# **Metamorph Files Folder To Hyperstacks**

# ImageJ macro, Marcel Boeglin Jul 2018 – Apr 2020 **User guide**

### **Introduction - Features:**

MetamorphFilesFolderToHyperstacks\_ ImageJ macro Author: Marcel Boeglin, July 2018 - January 2020

e-mail: boeglin@igbmc.fr

- <sup>22</sup> Opens Metamorph multi-position time-series z-stacks of up to 7 channels from input folder and saves them as hyperstacks to output folder.
- <sup>II</sup> Only **TIFF** and **STK** files are processed.
- <sup>2</sup> Image files for which no series name can be found are skipped.
- <sup>II</sup> Single file series (including output files of this macro the user manually copied to input folder) are skipped because output files would be identical to input files,
- <sup>¤</sup> Series having same channel sequence are grouped in channel groups. All series from a given channel group are processed using same parameters.
- Example 2 colors are determined automatically from filenames by use of red, green, blue, cyan, magenta, yellow and gray determinants but can be changed by the user. Only these 7 colors are possible.

For instance, the red determinants set is: {"543", "555", "561", "594", "CY3", "Y3", "DsRed", "mCherry", "N21", "RFP", "TX2"}.

Channels containing a red, green, blue or gray determinant in their names are assumed to be respectively red, green, blue or gray.

In future versions, it's planned to get channel colors from metadata in case Metamorph has been configured to not add the illumination settings to the filenames (which in that case contain just  $_{w1}$ ,  $_{w2}$ ,  $_{y}$ ,  $_{y}$ ).

If none of the color determinants works with your illumination settings, just add the missing determinant(s) to the appropriate set in the macro (advanced users).

para Dual camera channels handling:

It's Assumed channels are saved in separate images. As in single camera acquisitions, colors are derived from filenames.

Dual camera filenames are assumed to be of the type:

seriesName\_w1Lambda1 Lambda2... seriesName\_w2Lambda1 Lambda2...

One could expect that w1 corresponds to lambda1 and w2 to lambda2, but w1 may correspond to lambda2 and w2 to lambda1, depending on the configuration of Metamorph.

The dual channel separator (a space character in the example) may be different, for instance a "-", depending on how dual camera illumination settings have been named in Metamorph.

Dual channel order and separator are managed by the macro but are assumed to be the same for all series in a given folder.

If the dual channel colors attribution fails, arbitrary (probably unwanted) colors are assigned to the channels. In such a case, the user can choose the color of each channel for each channel group.

- <sup>22</sup> Output images are X, Y, Z and T calibrated if calibration data are available. If not, a unique xy-z-t calibration, applied to **all series**, can be set by the user to avoid manual calibration of output files.
- □ Allows control of z-range, position-range (s1, s2, s3, ...) and time-range of processed files.
- <sup>a</sup> Does optional resizing or cropping of input files.
- <sup>22</sup> Does optional maximum, average, minimum, median or sum z-projection of input files and color balance of output files.
- ¤ In case of heterogeneous z-dimensions between channels and 'Do z-projection' is unchecked, single-section channels are transformed into z-stacks having same depth as channels acquired as z-stacks by duplicating the single-section.
- Series handling dialog is not displayed if larger than screen (happens if the folder contains more series than can be added as checkboxes in a dialog). In this case all series that have passed the including and excluding file-filters are processed.
- From version 43u11, MetamorphFilesFolderToHyperstacks\_ can be run in easy or full mode:
   Easy modes use default parameters and skip dialog boxes.
   Full mode provides access to all parameters through up to eight dialog boxes.
- <sup>22</sup> From version 44u16, the "Data Reduction dialog displays a combobox allowing to extract regions in speed optimizing or memory economising mode."
- <sup>II</sup> From version 44u38, the main dialog displays a checkbox to open the **'Memory'** window during run.

#### ¤ Dependencies:

- "tiff\_tags" plugin from Joachim Wesner <a href="https://imagej.nih.gov/ij/plugins/tiff-tags.html">https://imagej.nih.gov/ij/plugins/tiff-tags.html</a> which is able to access full metadata needed to calibrate the z-interval.
- "MemoryMonitor\_Launcher": plugin that allows to open the 'Memory' window within a ImageJ macro without blocking it as when you use the run("Monitor Memory...") command.
- The **Fiji.app\macros**\ or \**ImageJ\macros**\ **StartupMacros.ijm** file must contain following macros set to allow MetamorphFilesFolderToHyperstacks\_ to be run in one of easy modes or in full mode.

### Macros set to be copied to StartupMacros.ijm (preferably to the end of the file):

```
/**
* This macro set runs MetamorphFilesFolderToHyperstacks .ijm
* in three different modes:
* - "EasyMode_Zproj [F1]"
* - "EasyMode noZproj [F2]"
* - "Full_Mode [F3]"
* Easy modes ask the user only for input and output folder.
* Full mode displays all dialogs allowing the user to set all the
  parameters for input folder files processing.
* Author: Marcel Boeglin - January 2020
macro "Metamorh Files FolderTo Hyperstacks:" {}
macro "EasyMode_Zproj [F1]" {
      mode = "EasyMode_Zproj [F1]";
      showMessageWithCancel("Run macro \""+mode+"\" ?");
      //print("\\Clear");
      print("\nRunning EasyMode Zproj:");
      runMacro("MetamorphFilesFolderToHyperstacks_.ijm", "EasyMode_Zproj");
}
macro "EasyMode noZproj [F2]" {
      mode = "EasyMode noZproj [F2]";
      showMessageWithCancel("Run macro \""+mode+"\" ?");
      print("\nRunning EasyMode_noZproj:");
      runMacro("MetamorphFilesFolderToHyperstacks_.ijm", "EasyMode_noZproj");
}
macro "Full_Mode [F3]" {
      mode = "Full Mode [F3]";
      showMessageWithCancel("Run macro \""+mode+"\" ?");
      print("\nRunning Full_Mode:");
      runMacro("MetamorphFilesFolderToHyperstacks_.ijm", "Full_Mode");//default = noZProj
      //runMacro("MetamorphFilesFolderToHyperstacks_.ijm", "fullMode doZProj");
}
```

### Performance: typically 120-180 GB per hour including output files saving.

To extract large numbers of regions in one or more positions from one or more series, choose the "Optimize Speed" option in the 'Data Reduction' dialog.

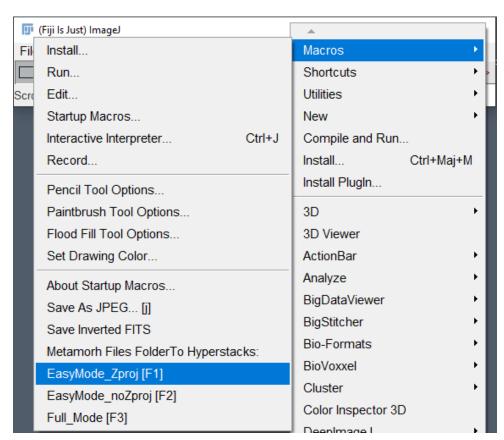
Else ("Economise Memory"), performance may be dramatically reduced because corresponding algorithm implies input files to be opened as many times as there are regions to be extracted. Note: processing files from or to an external drive (ethernet, usb or other) may also drastically increase process time.

### **Installation and short instructions:**

- Copy tiff tags.jar to the plugins folder of your ImageJ or Fiji installation.
- Copy "MetamorphFilesFolderToHyperstacks\_.ijm" (current version 43u16) to the "macros" folder.
- Copy the above macro sets to the StartupMacros.ijm file, restart ImageJ or Fiji.

In the "Plugins > Macros" menue, you will have 4 new commands (requires 43u11 or later):

- Metamorph Files to Hyperstacks does nothing
- EasyMode\_Zproj [F1] processes all series from input folder, z-projected output files;
- EasyMode\_noZproj [F2] processes all series from input folder, non z-projected output files;
- Full\_Mode [F3] gives access to all features (up to 10 dialogs).



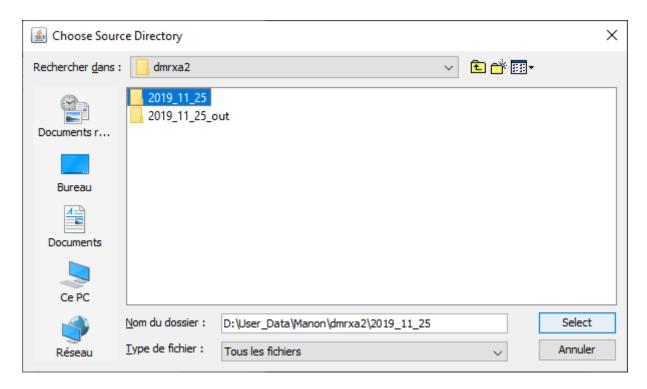
Easy modes just need you choose the input and output folders and the macro will run with default parameters.

Full mode takes less than a minute to answer the dialogs but may save tens of hours spent to fight with your computer to open, merge and save hundreds of images. Moreover, the region extraction procedure allows processing of images that you never would be able to process directly with ImageJ (when the size exceeds the ImageJ's capacity (about 2 G pixels for the XYCZT product of image's all dimensions)).

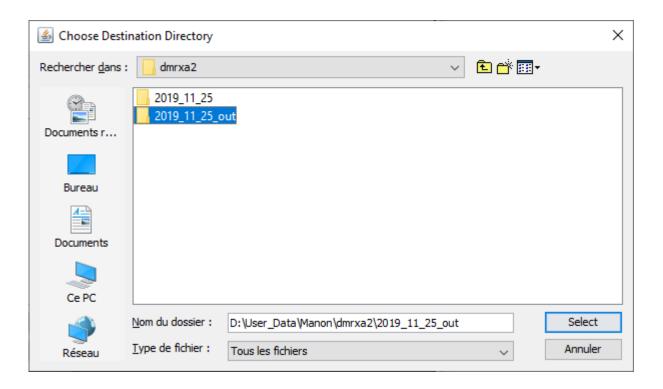
Full mode is described in details in next pages.

# **Detailed Procedure and description of dialog boxes:**

# 1. Choose input folder:

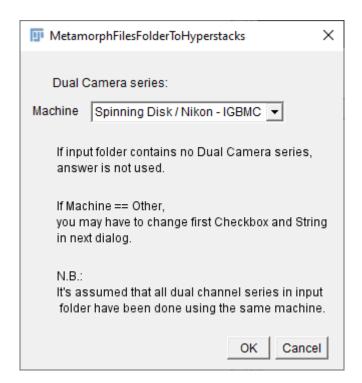


# 2. Choose output folder: shouldn't (and can't) be identical to input folder



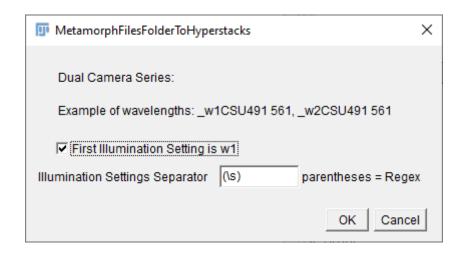
# 3. Dual Camera series, dialog 1:

- If you didn't do dual-camera acquisitions, skip this dialog by clicking the OK button or typing "Enter" without changing anything;
- Else select the machine used for dual-camera acquisitions.



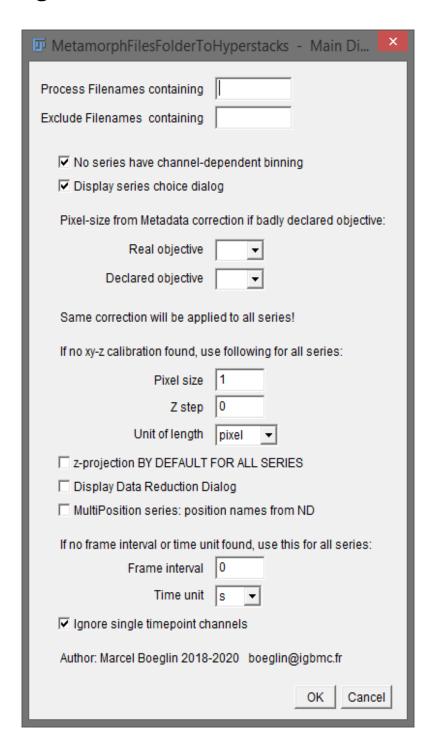
## 3. Dual Camera series, dialog 2:

- Defaults in this dialog are set automatically if you have selected a spinning disk from IGBMC's Imaging Center in dialog1,



so as you can validate it by pressing the OK button or the "Enter" key.

### 4. Main Dialog:



- > **File filters:** limit processing to filenames containing the including string but not the excluding string. Allows to process only a part of input folder; reduces also delay to display next dialogs and the number of components they contain.
- > **Missing files infos:** if some images have been lost, the macro doesn't stop but replaces any missing image with a black one where is written the path of the expected file. The path can be written in the output image's overlay (recommanded) or grabbed in its pixels.
- > **Channel-dependent binning:** the macro is designed to manage such situations but this feature has not been tested.

- > **Pixel-size from Metadata correction:** to be used only if Metamorph has been configured for pixel-size calibration. If the used objective is the declared one, let as is.
- > **If no xy\_z calibration data found** in metadata, a unique calibration can be used **for all series** to avoid having to do it manually afterwards by opening and saving again the output images.
- > **z-Projection BY DEFAULT FOR ALL SERIES:** if checked, channel groups handling dialog defaults will be do z-projection for all channel groups, but you can choose to do z-projections or not for each channel-group individually.
- > **Display Data Reduction Dialog:** select if you want to resize or crop images at import or if you want to limit processing to given positions, z or time ranges; you will get an extra dialog to indicate what you want to do.
- > **MultiPosition series: position names from ND:** select if you want to replace "s1", "s2", "s3" etc. suffixes from input names with the position names you entered in the Metamorph "Stage" tab, for instance "WT\_1mM", "WT\_10mM", "WT\_100mM" etc. If the Metamorph ND file for a series is not found or is unusable, the macro will not stop working. Output files position suffixes will just remain "s1", "s2", "s3" etc..
- > **If no frame interval or time unit is found** in metadata: you can set a unique time-calibration that will be used **for all output series**, avoiding to do it manually afterwards.
- > **Ignore single time point channels:** avoids processing images that are not parts of a time-lapse. Avoids issues if you have used the possibility of taking only first and last time-points for some channels.

### 5. Data Reduction:

- > **Resize at import:** use for instance 0.5 or less if output images are too big for installed memory or for ImageJ's capabilities.
- > Crop at import: does extraction of rectangular regions from input images in multi-channel and / or time series. For more details, refer to "Extracting regions from input images". If you have enough memory, choose the Optimize Speed processing mode. If you have 10 regions to extract, it will work ~ 10 times faster than the Economize Memory mode.
- > **Z-series:** if firstSlice > lastSlice, output stack will be reversed.

  ¤ Process range around median slice:
   useful if series have multiple stack sizes and you want to limit output images stack sizes.
- > **Multi-positions series:** to process for instance only positions s9, s10 and s11, set "firstPosition" to 9 and "lastPosition" to 11.

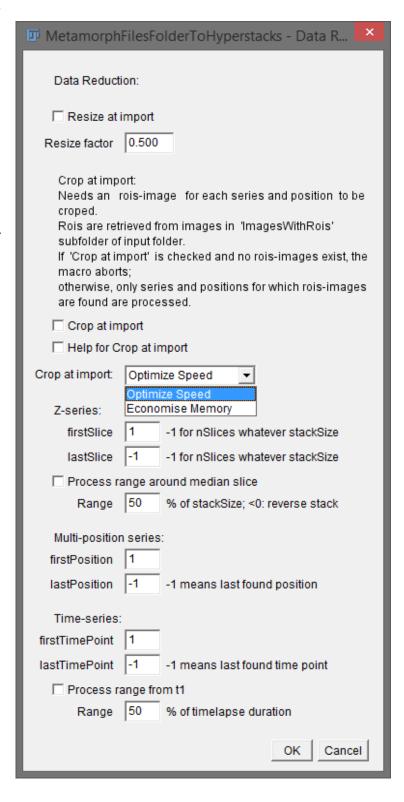
#### > Time-series:

to process for instance from t12 to end of timelapse, enter:

"firstTimePoint": 12 "lastTimePoint": -1

¤ Process range from t1:

if range is for instance 25 % and input folder contains a timelapse with 400 time points, only the 100 first time points will be processed. If the same folder contains another time-lapse of 100 time points, only those from 1 to 25 will be processed.



### 6. Process series:

This dialog lists all series found in input folder after file filters have been applied.

To skip a series, just uncheck it.

The list may be very long if not reduced by use of file filters in main dialog.

Remember that file filters allow to reduce processed files to a few series.

For instance, if the including filter (the upper one) in the Main Dialog is equal to a series name, only this series will be processed and the "Process series" dialog will even not be displayed.

If you chose **p Crop at import** and don't want to process all series in the **"ImagesWithRois"** folder, just uncheck the series you don't want to process.

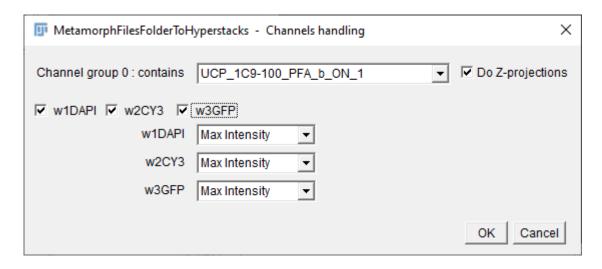
The chapter "Extract regions from input images" shows another way to limit the processing to given series.

#### Process series:

- **V** UCP 1C9-100 PFA b ON 1
- ✓ UCP 1C9-100 PFA b ON 2
- ✓ UCP 1C9-100 PFA b ON CCCP 1
- ✓ UCP 1C9-100 PFA b ON CCCP 2
- ✓ UCP 1C9-100 PFA b ON CCCP 3
- ✓ UCP 1C9-100 PFA b RT 1
- ✓ UCP 1C9-100 PFA b RT 2
- ▼ UCP 1C9-100 PFA b RT CCCP 1
- ▼ UCP 1C9-100 PFA b RT CCCP 2
- ▼ UCP 1C9-100 PFA b RT CCCP 3
- ▼ UCP 1C9-100 PFA nb ON 1
- ▼ UCP 1C9-100 PFA nb ON 2
- ✓ UCP 1C9-100 PFA nb ON CCCP 1
- ✓ UCP 1C9-100 PFA nb ON CCCP 2
- UCP 1C9-100 PFA nb ON CCCP 3
- ▼ UCP 1C9-100 PFA nb RT 1
- ▼ UCP 1C9-100 PFA nb RT 2
- ▼ UCP 1C9-100 PFA nb RT CCCP 1
- ▼ UCP 1C9-100 PFA nb RT CCCP 2
- ✓ UCP 1C9-100 PFA nb RT CCCP 3
- ✓ UCP 1E3-100 glyoxal b ON 1
- ✓ UCP 1E3-100 glyoxal b ON CCCP 1
- ✓ UCP 1E3-100 glyoxal b ON CCCP 2
- ✓ UCP 1E3-100 glyoxal b RT 1
- ✓ UCP 1E3-100 glyoxal b RT CCCP 1
- ✓ UCP 1E3-100 glyoxal b RT CCCP 2
- ✓ UCP 1E3-100 glyoxal nb ON 1
- UCP 1E3-100 glyoxal nb ON CCCP 1
- ✓ UCP 1E3-100 glyoxal nb ON CCCP 2
- UCP 1E3-100 glyoxal nb RT 1
- ✓ UCP 1E3-100 glyoxal nb RT CCCP 1
- ✓ UCP 1E3-100 glyoxal nb RT CCCP 2
- ✓ UCP 1F9-100 glyoxal b ON 1
- ✓ UCP 1F9-100 glyoxal b ON CCCP 1
- ✓ UCP 1F9-100 glyoxal b RT 1
- **V** UCP 1F9-100 glyoxal b RT CCCP 1
- ✓ UCP 1F9-100 glyoxal b RT CCCP 2
- UCP 1F9-100 glyoxal nb ON 1
- ✓ UCP 1F9-100 glyoxal nb ON CCCP 1
- UCP 1F9-100 glyoxal nb RT 1
- ✓ UCP 1F9-100 glyoxal nb RT CCCP 1

OK Cancel

# 7. Channels handling:



The macro groups the series in sets having identical channel sequences where the order of channels is taken in account to avoid mixing files from different series.

For instance, sequence (w1Dapi, w2CY3) is different from (w1CY3, W2Dapi).

If the input folder contains multiple channel sequences, the "Channels handling" dialog will contain a components group for each of them, allowing to control each channel sequence individually.

**Channel group n: contains:** This combo box is for information only; it lists the series names concerned by nth Channel group.

All choices done here will be applied to all series from current channel group.

**a Do Z-projections:** this checkbox allows you to do or not z-projection for this channel group.

Following is just an example, the situation may be different for you:

**¤ w1Dapi ¤ w2CY3 ¤ w3GFP:** unchecked channels will not be added to output images.

**w1DAPI** Max Intensity:

w2CY3 Max Intensity:

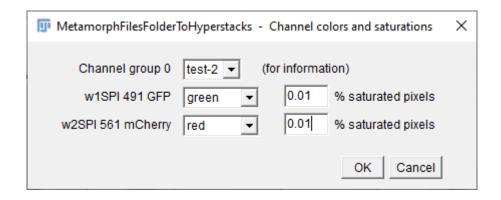
w3GFP Max Intensity:

These combo boxes are for choice of projection type of first, second, third, ... channel if **Do Z-projections is checked.** 

The available projection types are: Average Intensity, Max Intensity, Min Intensity, Median and Sum Slices.

If any channel is projected using Sum Slices, output images type is 32-bit for all series in the channel group.

### 8. Channels colors and saturations:



The macro determines automatically the default colors, but you can change output colors if you prefer different ones.

**Caution:** the macro uses ImageJ's color merge command. It fails if two channels have the same color. In this case, the macro tries to replace one of the duplicate colors by a different one, but you have no control on the procedure. If it fails due to an unknown bug, the execution will stop.

**Channel group n:** this combo box is for information only; it lists the series names concerned by nth Channel group.

The choices for a given channel group stand for all series it contains.

#### Following is just an example:

**w1SPI 491 GFP green 0.01** % saturated pixels:

the macro proposes "green" because the channel name contains "GFP".

**w2SPI 561 mCherry red 0.01 % s**aturated pixels:

the macro proposes "red" because the channel name contains "mCherry".

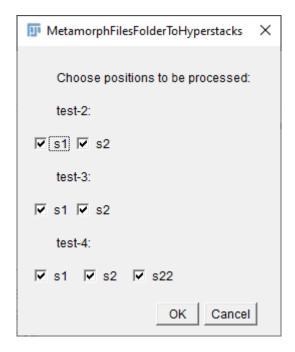
**Saturated pixels %:** the macro uses these values to set the maximum displayed values of each channel in output images, calculated in the median slice for z-stacks and median time point for time-lapse images.

**Note**: the macro never uses the "Apply" function from the Image > Adjust > Color Balance... or Brightness/Contrast... window.

This means that even if the output images appear saturated, their pixel values are **not** changed and are suitable for intensity measurements.

A simple click on "Reset" in the "B&C" or "Color" window will desaturate the active channel.

# 9. Choose positions to be processed:



#### For each selected series:

the dialog displays a checkbox for each detected position in given series. If you have filtered the positions, only those that have passed the filter will appear here.

To skip a position, just uncheck it.

This applies also if you chose **property Crop at import**. If you don't want to process some positions of some of the series contained in the **"ImagesWithRois"** folder, just uncheck them in this dialog.

Note: Dialogs with component groups as several checkboxes on a same row cannot be validated by pressing "Enter", you must click on "OK".

# 10. Extracting regions from input images:

- Create folder "ImagesWithRois" in input folder

# - For each series or position from which you want to extract regions of interest:

- Open an input image (any channel or time-point)
- If it's a stack, do a z-projection (Average, Max, Min etc.)
- Draw a rectangle for each region to be processed
- Add the rectangle to the overlay (press b or Ctrl b)
- Save the image with rois to folder "ImagesWithRois". Don't rename it, if its name doesn't contain the series name and the position name, it will not be processed.
- Close the projection-image.

### - Launch MetamorphFilesFolderToHyperstacks:

- In the "Main Dialog", check "Display Data Reduction Dialog"
- In the "Data Reduction" dialog check "a Crop at import"

This will result in limiting process to:

- series and positions for which an image is found in the "ImagesWithRois" folder.
- regions drawn in these images; all detected regions are processed.

#### **a** Output files:

For each region drawn in images with rois:

- Cropped image of merged channels and time-points of corresponding series.
- The output image-names contain the coordinates of the upper-left edge of the extracted region.

#### **¤** Tips:

#### In case of a timelapse and the sample moved during acquisition:

- Open an input image of the first timepoint to be processed, do z-projection if it's a stack.
- Repeat this for the last time point to be processed.
- Synchronize the two projections
- Draw the rois in the first time point projection: during the drawing process, the roi will be visible also in the last timepoint projection, so that you can make it large enough to contain the sample at the begin and the end of the time-lapse.
- Save the first timepoint projection to the "ImagesWithRois" folder.
- If you want to keep the last timepoint projection, you can save it also in the "ImagesWithRois" folder, it will not be taken in account if you check the "Crop at input checkbox.

#### To find easier which output image corresponds to a given region:

- Open the image with rois
- launch Image > Overlay > List Elements
- In the Overlay Elements table, double-click on a row activates corresponding roi; The X and Y coordinates of the roi can be compared with those of the cropped output image name which ends for instance with "\_x92y90.tif", where x and y are the coordinates of the upper-left corner of the extracted region.

#### Management of images with rois:

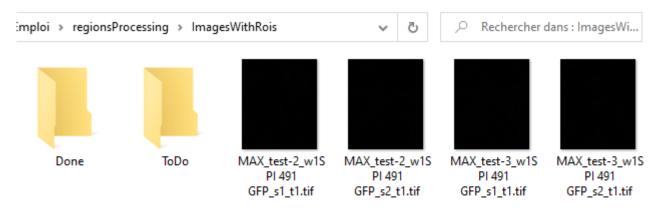
If you don't want to process all series or positions for which you have created images with rois in "ImagesWithRois" folder in a single run:

- Create subfolders "ToDo" and "Done" in "ImagesWithRois" folder,
- Push all the images you don't want to process in next run to "ToDo" folder (keep only those to be processed in "ImagesWithRois" folder). If a Crop at import is checked, only files in

- "Images WithRois" folder will be processed even if you have selected all series and positions;
- If the run terminated properly, push the images from "ImagesWithRois" to "Done".
- To process another batch, move corresponding images from "ToDo" to "ImagesWithRois" and run the macro again.

Notes: "ToDo" and "Done" can also be sub-folder of the input folder, not necessarily of "ImagesWithRois".

**"ToDo"** and **"Done"** folders may be named differently, but not the **"ImagesWithRois"** folder. **"ImagesWithRois" must** be a direct sub-folder of input folder.



Screen cast showing an example of how to organise management of images with rois

In case "ImagesWithRois" folder contains several images with rois corresponding to same series or position:

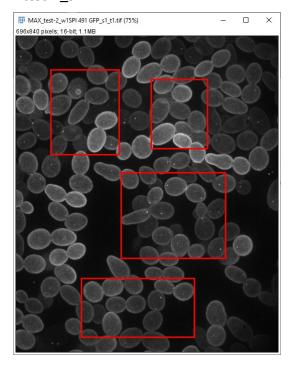
Only one of the images will be used to extract rois. The used image is the first one in the list built by MetamorphFilesFolderToHyperstacks. The list-order possibly depends on the operating system and may be different on Windows, Linux and MacOS.

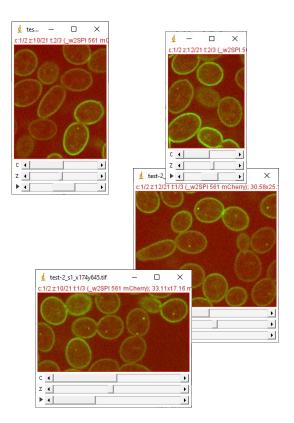
For this reason, it's recommended to avoid having several images with rois corresponding to the same series or position in the "ImagesWithRois" folder.

Images containing potential rois for the given series or position, can be stored in the "ToDo" folder.

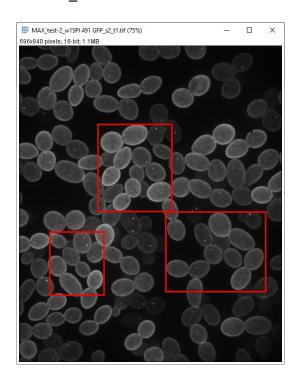
### Examples of images with rois for regions extraction and output images:

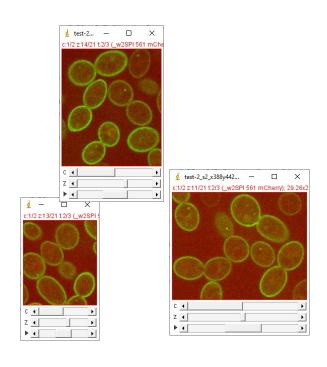
test-2\_s1





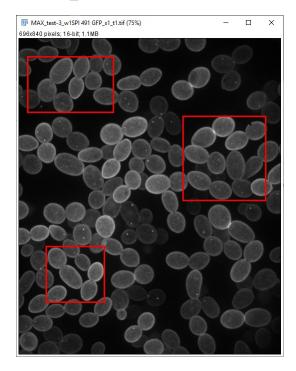
test-2\_s2

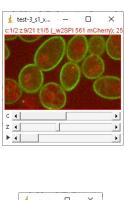


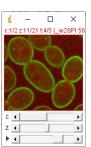


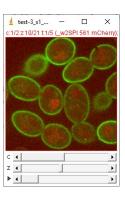
### Examples of images with rois for regions extraction and output images (continued):

test-3\_s1

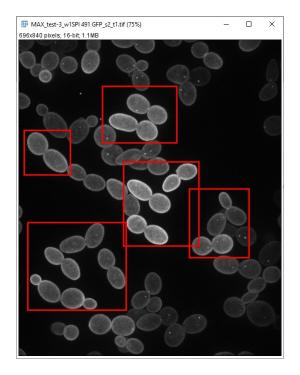


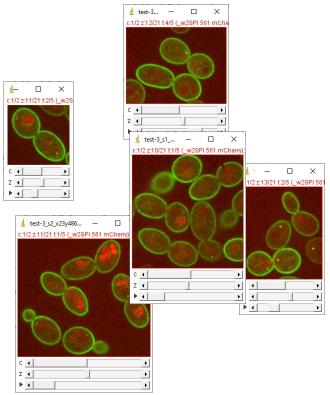




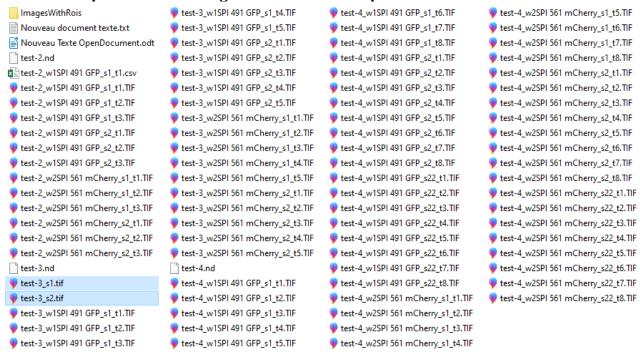


 $test-3_s2$ 





#### Content of input folder in the regions-extraction example:



In the example, the macro may works properly despite input folder contains output images. But it's recommended to avoid such situations which may cause unexpected issues.

#### Content of Input folder / ImagesWithRois:

MAX\_test-2\_w1SPI 491 GFP\_s1\_t1.tif
 MAX\_test-2\_w1SPI 491 GFP\_s2\_t1.tif
 MAX\_test-3\_w1SPI 491 GFP\_s1\_t1.tif
 MAX\_test-3\_w1SPI 491 GFP\_s2\_t1.tif

#### **Content of output folder:**



#### **Notes:**

From version 44a18, MetamorphFilesFolderToHyperstacks\_ macro uses the .nd file to rename the positions "s1", "s2", "s3", ... according to the names you entered in Metamorph's Multiposition tab.

A more efficient implementation of the tasks done by the macro would need to rewrite it in Java which is much more efficient and versatile than the ImageJ macro language. Java is strongly object-oriented and allows to store intermediate images and other objects in multidimensional arrays or collections and to recall them when necessary, simplifying drastically the code.

Translating the macro in Java would take as long as needed to design the macro because it would require a complete refactoring.

#### New in version 44:

Possibility of renaming the output files positions according to their names in the multiposition window of Metamorph.

Regions extraction can be done in two different modes:

#### - Optimize Speed:

Regions extraction needs each input file to be opened only once. Output files corresponding to the different regions remain in memory until they have all been completed in Z, C, T dimensions and saved to output folder. If some series or positions have large and overlapping region with a total area larger than input images, the memory needs may be more than available. In this case, you can use following mode:

#### - Economize Memory:

This mode is mainly designed for low regions numbers extraction, optimally one, to reduce output image size in case of huge multichannel z-stack timelapses. Used together with Z and T ranges reduction, this may allow processing series which could not be processed in full XYZCT domain.