	swblock_YldMassDry_PRED.tif				
Standard Error TIFF source: VESPER Krige result above	<control file="" name=""> where kriged is replaced by SE</control>	.tif	swblock_YldMassDry_SE.tif				
Confidence Interval (CI) metadata text file	<control file="" name=""> where control is replaced by Cl</control>	.txt	swblock_YldMassDry_CI.txt				



2.4 Import VESPER Results

Summary

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

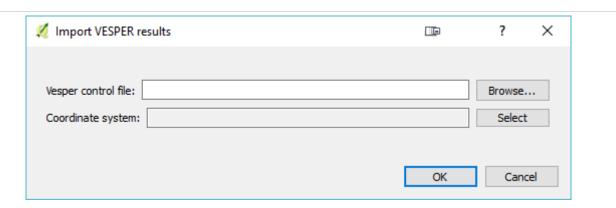
The output filenames will be created using the VESPER control file name as a base and demonstrated in the *Prepare Data for Kriging using VESPER Tool - File Naming Conventions* section on page 22.

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.

Dialog



Import VESPER Resul	mport VESPER Results					
VESPER Control File:		A VESPER control file used to identify VESPER results files and convert to rasters.				
Coordinate System for CSV File:	By default it is extracted from the input control file.	This is the coordinate system of the VESPER results.				

See Prepare Data for Kriging using VESPER File Naming Convention

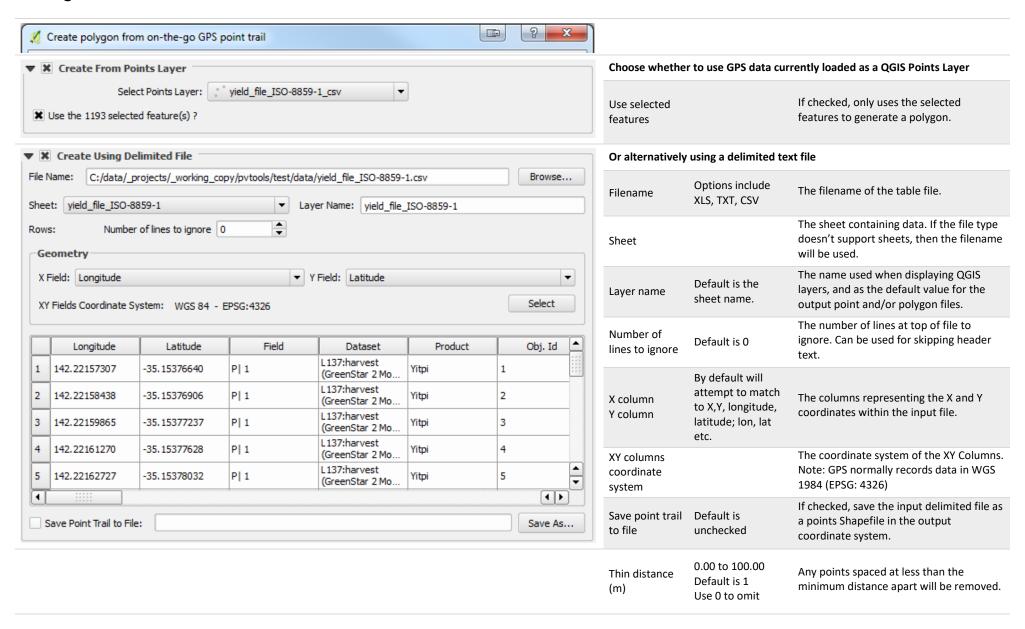


2.5 Create Polygon from On-The-Go GPS Point Trail Data

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 52 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.

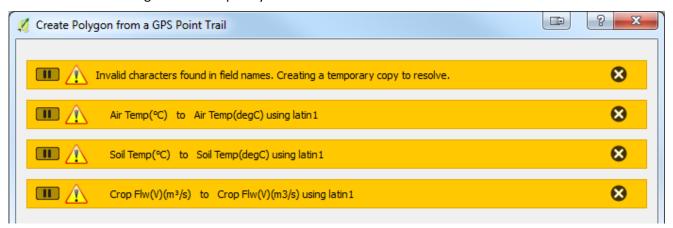
As the process involves a dot-to-dot approach, it is critical that the input file of points are in order (i.e. sorted by an increasing time sequence - as would be normal for data collected by an on-the-go sensor such as yield monitor). For efficiency, points can be thinned by removing points closer than 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.



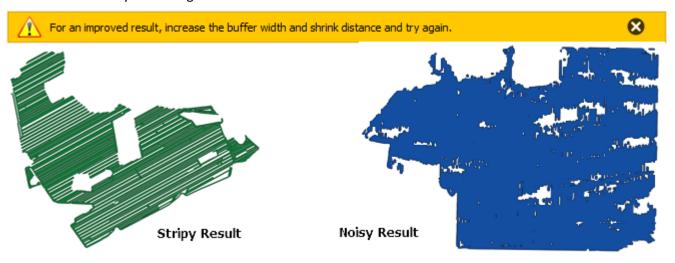


Notes

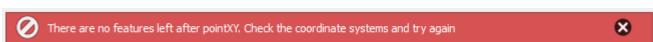
The following notification occurs when a delimited text file contains invalid characters in the column name. A copy of the file will be made in the PAT temporary folder containing corrected column names as specified in the PAT log message and to file. For more information on the location of the saved 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 all in the range 0 to
- 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.

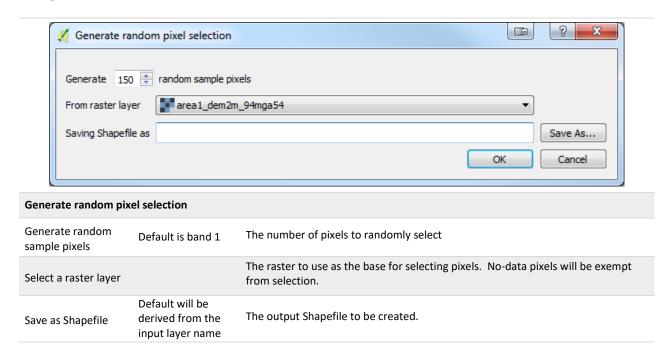


Rescale or normalise a raster						
Select a raster layer for band X	Default is band 1	The raster layer and band to rescale or normalise.				
Method	Default is rescale	Options: Rescale – adjust values to a fixed range Normalise – adjust values to a mean of zero and a standard deviation of one				
Rescale between (when selected)	Default 0 to 255	The range of values used with rescaling.				
Save as Shapefile	Default will be derived from the 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			X			7	
		5		X		5		
			3	X	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 statistics for points ?	×			
Select a points layer: ▼		Select point layer		A point layer in QGIS used to extract pixel statistics values.
No features selected		Use selected features	Default is unchecked	If checked , extract pixel values only for the selected points. If unchecked , all the points will be used
Use raster layer Only process rasters with one pixel size. Adding the first raster layer will set this pixel size 0 Raster(s) Move Up Move Dov Remove	wn	Use raster layer	Only raster layer a time and this i The Add, Delete, used to manage The order of ras	to extract values from. rs of the same pixel size can be analysed at s set by the first raster layer added. , Move Up, Move Down, buttons can be the list of rasters. ters is retained in the output CSV file with ns for each raster and statistic combination.
Apply to all rasters Select one or more statistics to calculate using one of the following pixel neighbourhood sizes Include current pixel value (uses 1x1 neighbourhood by default) Mean Minimum 3x3 7x7 Standard deviation Maximum 5x5 9x9		Calculate pixel neighbourhood statistics	the list. Current pixel just and/or Statistics to be of	red statistical methods for each raster in st extracts the pixel value calculated using the set neighbourhood size. mn naming format is explained below.
Coefficient of variation Pixel count		Neighbourhood size (pixels)	Default is 3x3	The size of the neighbourhood footprint to use while calculating statistics.
Saved CSV file: Save As 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 on page 14.

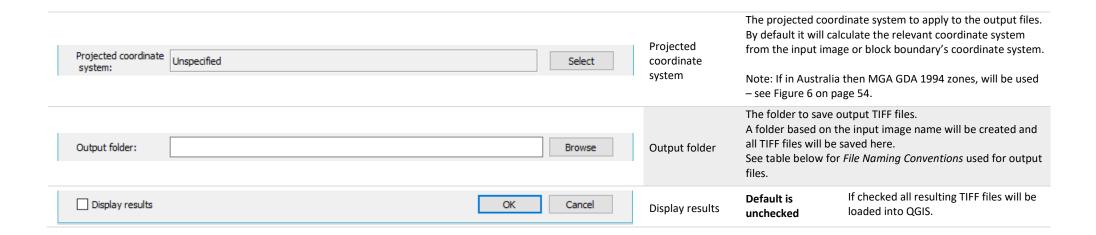
Table 3 The list of supported indices

Normalized Difference Vegetation Index (NDVI) Green Normalized Difference Vegetation Index (GNDVI) Normalised Difference Red-Edge Index (NDRE)

Plant Cell Density Index (PCD) Chlorophyll Red Edge Index (ChIRE) (Gitelson 2004; Gitelson et al. 2005)

$$\begin{aligned} \text{NDVI} &= \frac{(NIR - Red)}{(NIR + Red)} \\ \text{GNDVI} &= \frac{(NIR - Green)}{(NIR + Green)} \\ \text{NDRE} &= \frac{(NIR - Red \, Edge)}{(NIR + Red \, Edge)} \\ \text{PCD} &= \frac{NIR}{Red} \\ \text{Chl}_{red \, edge} &= \left(\frac{NIR}{Red \, Edge}\right) - 1 \end{aligned}$$

✓ Calculate image indices for blocks	Tip	?	×			
Select image:			•	Select Image	The image layer co- for calculate indice	ntaining the appropriate bands required s.
Nodata value: 0				No-data value	Default from image.	Can be used to specify a different nodata value.
Use a block			~	Use a block boundary	Default is unchecked	A layer in QGIS containing polygon(s) representing blocks.
boundary No features selected				Use selected features	Default is unchecked	If checked , only the selected polygons will be used.
Block ID column:			•	Block ID column		ng the block id or name. This will be used gons with the same block id or name as
Resample to 2.00 • metre pixels				Resample pixel size (m)	0.00m to 6 km Default is 2 m	The pixel size to apply to the raster outputs. This is expressed in meters. Recommended Values: Viticulture: 2 m Sugarcane: 2 m Broadacre grains: < 200 ha 5 m > 200 ha 10 m
Specify image bands used for index calculations Green ▼ Red-Edge ▼ □ ND	the indices to calculate VI			Specify image bands used for index calculations	will enable/disable requirements. For most images Green is l	umbers to band types. The mapped bands indices based on individual index band 2, Red is band 3, ared is band 4
Red ▼ Near Infrared ▼ GN Non-vine mask ▼	DVI Chire					band where pixels <u>not</u> containing vine o-data. Not required for broadacre data.
				Select the indices to calculate	haven't been mapp	ed (greyed out) the bands it requires



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 250cm pixels for the Plant Cell Density (PCD) Index created without specifying a block id column.



2.10 Resample Image Band for Blocks

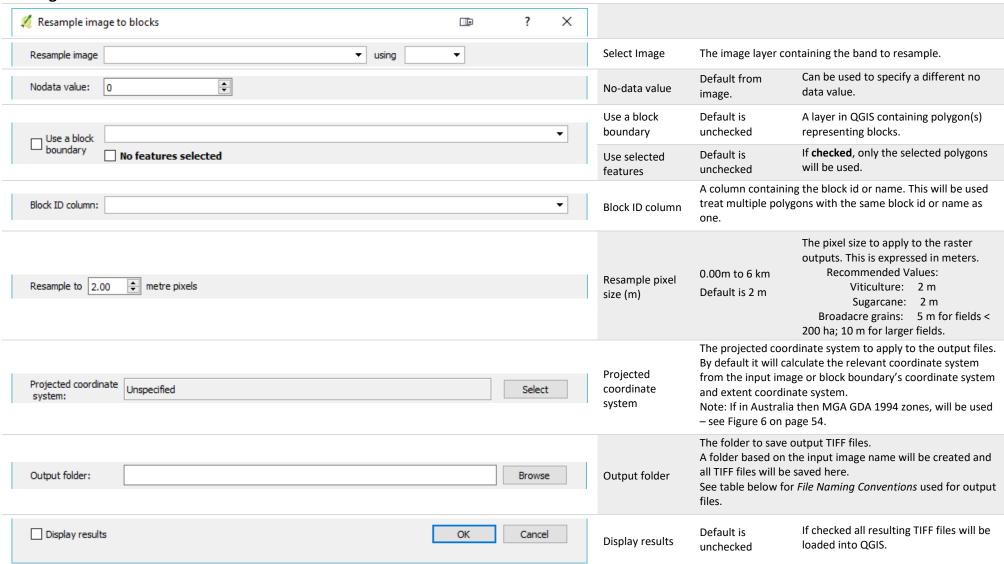
Summary

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* on page 14.

Dialog



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:	Image Names:				
 	B1_Band6_2m.tif	The TIFF file where Band 6 is resampled to 2m for block id/name of B1.			
	Band7_250cm.tif	The TIFF file where Band 7 is resampled 250cm pixels without specifying a block id column.			



2.11 Create Zones with k-means Clusters

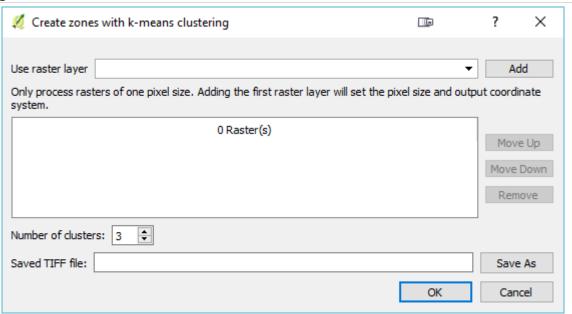
Summary

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.

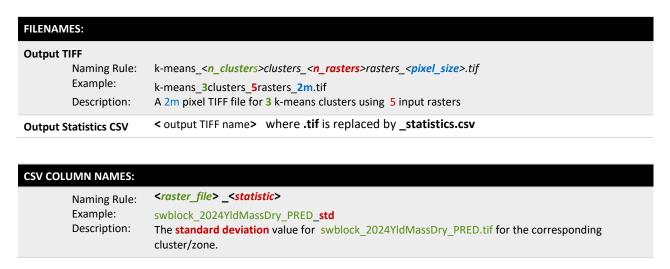
Dialog



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. 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 Default is 3 Number of clusters: appropriate. The name and path of the output TIFF file representing the zones. In addition, a statistics CSV file will be written to disk along with the TIFF, and results printed to PAT's log Save TIFF file messages panel.

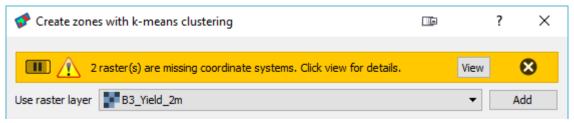
File and Column Naming Conventions

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

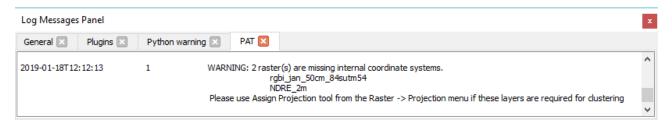


Notes

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.





Summary

This tool is used to generate the data needed to implement the moving window t-test method of Lawes and Bramley (2012) for analysis of simple strip trials. It may be useful to users who, for example, have implemented 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 using the line nominated distance between points. In addition, two sets of points are placed at the offset distance either side of the line as shown in Figure 4 which will be used as a control. This allows the treatment strip data, to be compared to data from the neighbouring (control) area.

Each point has values assigned to the columns Strip Name, TrialID, PointID and DistOnLine. Treatment strip line will be assigned a unique identifier in the column TrialID and the value of 'Strip' in the Strip Name column. Points will be 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. The PointID of the control points are paired with the corresponding treatment point. The value of the Strip Name column is assigned based on the cardinal direction of the control offset line as shown in Figure 4. The DistOnLine represents the distance in metres of the point from the start of respective line. The distances for the control strips are calculated from the start of the control line, not the strip line.

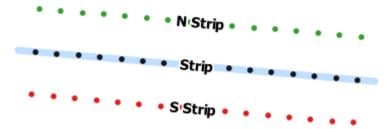
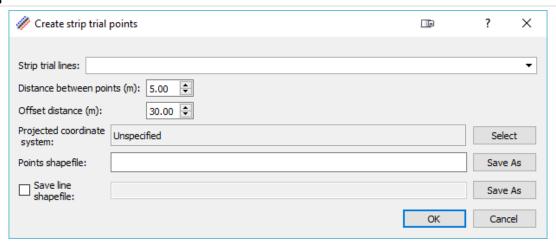


Figure 4 Output from the Create Strip Trial Points tool. the Blue line represents the treatment strip and the dots; the respective treatment and control points

Dialog



Create strip trial points

The layer containing the strip trial lines. Strip trial lines:

•					
Distance between points (m):	1 – 500.00 Default is 5.0	The distance used to space points <u>along</u> the line. In this instance, <u>users are advised</u> to select a value appropriate to their trial rather than accepting the default.			
		If used in conjunction with a strip trial t-test then it should be the pixel size of the input rasters used for the t-test.			
		The distance used to offset the points <u>from</u> the line.			
Offset distance (m)	1 – 500.00 Default is 30.0	If used in conjunction with a strip trial t-test then this should be far enough away from the strip trial to remove the trials treatment influence			
	•	dinate system to apply to the output files.			
Projected	By default it will calculate the relevant coordinate system from input layer's coordinate system.				
coordinate system:	Note: If in Australia	then MGA GDA 1994 zones, will be used – see Technical Notes on page 54.			
Points shapefile:	Derived from layer name	The name and path of the output points shapefile.			
	Default is				
Save line shapefile:	unchecked. Derived from	The name and path of the output points shapefile.			
	layer name.				



Summary

This tool enables a user to analyse strip trials to determine the effectiveness of effects within the strip. For example, it could be used analyse the value of a N-Rich strip treatment.

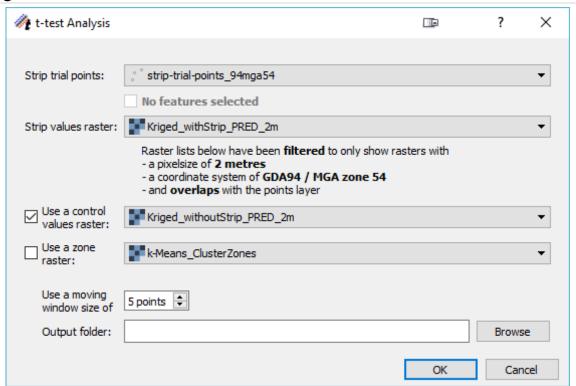
Using the Create Strip Trial Points values will be extracted from input rasters and a pairwise calculations for a moving average t-test, Response Index(RI) and treatment differences are calculated and graphed as described in Lawes and Bramley (2012).

The input rasters include

- Strip values raster: A map produced through following the high density on-the-go sensor mapping workflow (Figure 1)
- Control values raster (optional): A map produced through following the high density on-the-go sensor mapping workflow after points relating to the strip trial have been removed.
- Zone raster (optional): A zone raster. This can be created using the *Create Zones with k-means* Clusters tool.

For each input strip a map image will be produced long with a graph and CSV file, for each strip combinations 1) Strip vs Control 1, 2) Strip vs Control 2, 3) Strip vs Average of Control 1 & 2.

Dialog



Run Strip Trial t-	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</i> The raster layer containing values to extract for analysis				
Strip values raster:					
Use a Control Default is The raster layer values raster: unchecked		The raster layer created after treatment points have been removed.			
Use a Zone raster:	Default is unchecked	The raster containing the zone values. If not provided a zone of 1 will be applied to the entire strip.			
Use a moving Derived from The name and path of the window size; layer name		The name and path of the output points shapefile.			
Output folder		e output map, graph and CSV files. 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 (_).

	A map showing the Trial points
ming Rule:	line <trialid>_map.png</trialid>
mple:	line3_map.png
scription:	The map for the strip with a TrialID of 3
	The results displayed as graphs
ming Rule:	line_ <strip>_graph.png</strip>
mple:	line3_N-S-strip_graph.tif
scription:	The graph result for Trial 3 which compares the N-S control strip to the treatment strip
3	mple: cription: ning Rule: mple:

CSV files: The csv file used to create the graphs.

> Naming Rule: line<**TrialID>_<strip>**.csv Example: line3_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:	
<strip> Value</strip>	The number extracted from the Strip value raster .
<strip> Control</strip>	The number extracted from the Control values raster.
	Applies only when a Control values raster is used.
<strip> Zone</strip>	The zone value extracted from the Zone raster .
	Creates a Zone value of 1 if not specified.
controls_mean	The mean of the input control strips for each point.
treat_diff	The difference between the treatment and <u>controls_mean</u> columns
av_treat_diff	The average of the column treat diff using a moving window
pep8 formatting - processing.py, table_ops.py 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



Summary

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 5. 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 5 – maps showing the response to the different treatments over the entire block, and maps indicating the significance of differences between these treatments.

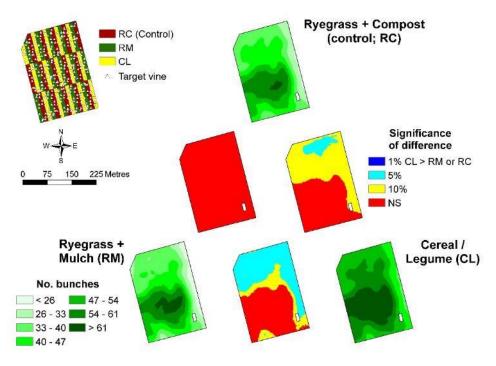


Figure 5 An example of a whole-of-block experiment conducted in a 4.8 ha Clare Valley vineyard. Panten and Bramley (2011) provide further information

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 5. 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.

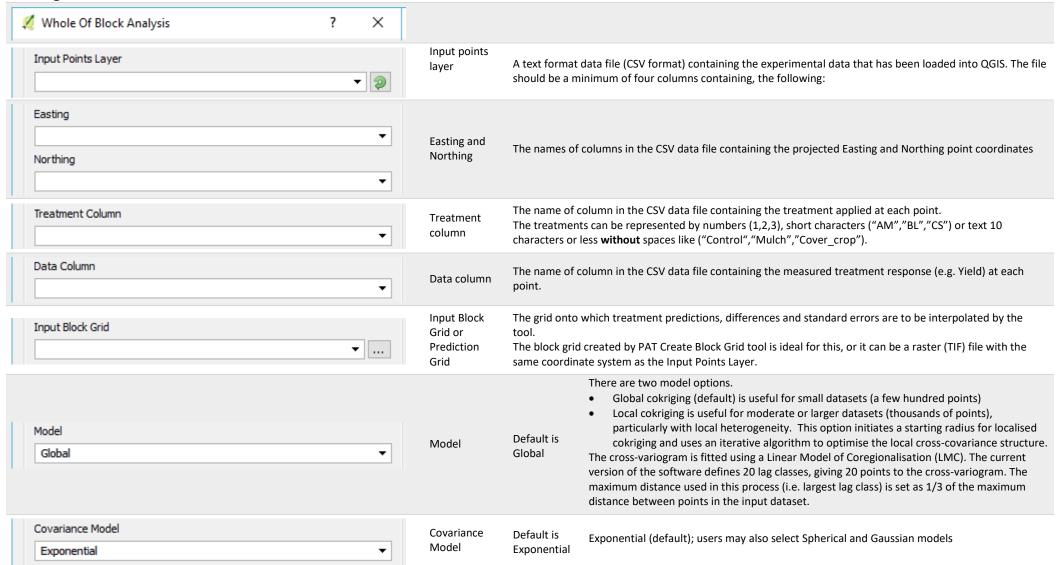
Algorithm for the Local Cokriging System – a brief commentary on the data radius used

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 (Jin et al., 2019).

Dialog



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

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
<pre><data col="">_tr_<tmt1>_<model><covariance model="">tif</covariance></model></tmt1></data></pre>	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
<data col="">_tr_<tmt1>_var_<model><covariance model="">tif</covariance></model></tmt1></data>	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.



Summary

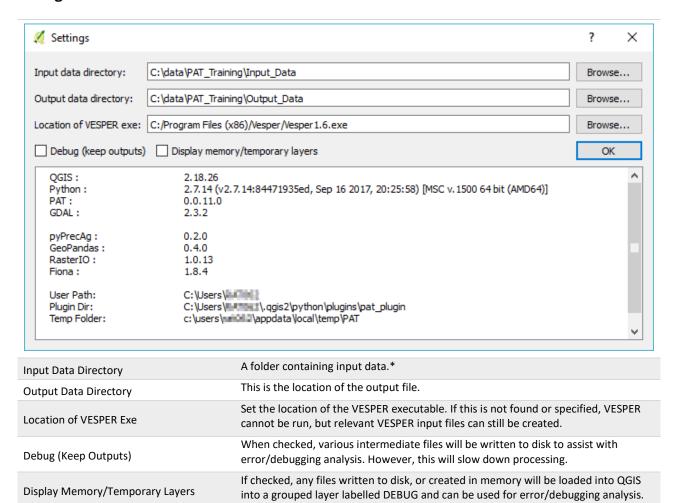
This tool is used to display and edit PAT settings.

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



^{*} By default this will be the PAT sub-folder in the user's home directory. To quickly navigate here type *%homepath%/PAT* in the address bar of Windows Explorer.

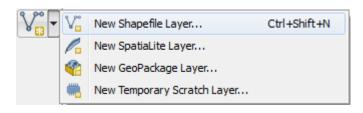
3 Technical Notes

- 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 are 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

 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.

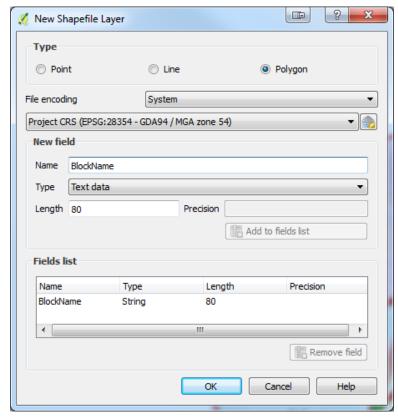


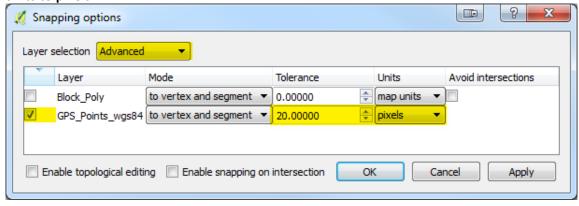
More information on coordinate systems can be found at

- 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
- 4. If required, load other vector or raster data, like imagery, which can be used as reference.
- 5. Setup your snapping environment.

WGS 84.

- a. Open *snapping* options (*Settings* menu -> *Snapping Options*)
- b. Change *layer selection* to *advanced*.
- c. **Tick** the layer containing the **point layer** loaded in step 3.
- d. Change the *tolerance* to 20 and set units to pixels





^{*}More information on selecting a coordinate system can be found on page 54.

6. Select/activate the **polygon** layer in the layers panel and click the **toggle editing** icon from the **digitizing toolbar**.

- 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 (+), 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.

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

To add, move or delete a vertex, toggle to node mode using the *node tool* 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.

To split a polygon use the **split features** tool from the **advanced digitizing toolbar** and sketch the 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 6 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)

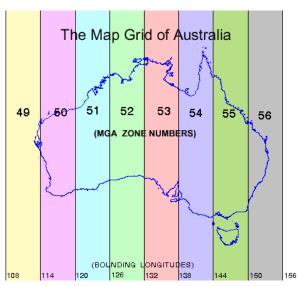


Figure 6 The Map Grid of Australia. (www.environment.gov.au/erin/tools/mga2geo-gda.html).

1. Determine your projected coordinate system.

Within Australia use GDA 1994 Map grid of Australia (MGA)

EPSG: 283xx

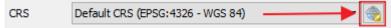
Outside Australia use the WGS84 UTM Zone system.

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.

2. Reproject your shapefile

- 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. c.



- 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

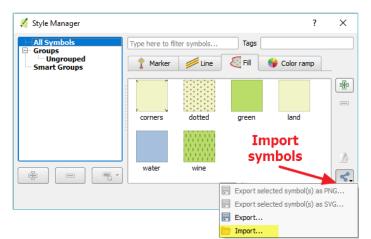
The shapefile will be reprojected and loaded into QGIS and is now ready for use. For further information on coordinate systems see

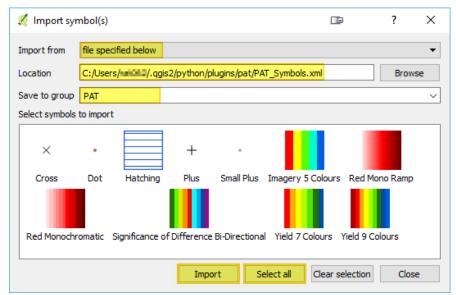
 $https://docs.qgis.org/2.18/en/docs/gentle_gis_introduction/coordinate_reference_systems.html$

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.

- 1. In QGIS, launch the **Style Manager** from the settings menu.
- 2. From the lower right 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 the Save to group.
- 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 Analysing strip trial data using the moving window t-test

Following use of the Create Strip Trial Points and Extract pixel statistics for points tools, the user should have an output dataset which contains a minimum of five columns: Columns of x and y coordinates corresponding to locations along the trial strip and three columns of data corresponding to the extracted pixel values from the strip 'centre line' and from right and left offsets to that centre line (see page 41). The data are the input to the moving window t-test of Lawes and Bramley (2012). These data can then be analysed in a spreadsheet or statistics package. For users of Excel, this may be done as described below. Note that in this example, we demonstrate testing of the difference between the strip and one offset line only and use a simple case in which no prior information (e.g. location of different management zones) is available. In practice, users may which to compare the strip with both of the offset lines or alternative use the mean of the two offset lines for comparison with the strip.

1. Data setup:

Column A Record number

Column B x coordinate or Easting Column C y coordinate or Northing Column D Data from the trial strip

Data from the offset comparison line Column E

It is assumed here that the column headings are in row 1 with the data starting in row 2.

2. Calculate a treatment difference

In Column F, insert a formula to calculate a treatment difference, e.g.:

Copy this formula down to the last row the dataset.

3. Select the size of your moving window.

The choice of this will need to be considered in relation to the grid pixel size, the length of the trial strip, and the distance setting used in the Create Strip Trial Points tool. In the example given in Lawes and Bramley (2012), they extracted values from a yield map at approximately 5 m intervals along their strip, which was approximately 700 m in length, and then used a window of 5 data values (i.e. approximately 25 m) for the analysis.

4. Calculate the treatment difference over the moving window

This will be calculated in Column G. Assuming that the size of the window is 5 data points, enter values of 0 in the first 4 rows of the column. In cell G6, calculate the average treatment difference from the difference data in Column F (step 2):

= Average(F2:F6)

Copy this formula down to the last row the dataset.

5. Calculate the significance (P value) of the difference calculated at step 4 using the two-tail paired student t-Test.

This will be calculated in Column H. In the cell of H6, enter the following formula:

=TTEST(D2:D6,E2:E6,2,1)

Copy this formula down to the last row the dataset.

The data generated by steps 4 and 5 enable the kinds of graphs displayed in Lawes and Bramley (2012) to be produced showing how the response to the treatment in the strip differs from the areas outside the strip, and the statistical significance of that difference, along the strip length.

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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