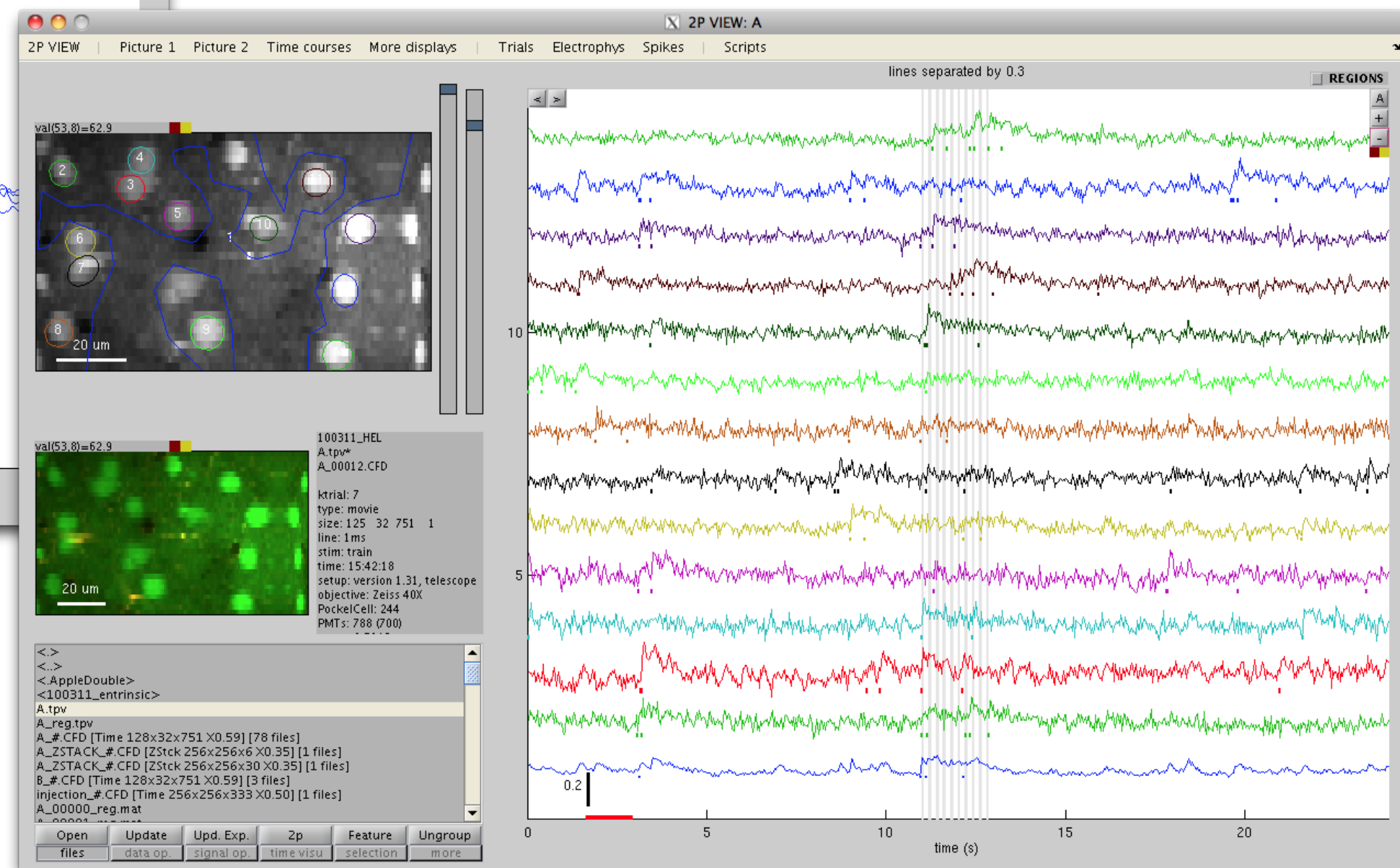
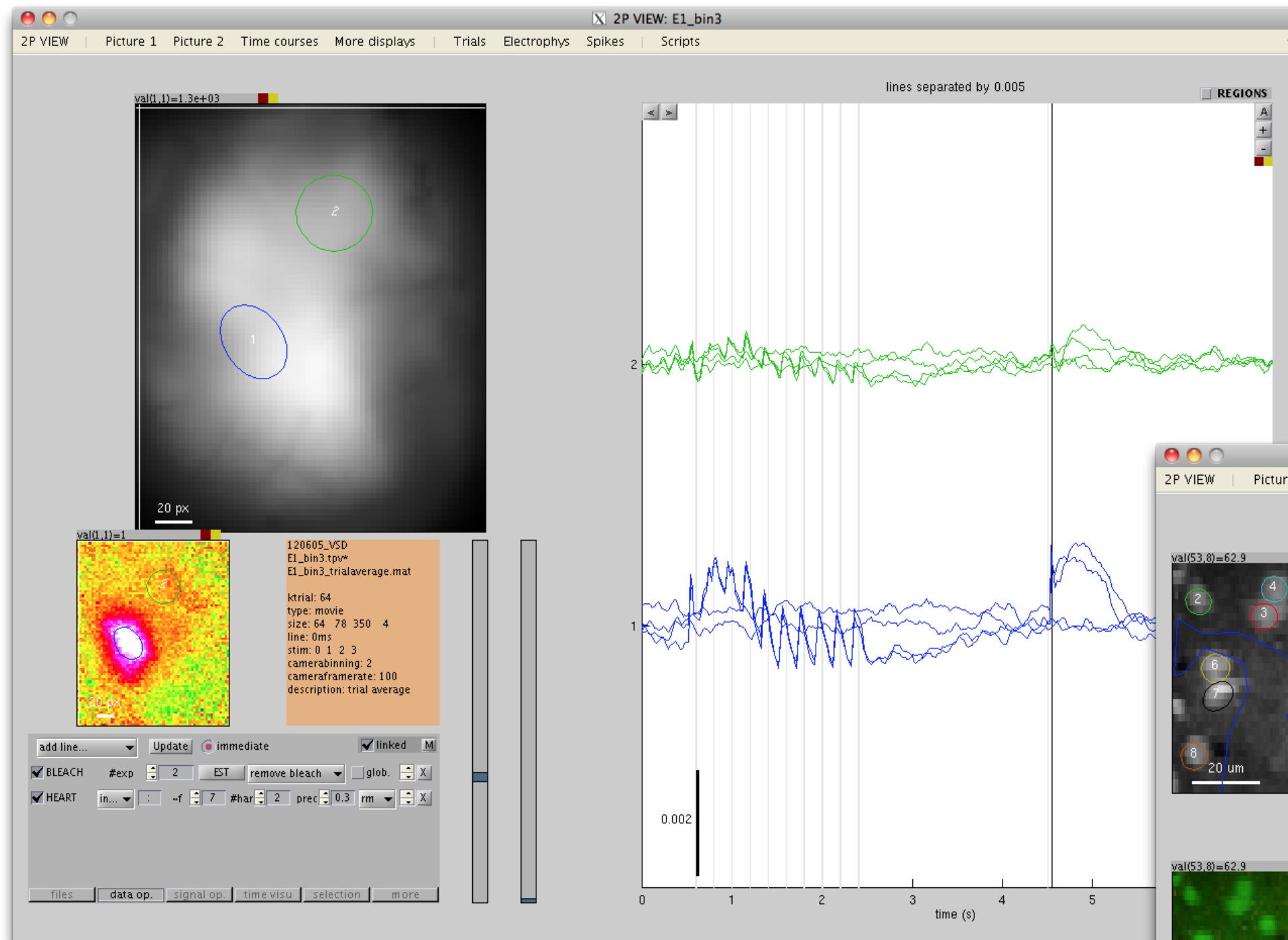


# 2PVIEW documentation



# Installation

## INSTALLATION

- Start Matlab
- Go in menu Files > Set path... and add the 4 directories 'brick', 'fn4Dtoolbox', 'twophoton' and 'oifun' (do not check the box 'add sub-directories')

## START THE PROGRAM

- type tpview in the Matlab command

# I. Quick Tour

- 1) Manipulation of the data
- 2) Display of the data
- 3) Other useful tools

# 1) Manipulation of the data

a. Open the data

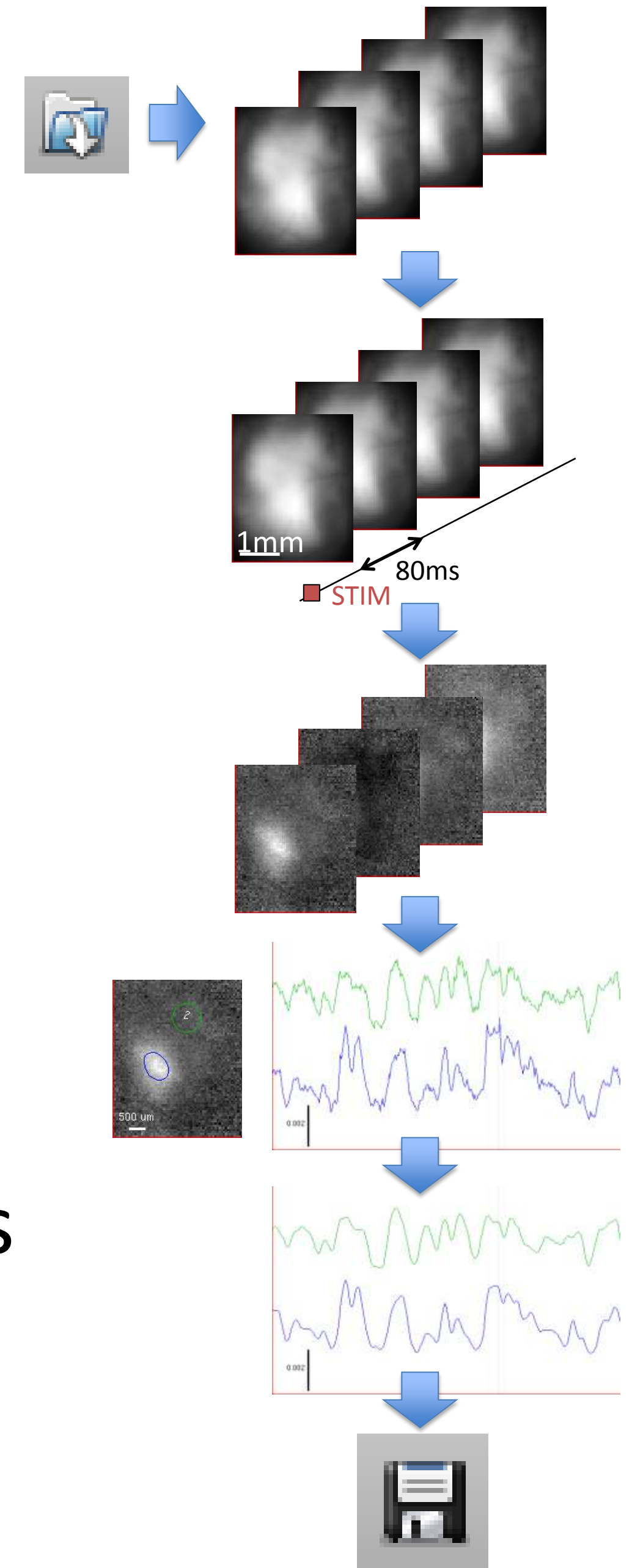
b. Header information

c. Operations on movie data

d. Extraction of time courses

e. Operations on time courses

f. Save



# 1)a. Open the data

The screenshot displays the 2P VIEW software interface. The top menu bar includes '2P VIEW', 'Picture 1', 'Picture 2', 'Time courses', 'More displays', 'Trials', 'Electrophys', 'Spikes', and 'Scripts'. The '2P VIEW' menu is open, showing options like 'Resize frames', 'Preset positions', 'Edit code', 'Reinit menus', 'Object in base workspace', 'Save PNG', 'Copy sub-part...', 'More', 'Preferences...', 'New window', 'Open...', 'Bin and open...', 'Save', 'Save as...', 'Comments', 'Auto-repair (light)', 'Auto-repair (heavy)', 'More repairs...', 'Edit code', 'access tpview object', 'Trial-specific selection', 'load data at opening', 'avoid loading data', and 'read data'. A red box highlights the 'Open...' option, with a callout bubble saying 'Open from the menu'. Below the menu, a small image shows a field of view with a scale bar of 500 um. To the right of the image, a text box contains the following information: '120605\_VSD', 'E1\_bin3.tpv\*', 'E1\_bin3\_trialaverage.mat', 'ktrial: 64', 'type: movie', 'size: 64 78 350 4', 'line: 0ms', 'stim: 0 1 2 3', 'camerabinning: 2', 'cameraframerate: 100', and 'description: trial average'. At the bottom left, a 'files' panel is open, showing a list of files: '<correlations>', '<data>', '<forpa>', '<vdaq>', 'E0(barrel).tpv', 'E1\_bin3.tpv', 'E1\_bin3\_corrected.tpv', 'E1\_bin4.tpv', 'E1\_bin4\_corrected.tpv', and '.DS\_Store'. A red box highlights the 'files' panel, with a callout bubble saying 'or use the « files panel »'. The main display area shows two time-series plots. The top plot is labeled 'lines separated by 0.0075' and shows a green line with a scale bar of 0.002. The bottom plot shows a blue line with a scale bar of 0.002. The x-axis for both plots is 'time (s)' ranging from 0 to 6.

2P VIEW: E1\_bin3

2P VIEW | Picture 1 | Picture 2 | Time courses | More displays | Trials | Electrophys | Spikes | Scripts

lines separated by 0.0075

REGIONS

Open from the menu

120605\_VSD  
E1\_bin3.tpv\*  
E1\_bin3\_trialaverage.mat

ktrial: 64  
type: movie  
size: 64 78 350 4  
line: 0ms  
stim: 0 1 2 3  
camerabinning: 2  
cameraframerate: 100  
description: trial average

500 um

or use the « files panel »

<correlations>  
<data>  
<forpa>  
<vdaq>  
E0(barrel).tpv  
E1\_bin3.tpv  
E1\_bin3\_corrected.tpv  
E1\_bin4.tpv  
E1\_bin4\_corrected.tpv  
.DS\_Store

Open | Update | Upd. Exp. | 2P | VSD | Unaroup

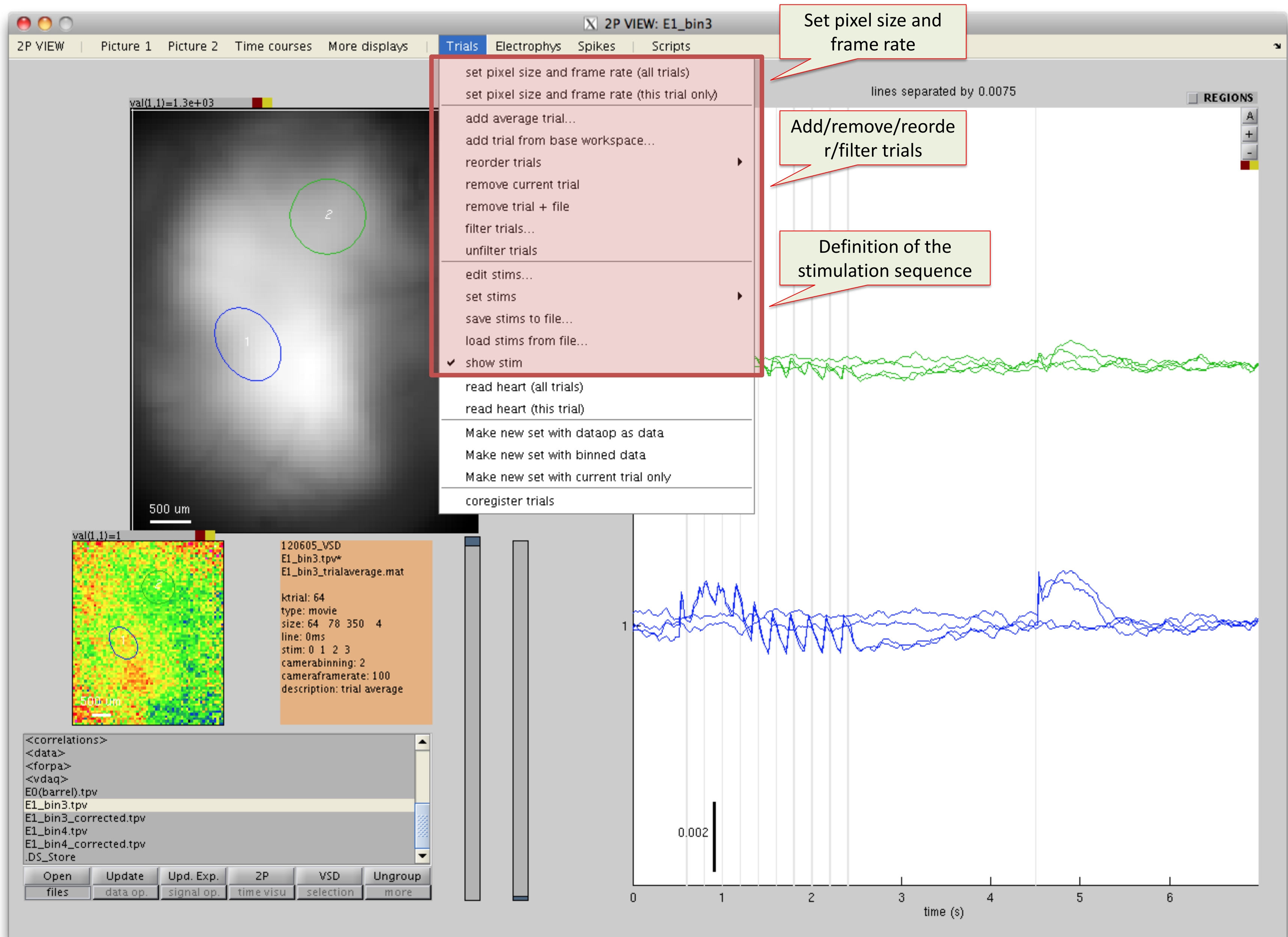
files | data op. | signal op. | time visu | selection | more

0.002

0 1 2 3 4 5 6

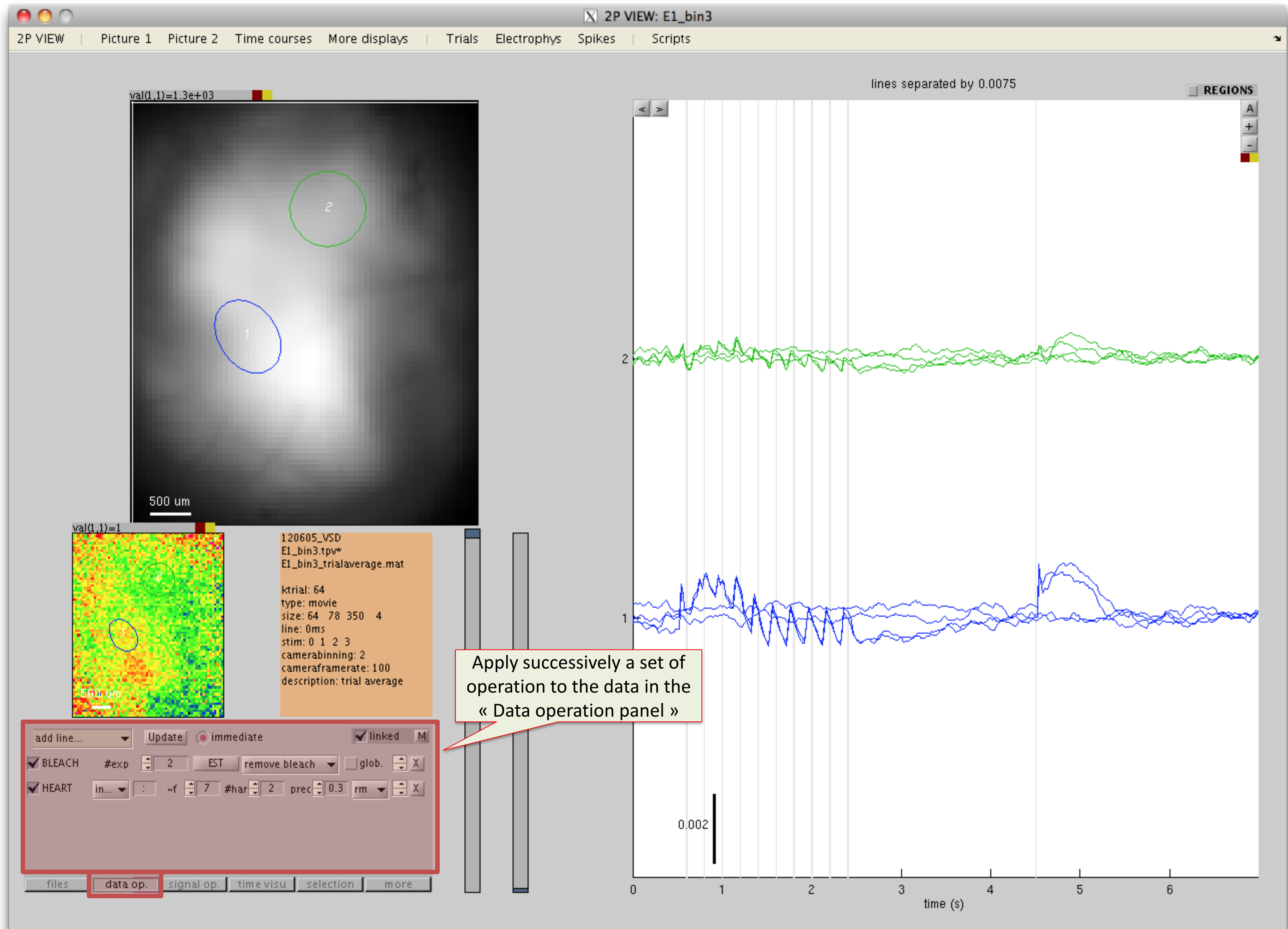
time (s)

# 1)b. Header information

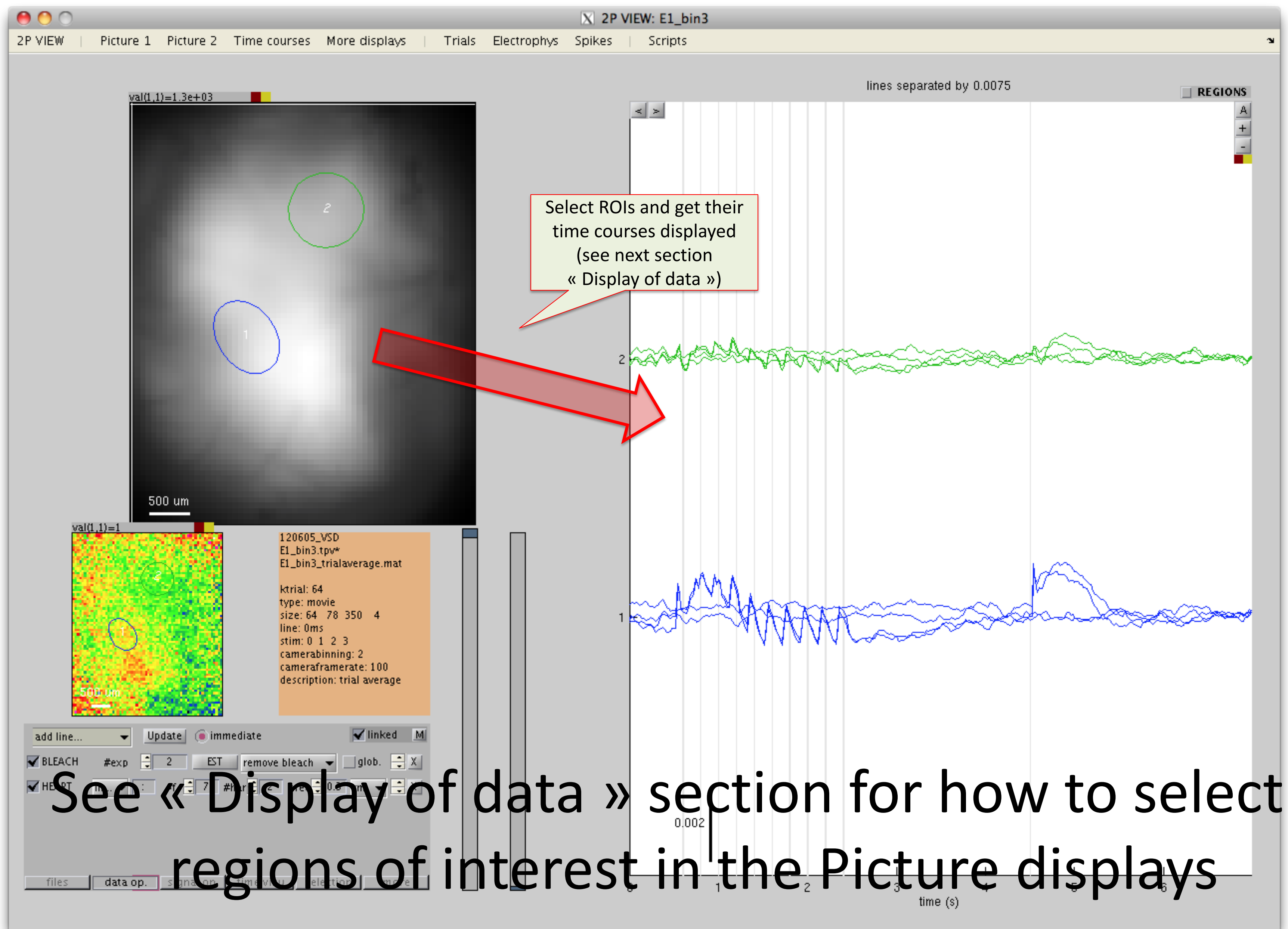




# 1)c. Operations on movie data

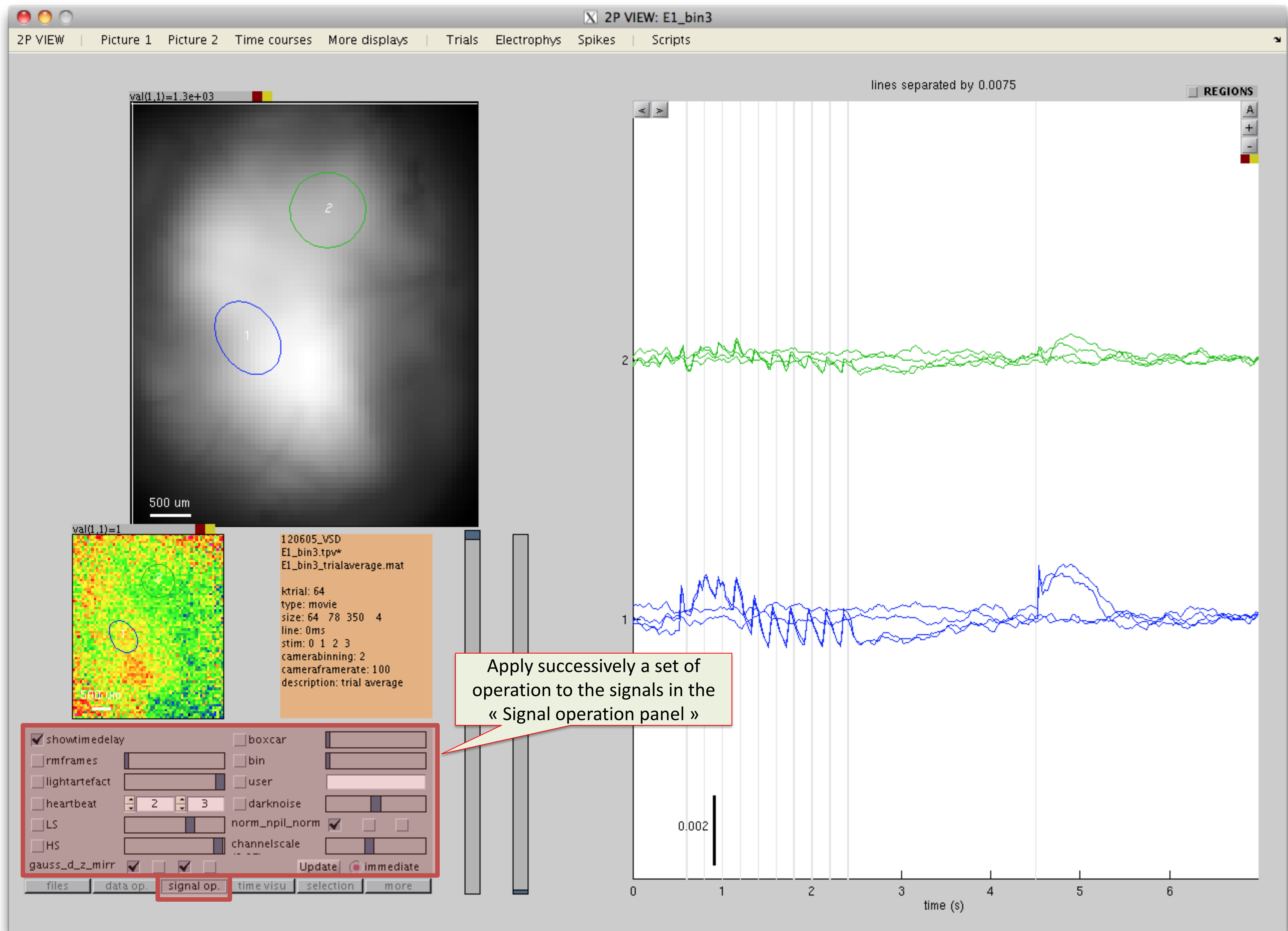


# 1)d. Extraction of time courses





# 1)e. Operations on time courses



# 1)f. Save

The screenshot displays the 2P VIEW software interface with the 'Trials' menu open. The menu options are as follows:

- Resize frames
- Preset positions
- Edit code
- Reinit menus
- Object in base workspace
- Save PNG (Ctrl+P)
- Copy sub-part...
- More
- Preferences...
- New window (Ctrl+N)
- Open... (Ctrl+O)
- Bin and open...
- Save (Ctrl+S)**
- Save as...
- Comments
- Auto-repair (light)
- Auto-repair (heavy)
- More repairs...
- Edit code
- access tpview object
- ✓ Trial-specific selection
  - load data at opening
  - avoid loading data
  - read data

A callout box points to the 'Save' option with the text: "Save / export / write in a comments file".

The 'Electrophys' menu is also open, showing options such as:

- set pixel size and frame rate (all trials)
- set pixel size and frame rate (this trial only)
- add average trial...
- add trial from base workspace...
- reorder trials
- remove current trial
- remove trial + file
- filter trials...
- unfilter trials
- edit stims...
- set stims
- save stims to file...
- load stims from file...
- ✓ show stim
- read heart (all trials)
- read heart (this trial)
- Make new set with dataop as data**
- Make new set with binned data**
- Make new set with current trial only**
- coregister trials

A callout box points to the 'Make new set with current trial only' option with the text: "Save a modified version of the data".

The main window displays a time course plot of fluorescence data (blue line) over 6 seconds. A scale bar indicates 0.002. The plot is titled 'lines separated by 0.0075'. A 'REGIONS' panel on the right shows a color-coded map of the field of view.

Metadata for the current trial is displayed in the bottom left:

```
120605_VSD
E1_bin3.tpv*
E1_bin3_trialaverage.mat

ktrial: 64
type: movie
size: 64 78 350 4
line: 0ms
stim: 0 1 2 3
camerabinning: 2
cameraframerate: 100
description: trial average
```

At the bottom, there are checkboxes for 'showtimedelay', 'rmframes', 'lightartefact', 'heartbeat', 'LS', 'HS', and 'gauss\_d\_z\_mirr'. There are also sliders for 'boxcar', 'bin', 'user', 'darknoise', 'norm\_npil\_norm', and 'channelscale'. The 'Update' button is highlighted.

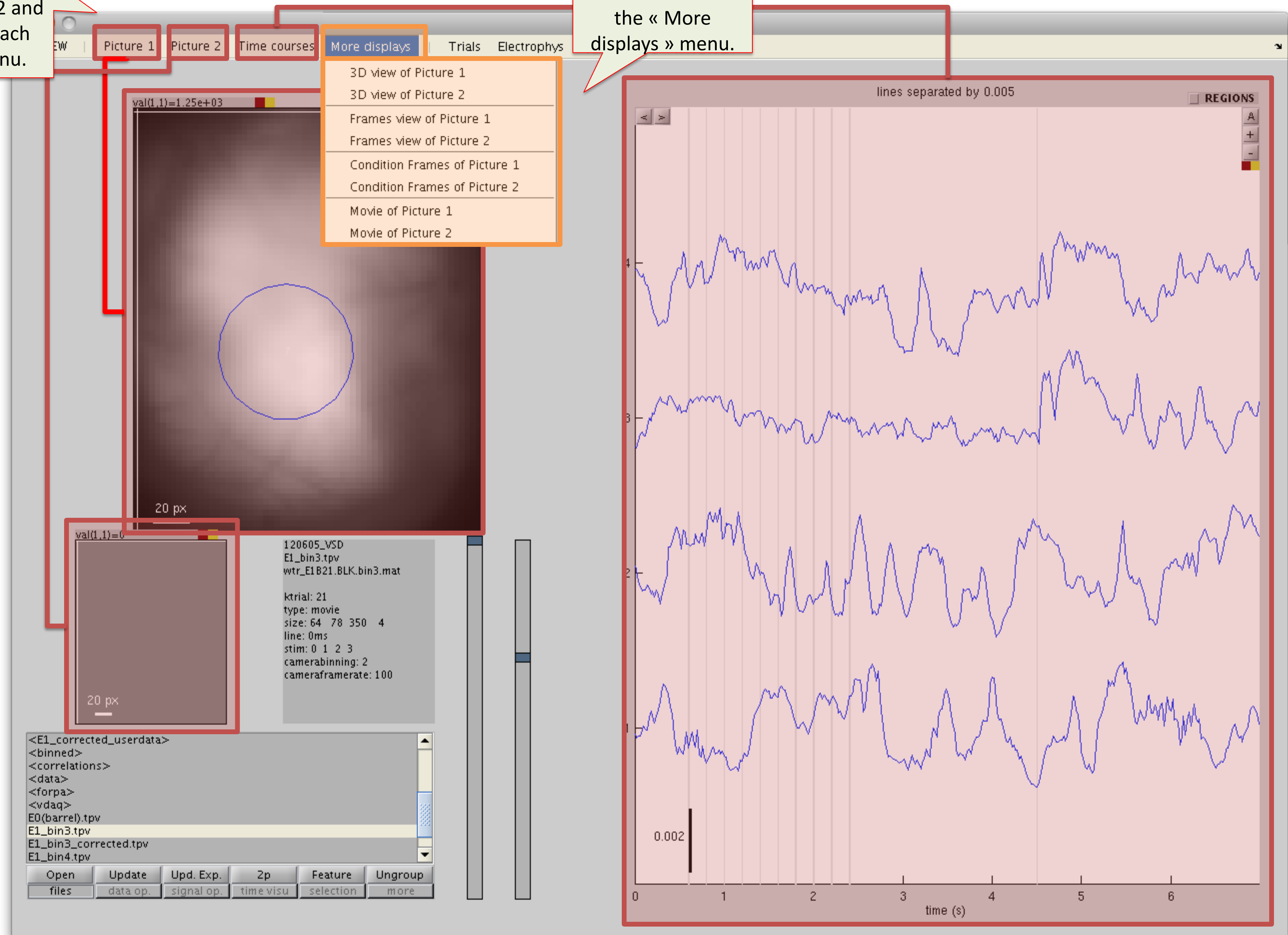
## 2) Display of the data

- a. Which displays are available?
- b. What to show in each display?
  - Picture1 and Picture2
  - Time Courses
- c. Display options
  - Picture1 and Picture2
  - Time Courses
- d. Mouse actions
  - Principles of region selection
  - Table of mouse actions

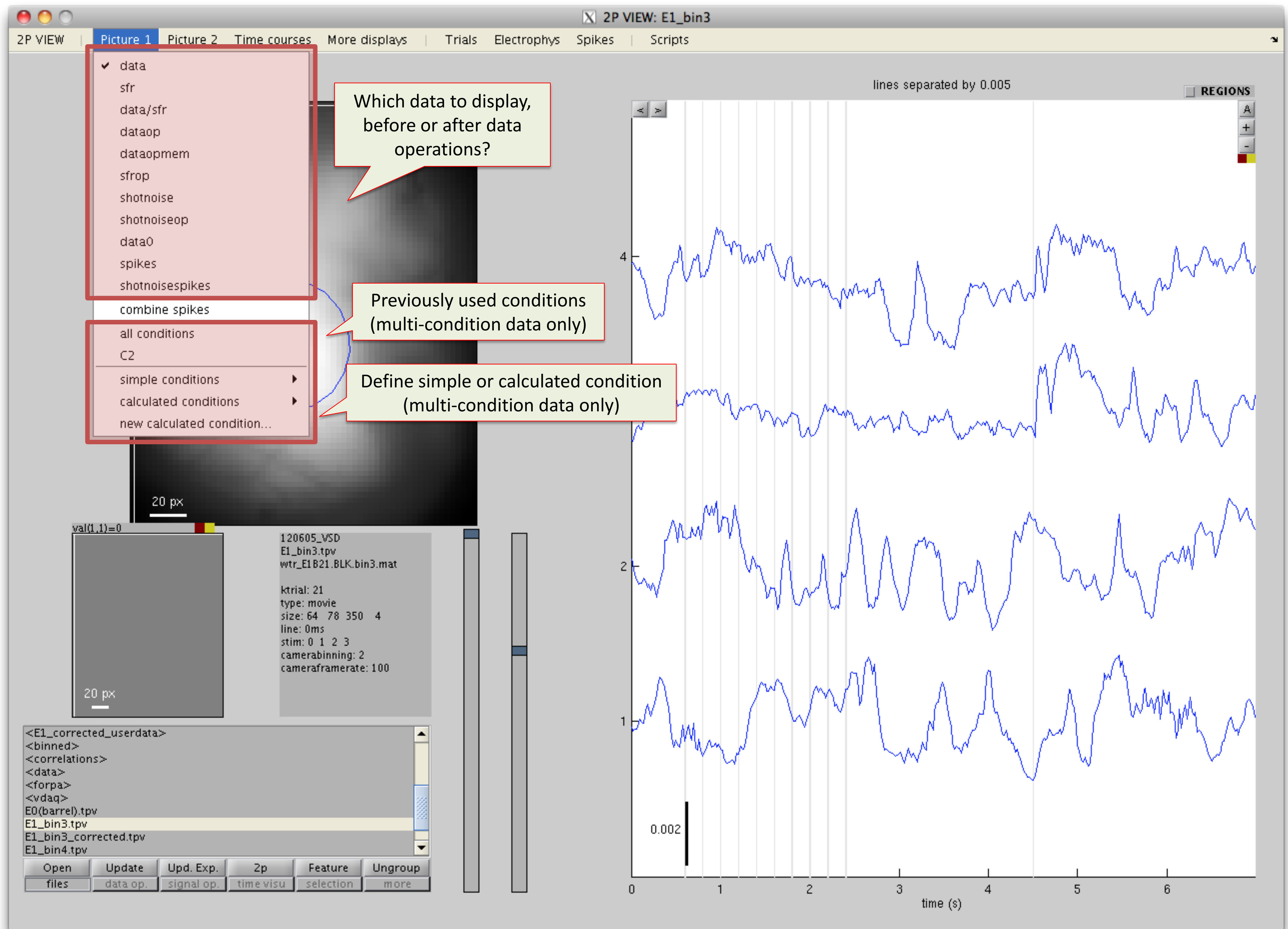
# 2)a. Which displays are available?

Three default displays:  
Picture1, Picture2 and  
Time courses. Each  
has its own menu.

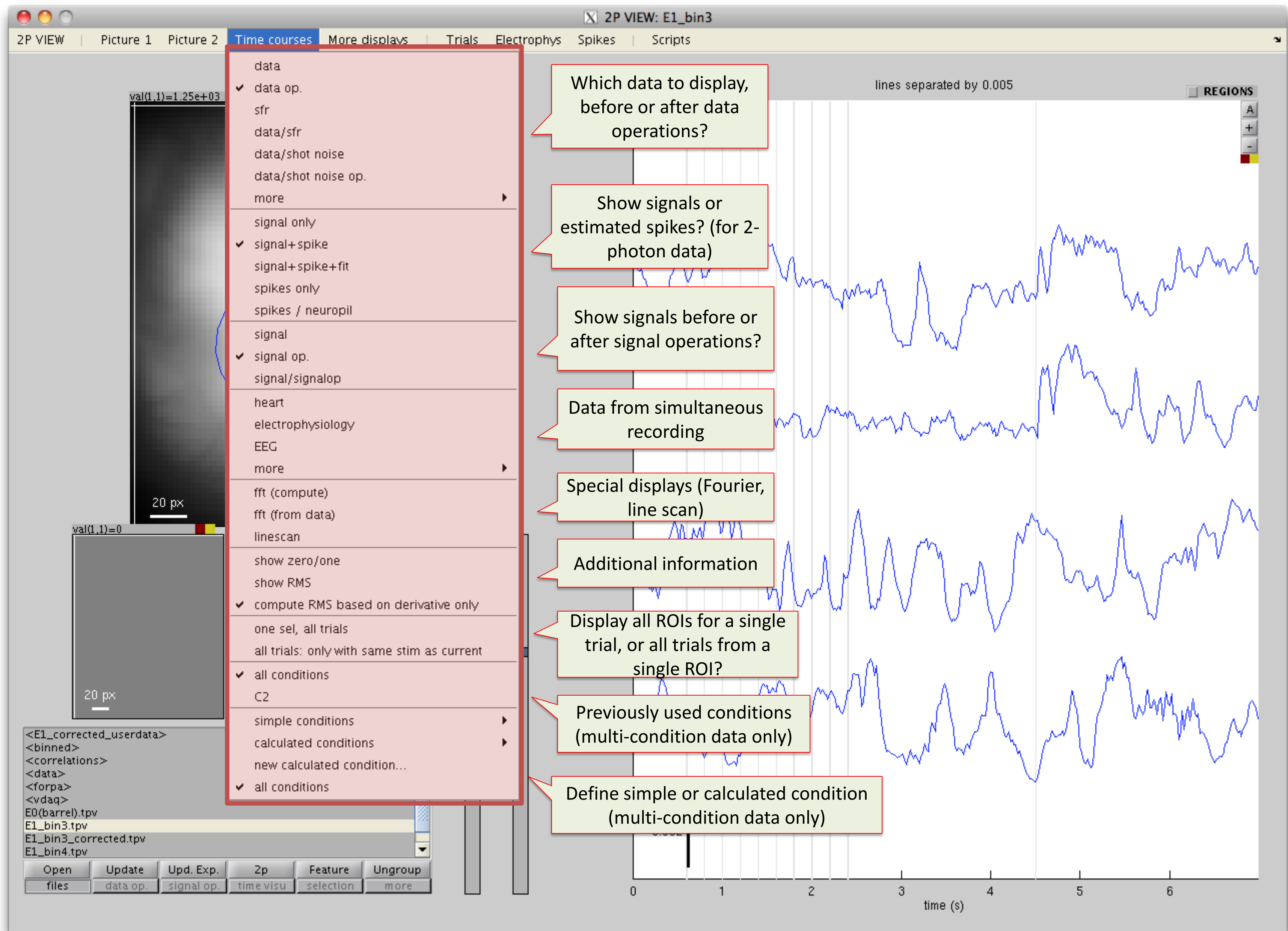
Additional displays  
are available from  
the « More  
displays » menu.



# 2)b. What to display in Picture1 or Picture2



# 2)b. What to display in Time Courses?





# 2)c. Display options (Picture1 and Picture2)

Left-click on the red button to adjust the clipping

Right-click on the red button to make a menu of display options appear

- shape select poly
- shape select free
- shape select rectangle
- ✓ shape select ellipse
- shape select ring

- advanced selection
- ✓ display selection marks
- color selection marks
- reset selection display
- reset selection

- navigation
- scrollwheel zooming

- features
- color map
- clipping mode
- autoclip mode
- user clip
- binning

- distance tool
- show color bar

- duplicate in new figure
- duplicate in ...
- display in base workspace
- save picture
- repair communications

cameraframerate: 100

Shape of the ROI selection tool

Selection and display of ROIs

Rules for mouse control of zoom and pan

Display options: scale bar, color map, rules for changing the clipping, binning...

Additional tools: distance tool; show the color bar

<E1\_corrected\_userdata>  
<binned>  
<correlations>  
<data>  
<forpa>  
<vdaq>  
E0(barrel).tpv  
E1\_bin3.tpv  
E1\_bin3\_corrected.tpv  
E1\_bin4.tpv

Open Update Upd. Exp. 2p Feature Ungroup  
files data op. signal op. time visu selection more

2P VIEW: E1\_bin3  
Electrophys Spikes Scripts

lines separated by 0.005

REGIONS

A  
+  
-  
■

20 px

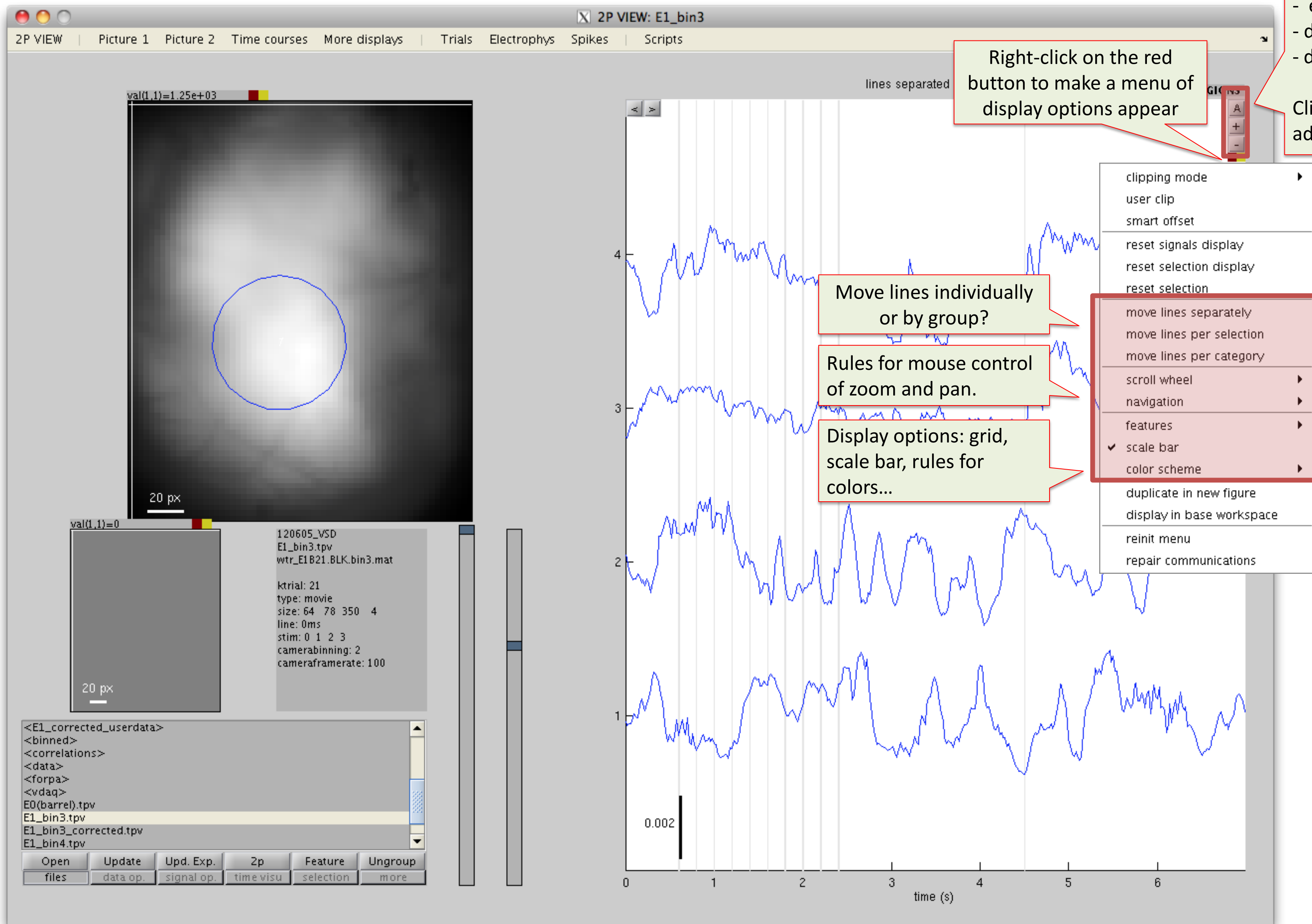
val(1,1)=0

20 px

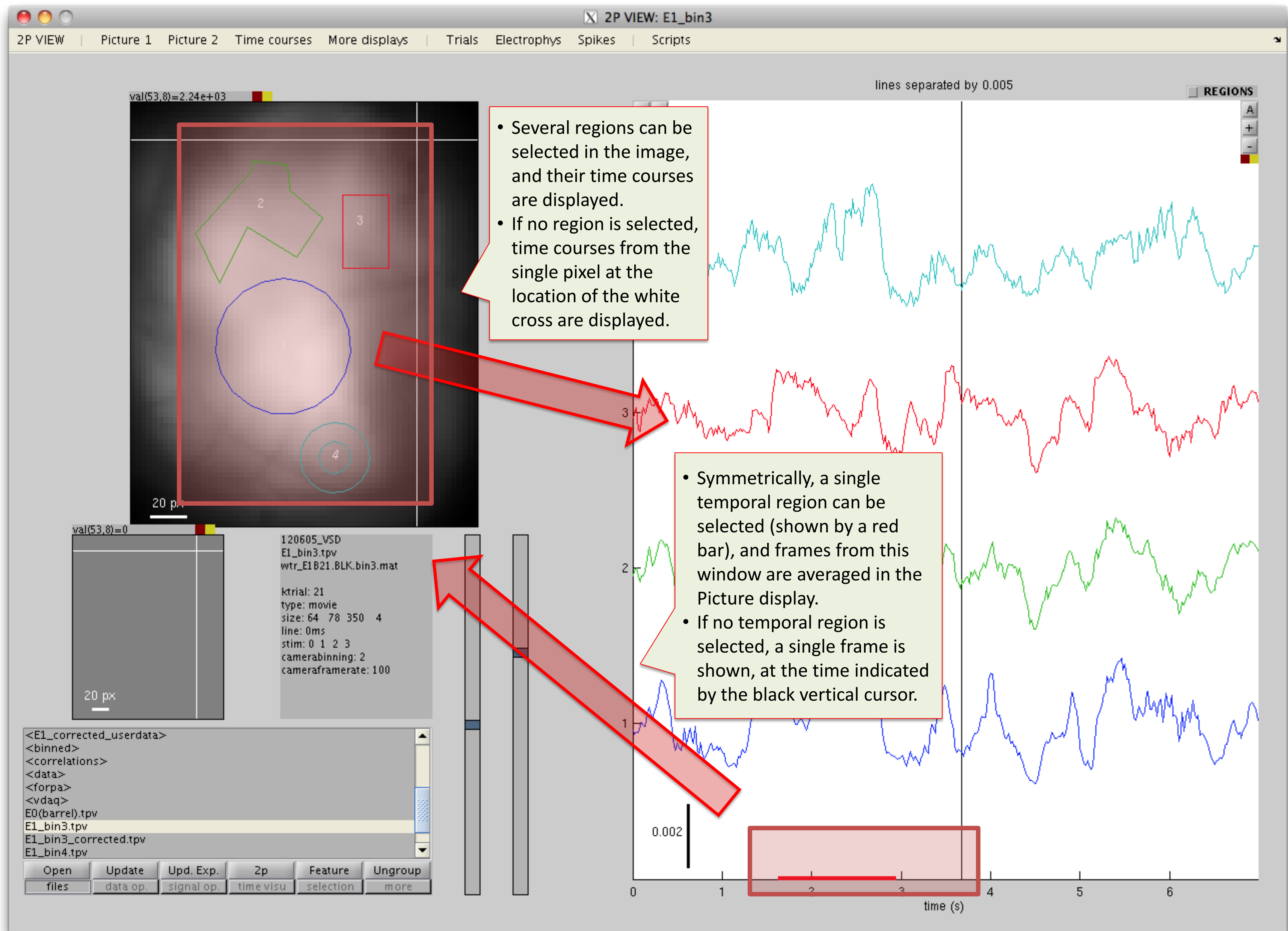
0.002

time (s)

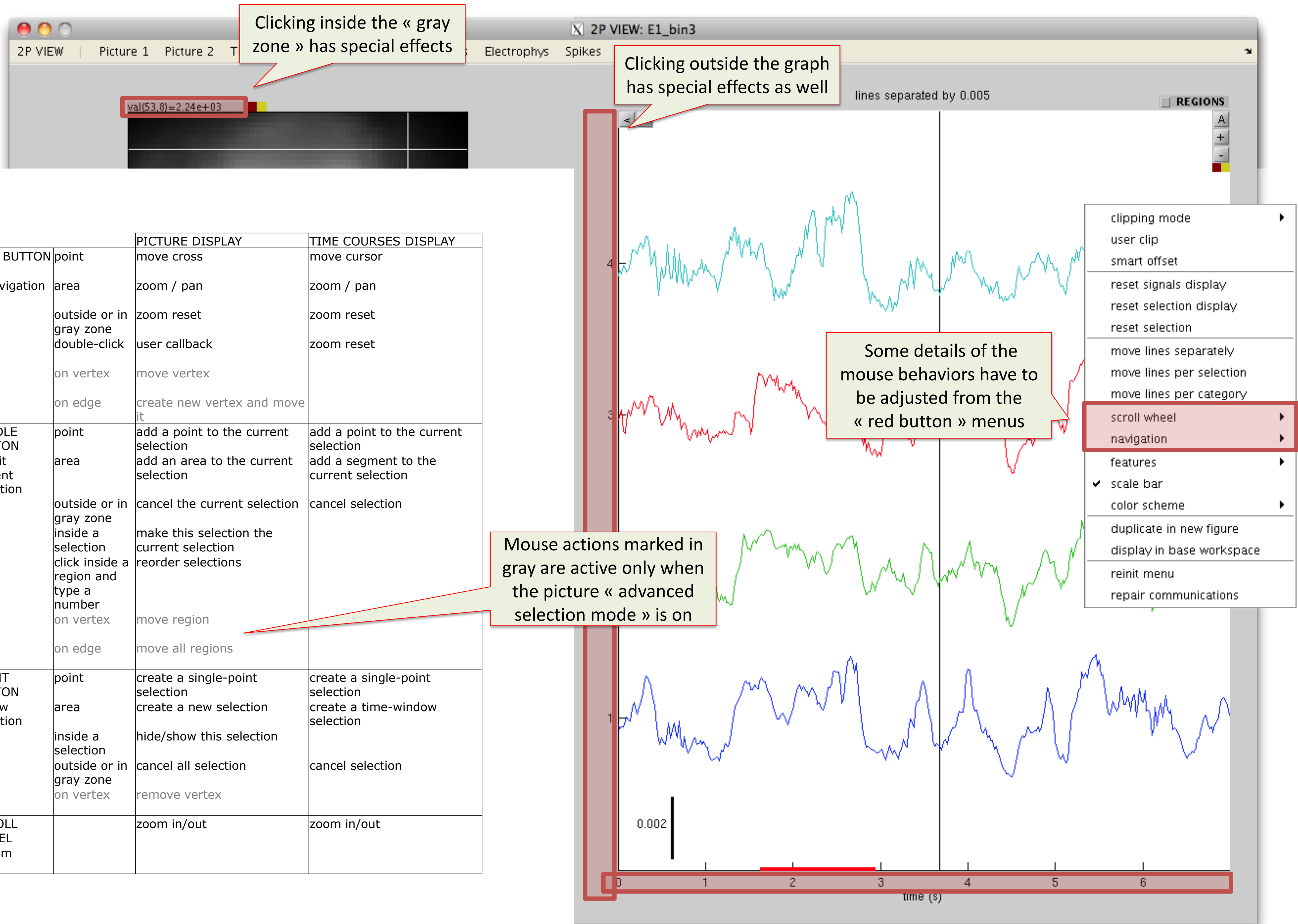
# 2)c. Display options (Time Courses)



# 2)d. Principles of region selection



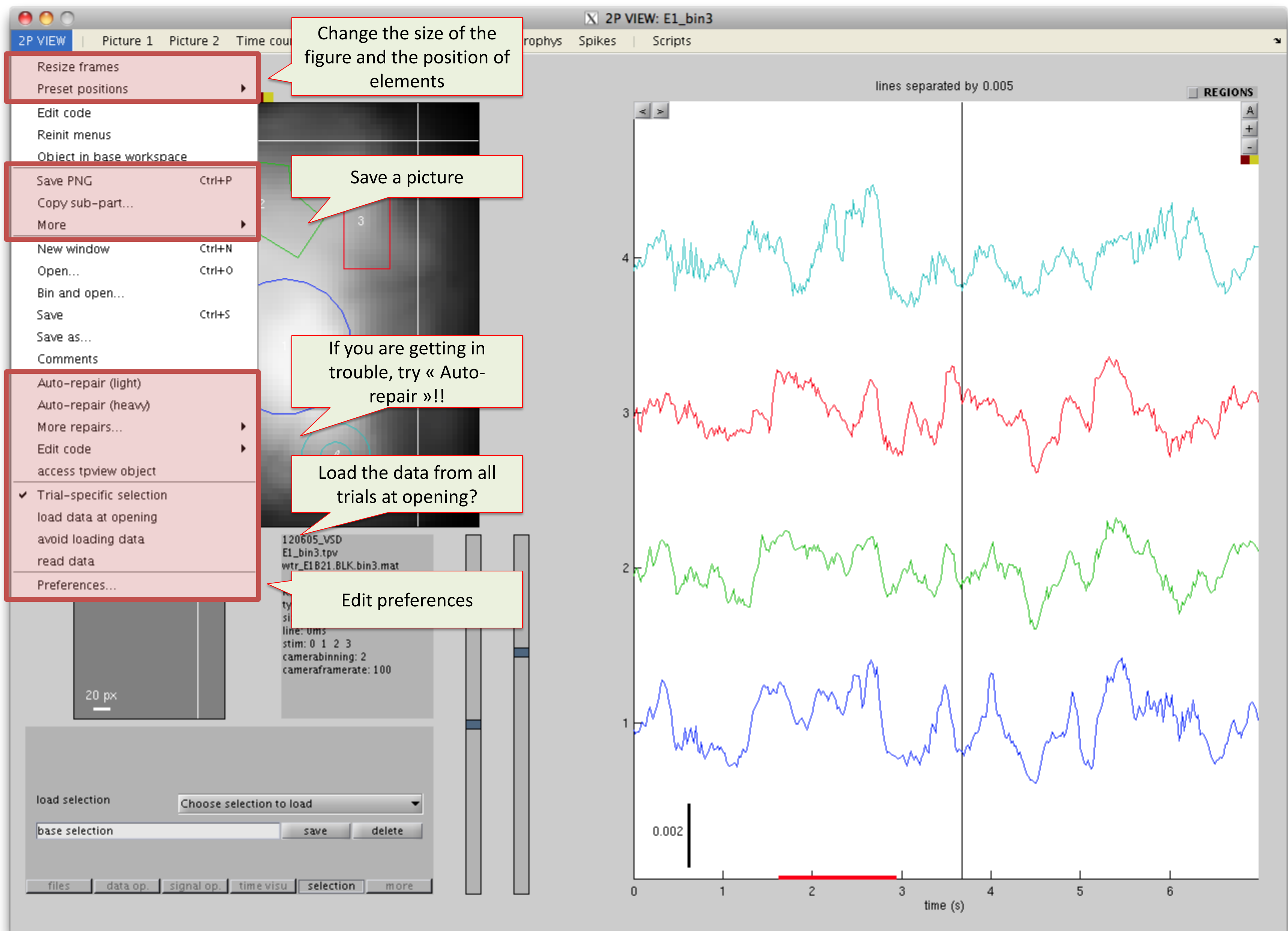
# 2)d. Mouse actions



# 3) Other useful tools

- a. The 2Pview menu

# 3)a. The « 2Pview » menu

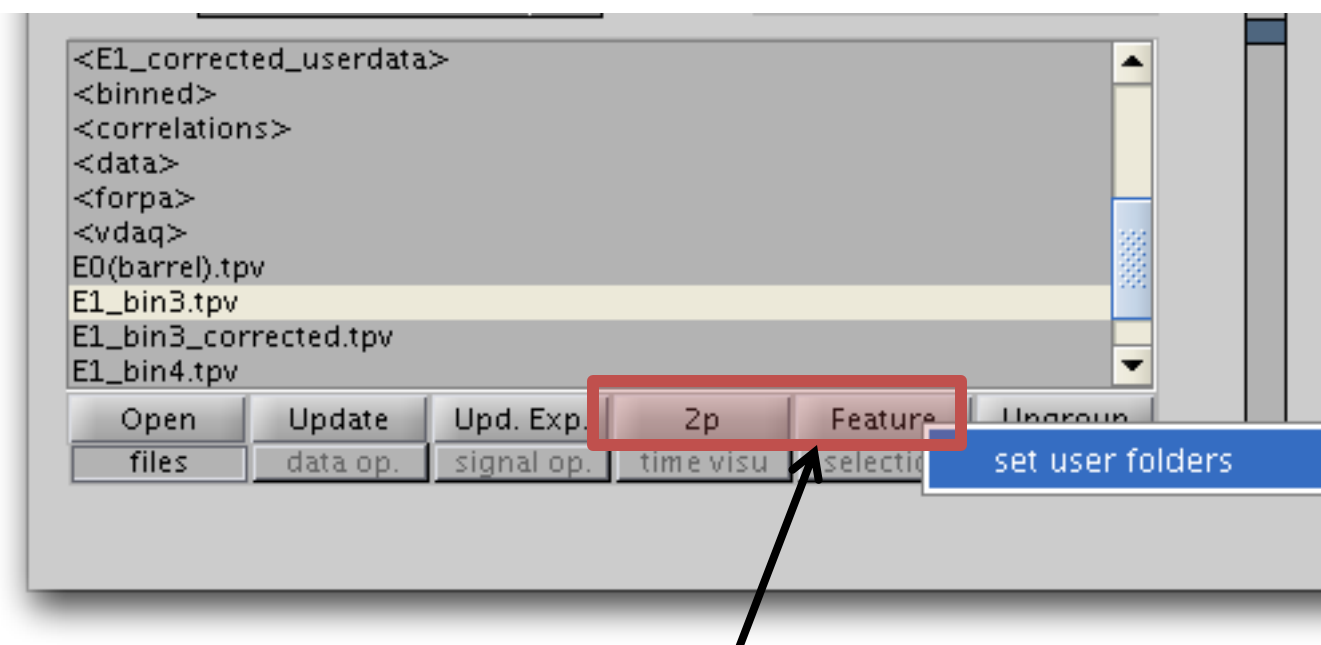




## II. In details

- 1) Manipulation of the data
- 2) Display of the data
- 3) Other useful tools

# 1)a. The « file panel »

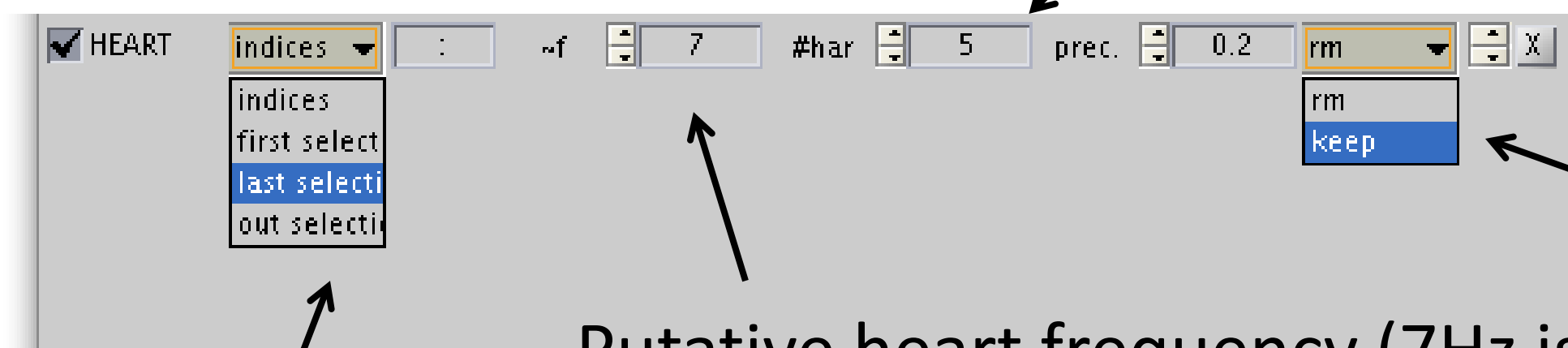


These two buttons provide the ability to create shortcuts to your two preferred data folders. Right-click on any of them to define these folders.

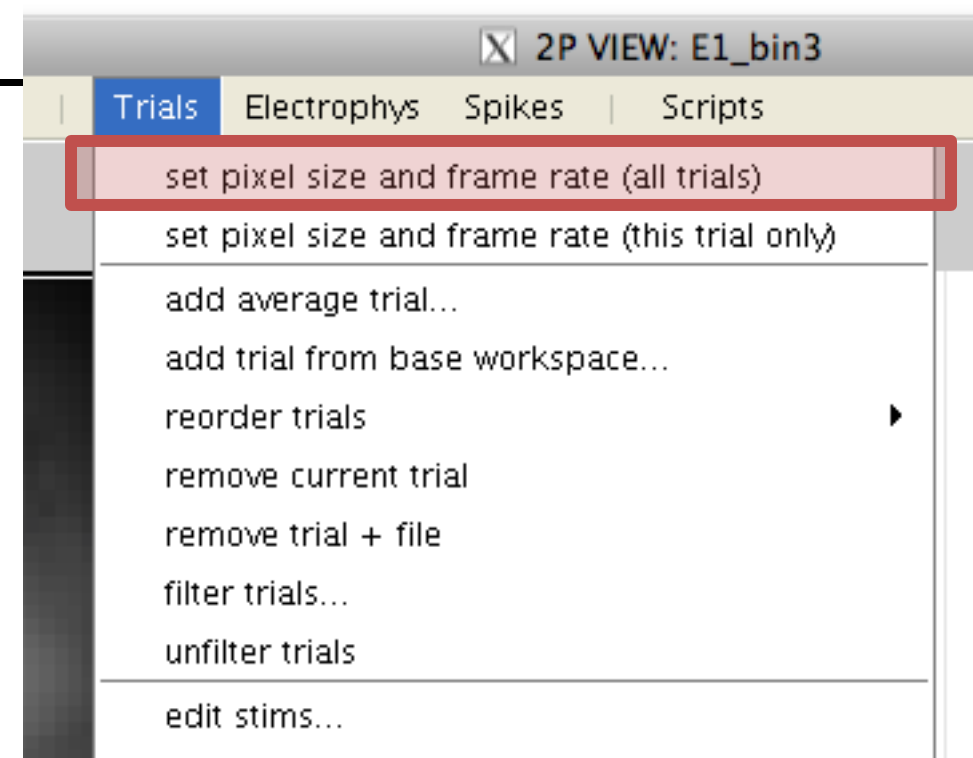
# 1)c. Heart correction

These are the recommended parameters for heart correction on rat.

Number of harmonics to use (1 would result in a sinus estimation, more harmonics capture more high frequencies)



Putative heart frequency (7Hz is fine for rat. Note that it is necessary that the frame duration has been defined accurately before (here:

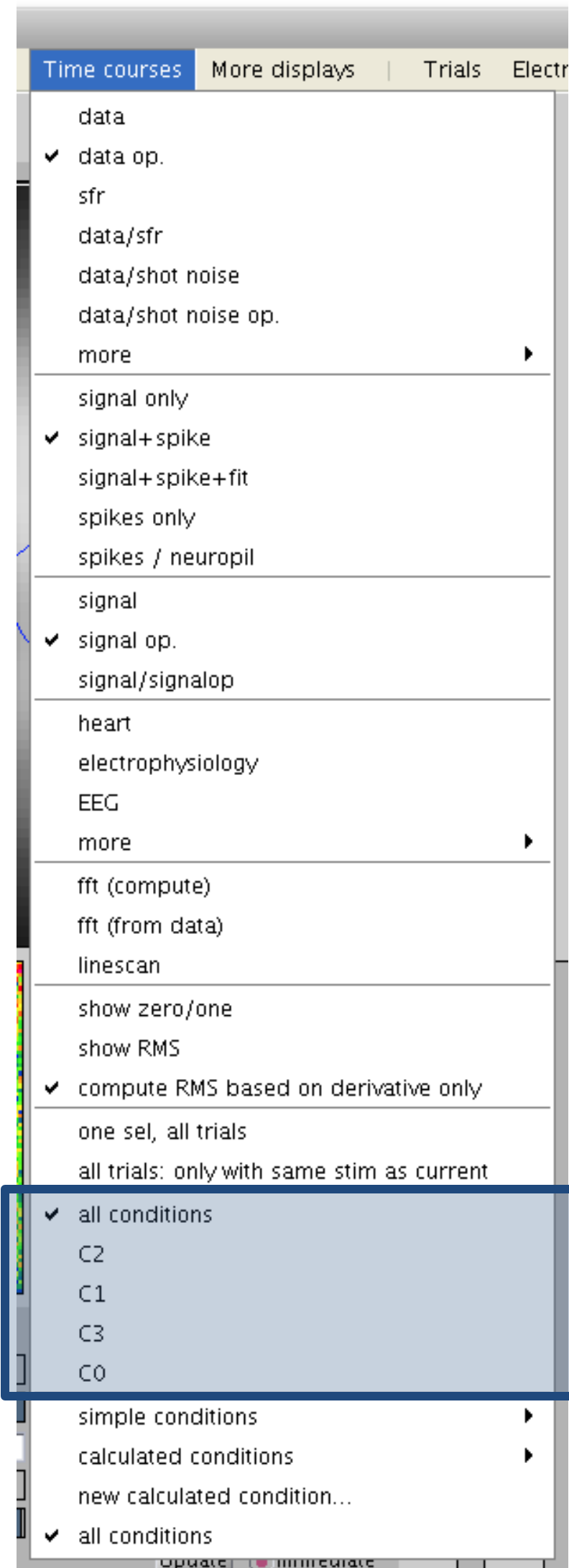


Choose whether to remove the estimated heart or keep only it. It is advised to first select 'keep' in order to check that it has been correctly estimated, then select 'rm' to perform the correction.

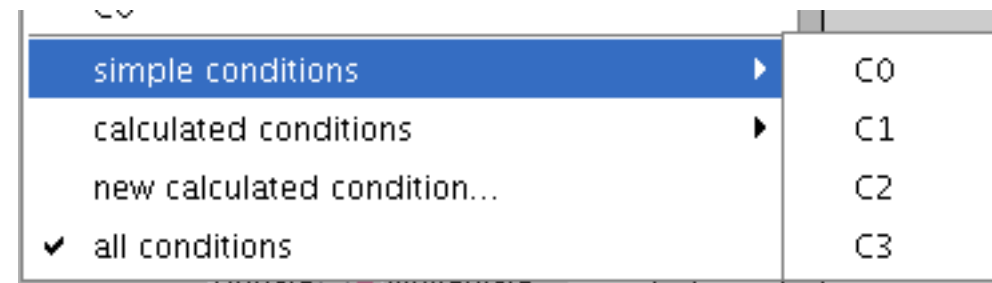
First select a region of interest and choose option 'last selection' to tell that heart beat times should be estimated using that region.

# 2)b. Which condition(s) to display?

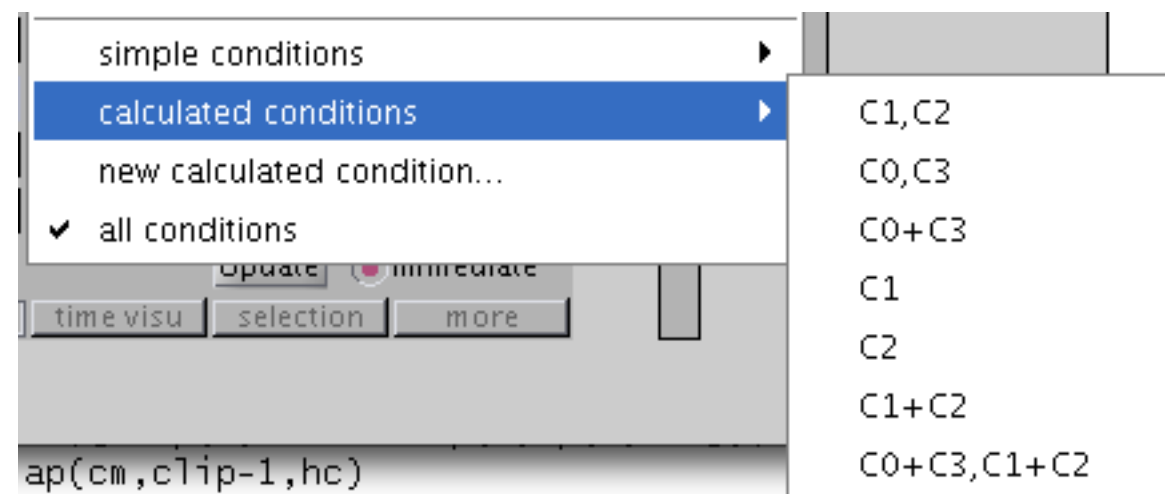
When data has multiple conditions, it is possible to choose which to display in Picture1, Picture2 or Time Courses. Several options exist:



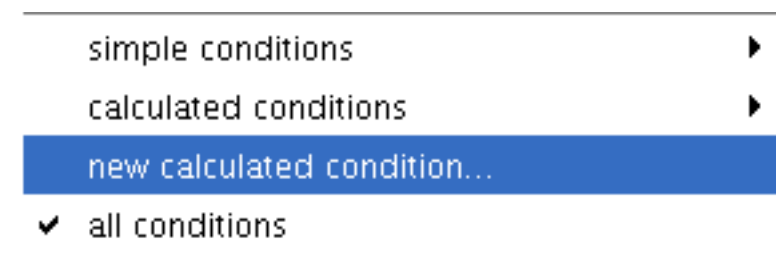
1) Choose one of the recently used conditions



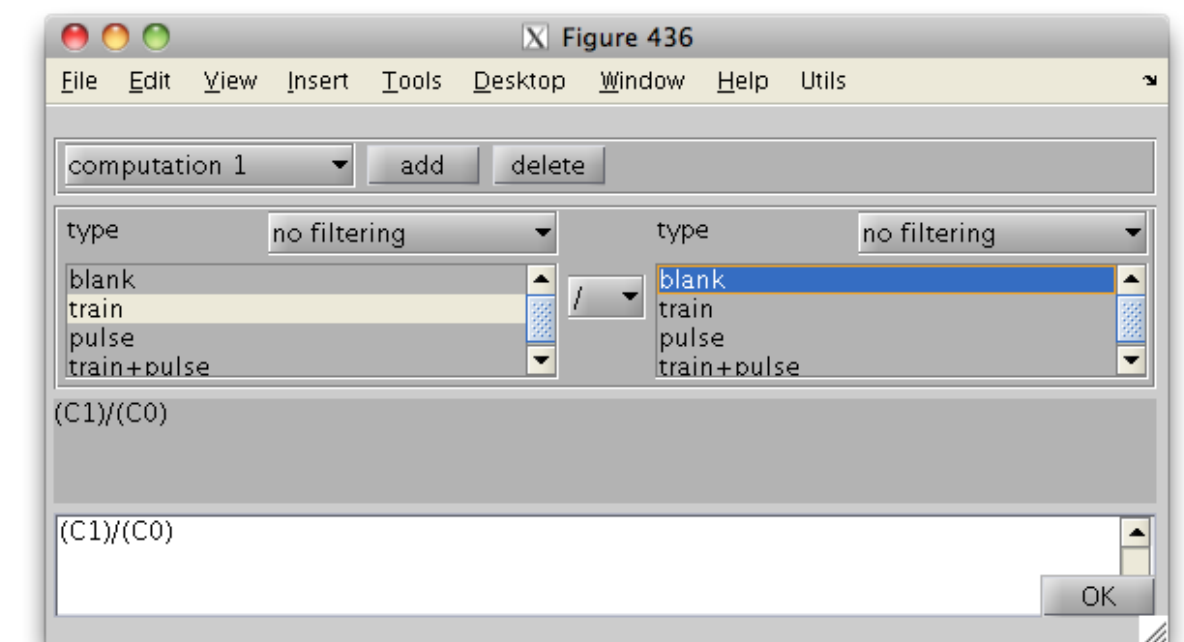
2) Choose a simple conditions



3) Choose a recent calculated condition



4) Create a new calculated condition: a new window opens that lets you define it



5) Display all simple conditions together

