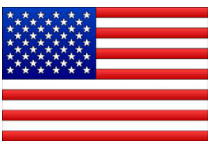


AFNI & SUMA

Concepts, Principles, Demos



Analysis of Functional NeuroImages

by

Robert W Cox, PhD

Released under the GNU General Public License Version 2 (GPL)
[or any later GPL version]

AFNI is a research tool.

*Clinical uses are **not supported or advised.***



AFNI User



<http://afni.nimh.nih.gov/afni>

Some Goals of fMRI Analyses

- **Task-based experiments**
 - Per subject: estimate amplitude of BOLD response to each different type of stimulus
 - Find+model inter-regional correlations between fluctuations in BOLD responses

- **Resting-state experiments**
 - Measure spatial patterns in coherent fluctuations in spontaneous BOLD

- **Group level**
 - Combine and contrast per subject results

Conceptual Basis - 1

- **Time shifting** = pretend get 3D snapshot

- **Despiking** = remove large blips

- **Image Registration** (AKA alignment)
 - intra-EPI time series, and EPI-Structural

- **Blurring in space** = lower resolution :-(
& less noise :-) & more group overlap :-)

- **Masking** = ignore non-brain voxels

- **Scaling** = normalizing data amplitude
 - Makes inter-subject comparisons more valid

Conceptual Basis - 2

★ Time series regression

- model of the BOLD response in the data = Hemodynamic Response Function ⊗ stimulus timing
- plus baseline (null hypothesis) model
- plus physiological noise
- plus allowing for serial correlation

★ Talairach-ing = Spatial Normalization

- Talairach, MNI-152, ...
- affine and nonlinear spatial transformations

Conceptual Basis - 3

★ **Group Analyses** = Putting it all together

★ – ANOVA, LME, Meta-Analyses, ...

• **Blobs** = Spatial models of activation

★ – Assigning statistical significance to blobs

• **Connectivity** = Inter-regional analyses

– SEM, PPI, SVAR, DCM, Granger, ...

– Resting state fMRI (Connectome!)

• **Dimensional factorization**

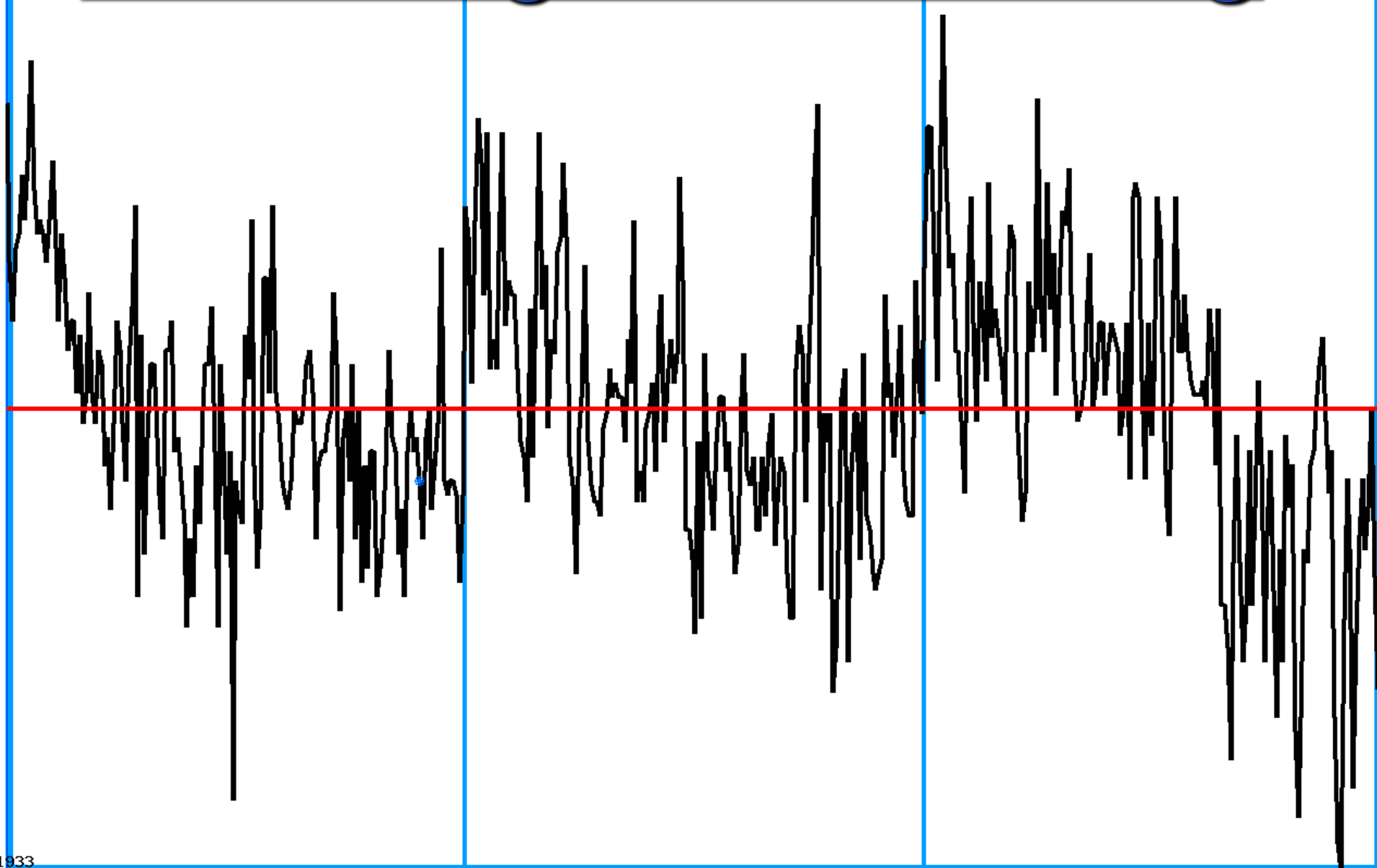
Components, such as PCA, ICA

Conceptual Basis - 4

- **Data Formats** = NifTI-1.x is your friend
- **Software** for fMRI analyses: *open-source
 - **AFNI***, BrainVoyager, FSL*, SPM*, ...
 - Whichever you use, **don't blindly assume** the software works perfectly all the time
- Most important thing I will say today
Understand and check the steps applied to your data!
- 2nd most important: **Is no "best" way to analyze data, just "reasonable" ways**

102.1128
[+4.09346]

Linear Regression = Fitting

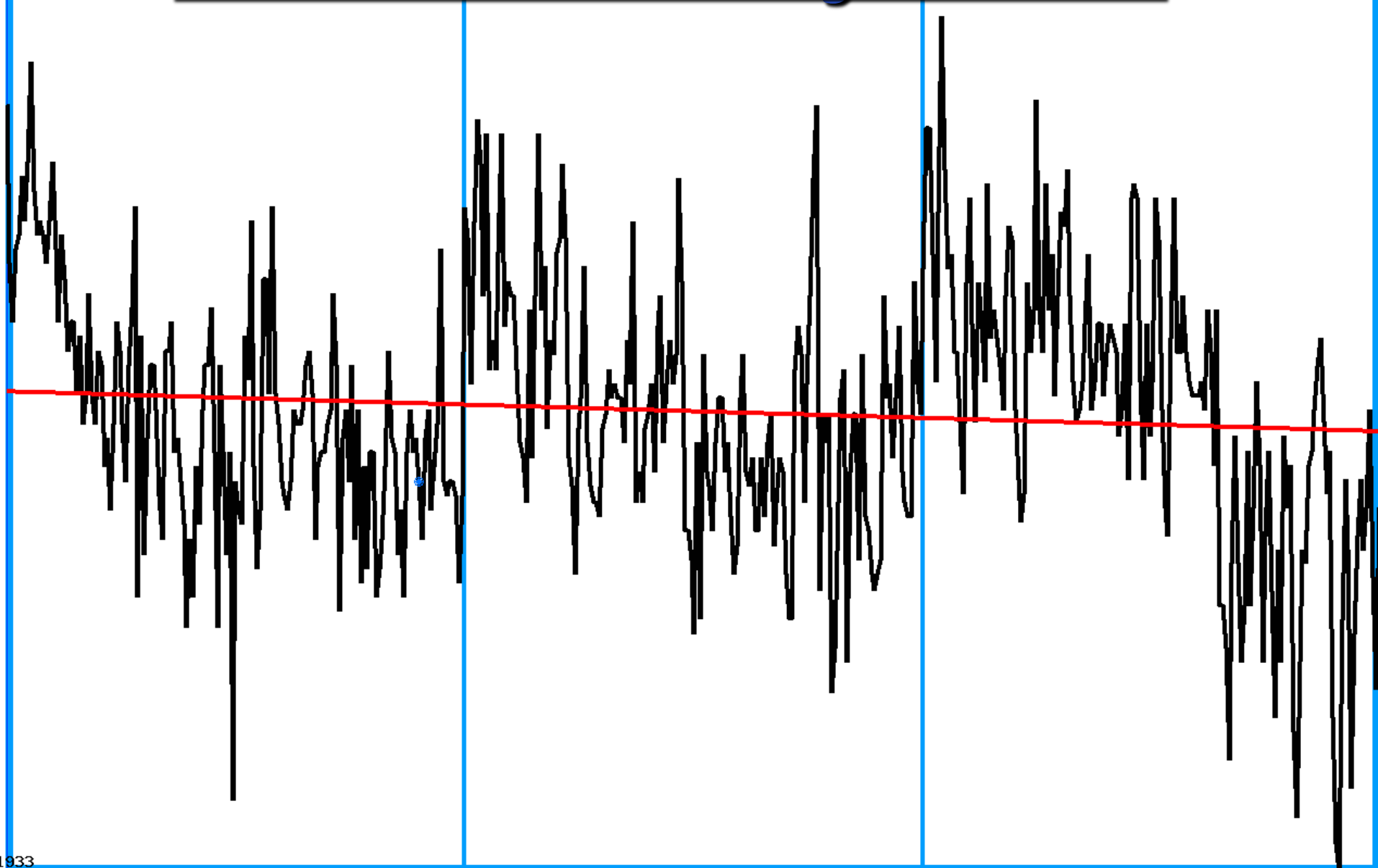


98.01933

AXIAL I: 48 indx=135 val=99.69785 @t=270
J: 27 Grid: 150 Scale: 214 pix/datum Mean: 100.0001 Tran OD = -none-
AFNI K: 28 # 0:449 Base: separate Sigma: 0.572177 Tran 1D = L1_Fit

102.1128
[+4.09346]

Fit = Linear Polynomial



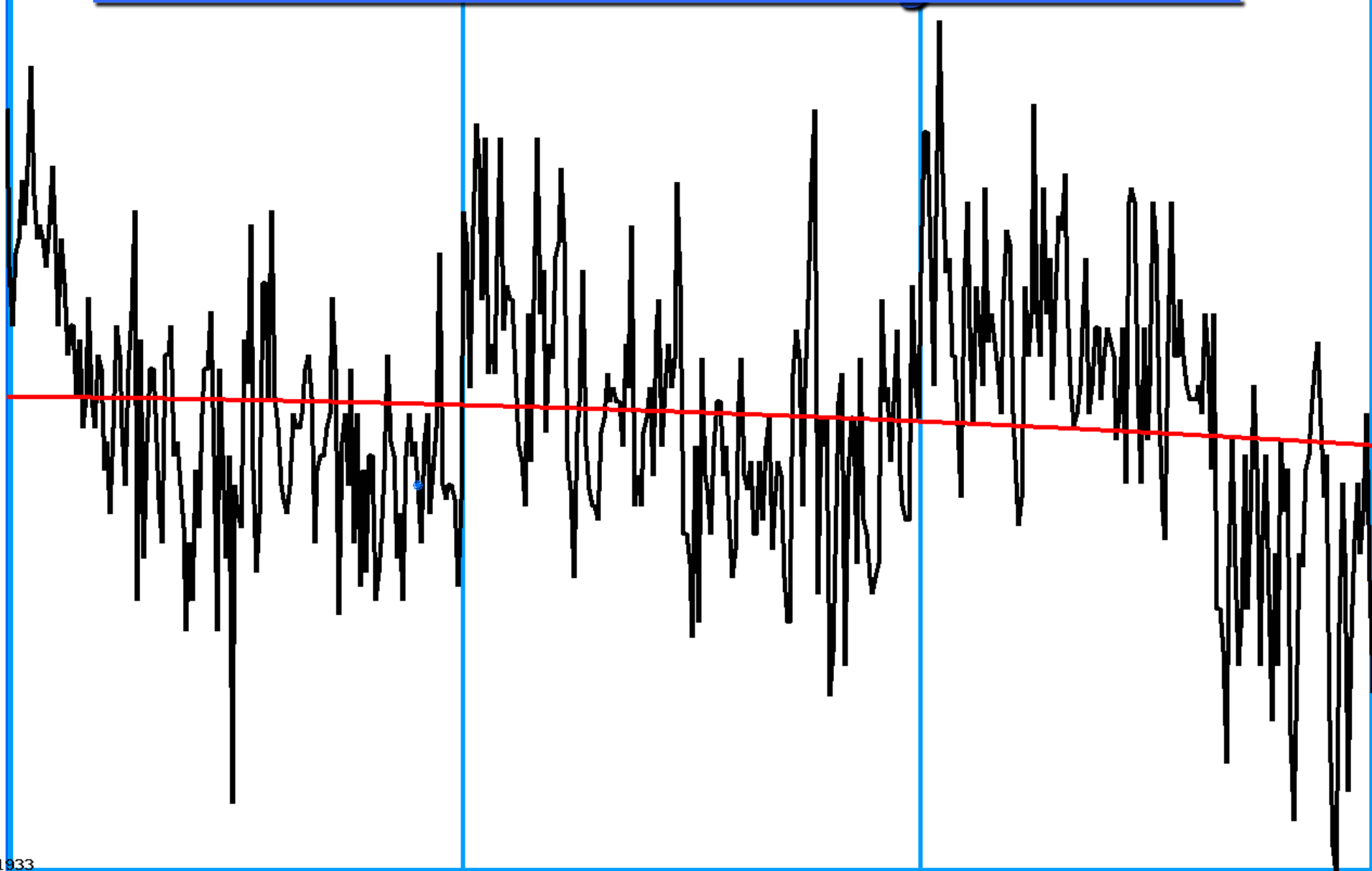
98.01933



I: 48 | [indx=135](#) | [val=99.69785](#) | [@t=270](#)
J: 27 | Grid: 150 | Scale: 214 pix/datum | Mean: 100.0001 | Tran OD = -none-
K: 28 | # 0:449 | Base: separate | Sigma: 0.572177 | Tran 1D = L1_Fit

102.1128
[+4.09346]

Fit = Quadratic Polynomial

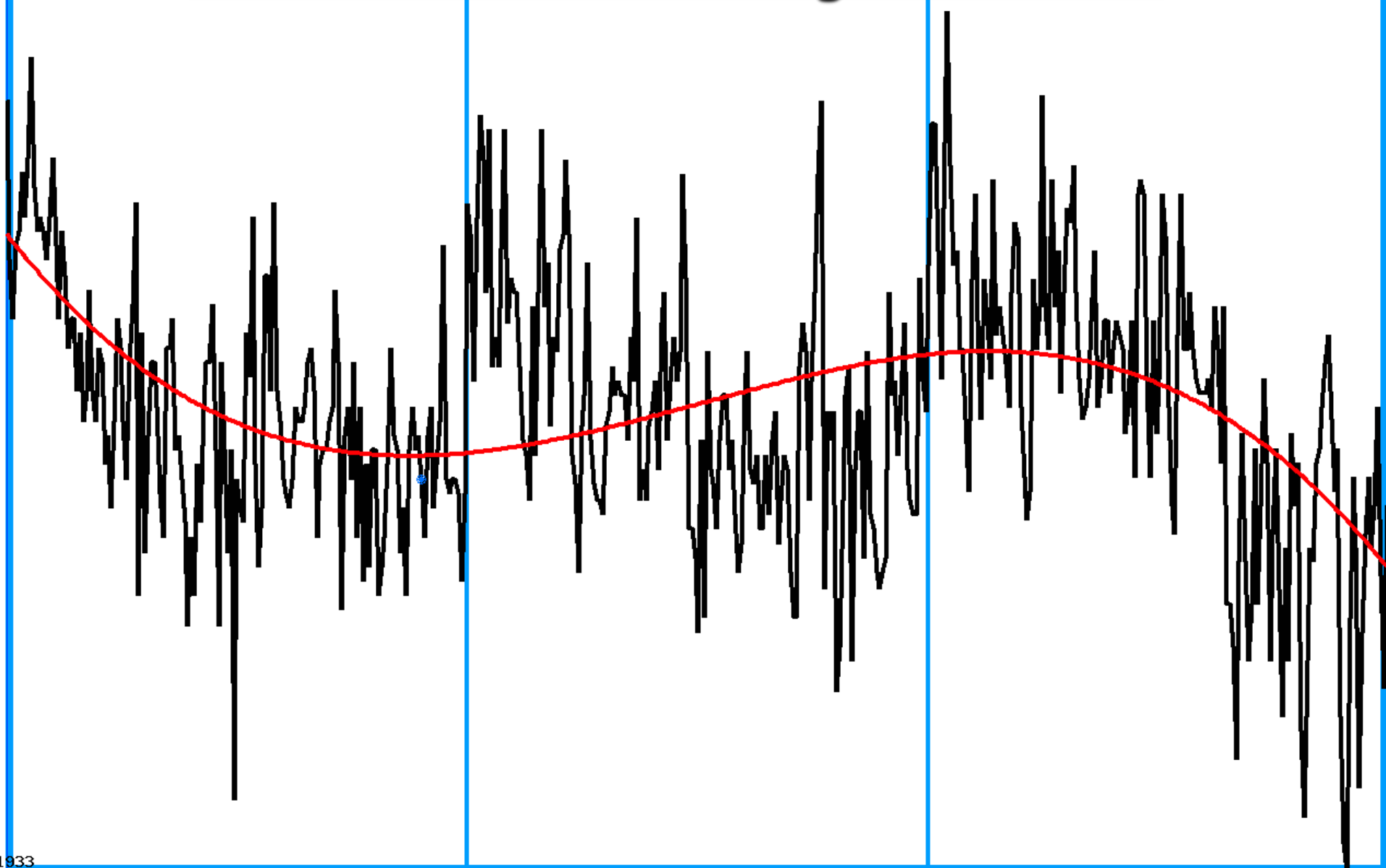


98.01933

AXIAL I: 48 indx=135 val=99.69785 @t=270
AFNI J: 27 Grid: 150 Scale: 214 pix/datum Mean: 100.0001 Tran OD = -none-
K: 28 # 0:449 Base: separate Sigma: 0.572177 Tran 1D = L1_Fit

102.1128
[+4.09346]

Fit = Cubic Polynomial



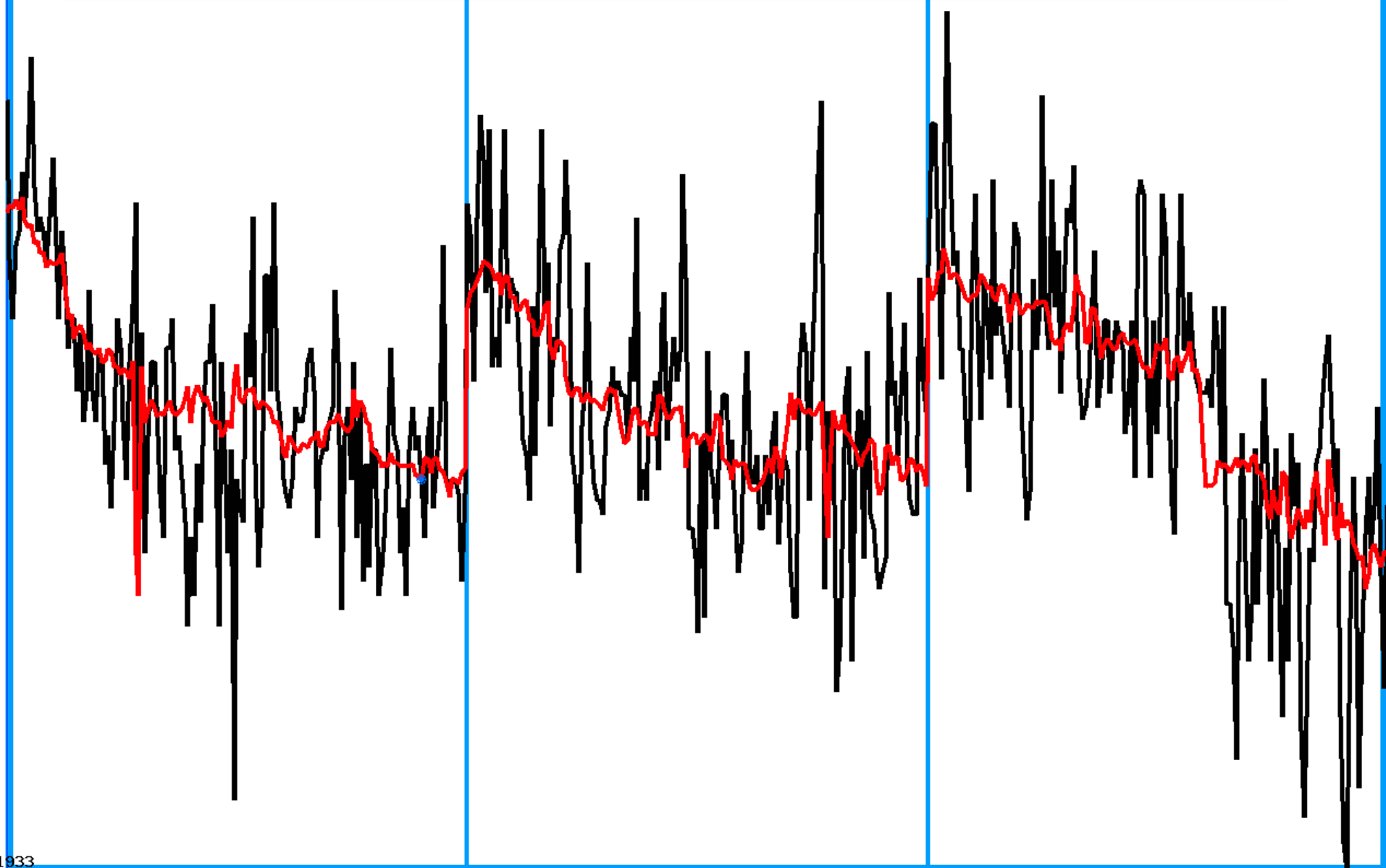
98.01933



I: 48 | [indx=135](#) | [val=99.69785](#) | @t=270
J: 27 | Grid: 150 | Scale: 214 pix/datum | Mean: 100.0001 | Tran OD = -none-
K: 28 | # 0:449 | Base: separate | Sigma: 0.572177 | Tran 1D = L1_Fit

102.1128
[+4.09346]

Fit = Cubic + Motion Params



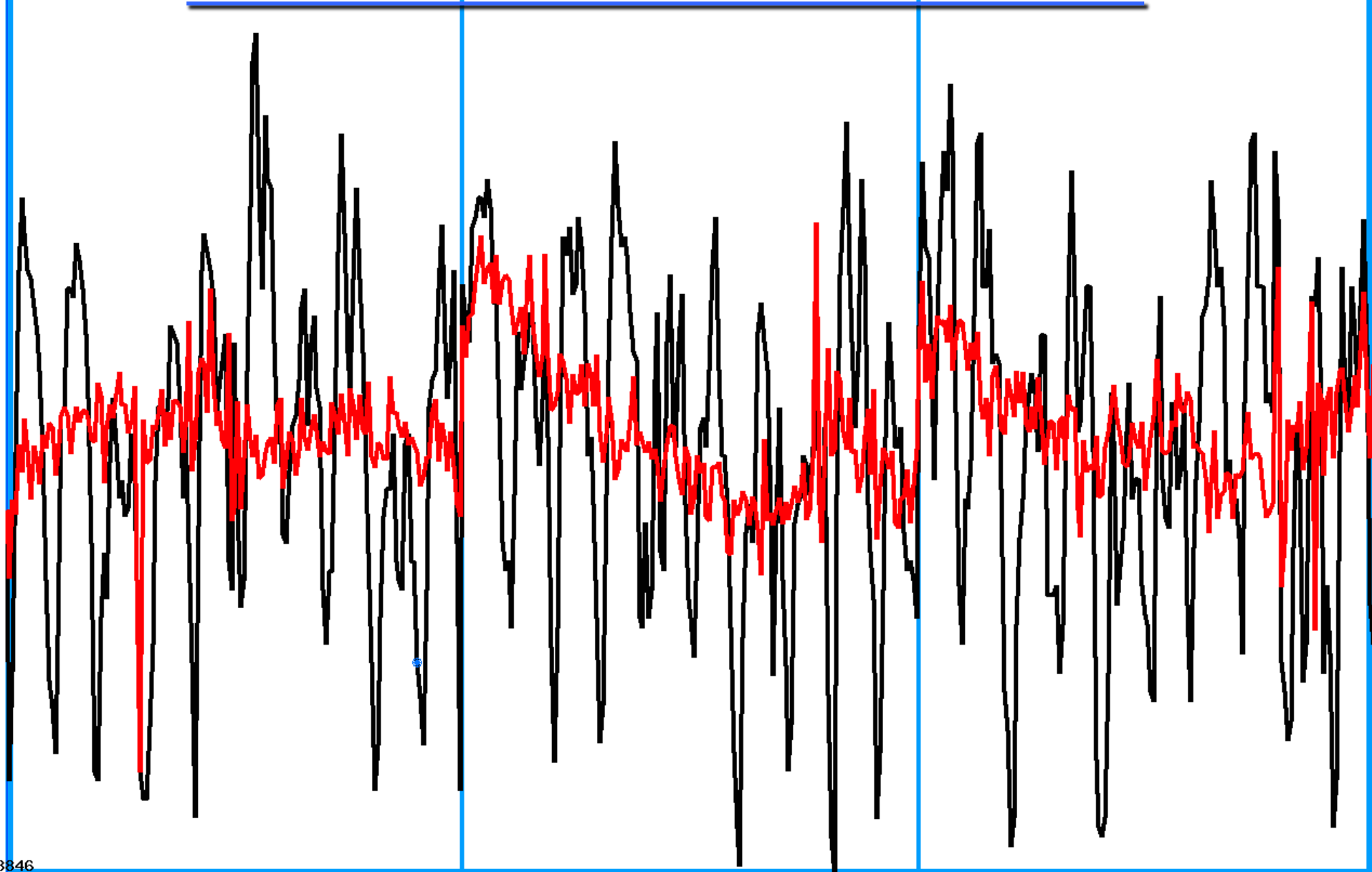
98.01933



I: 48 | [indx=135](#) | [val=99.69785](#) | @t=270
J: 27 | Grid: 150 | Scale: 214 pix/datum | Mean: 100.0001 | Tran OD = -none-
K: 28 | # 0:449 | Base: separate | Sigma: 0.572177 | Tran 1D = L1_Fit

108.7157
[+15.92727]

Fit in a Different Voxel



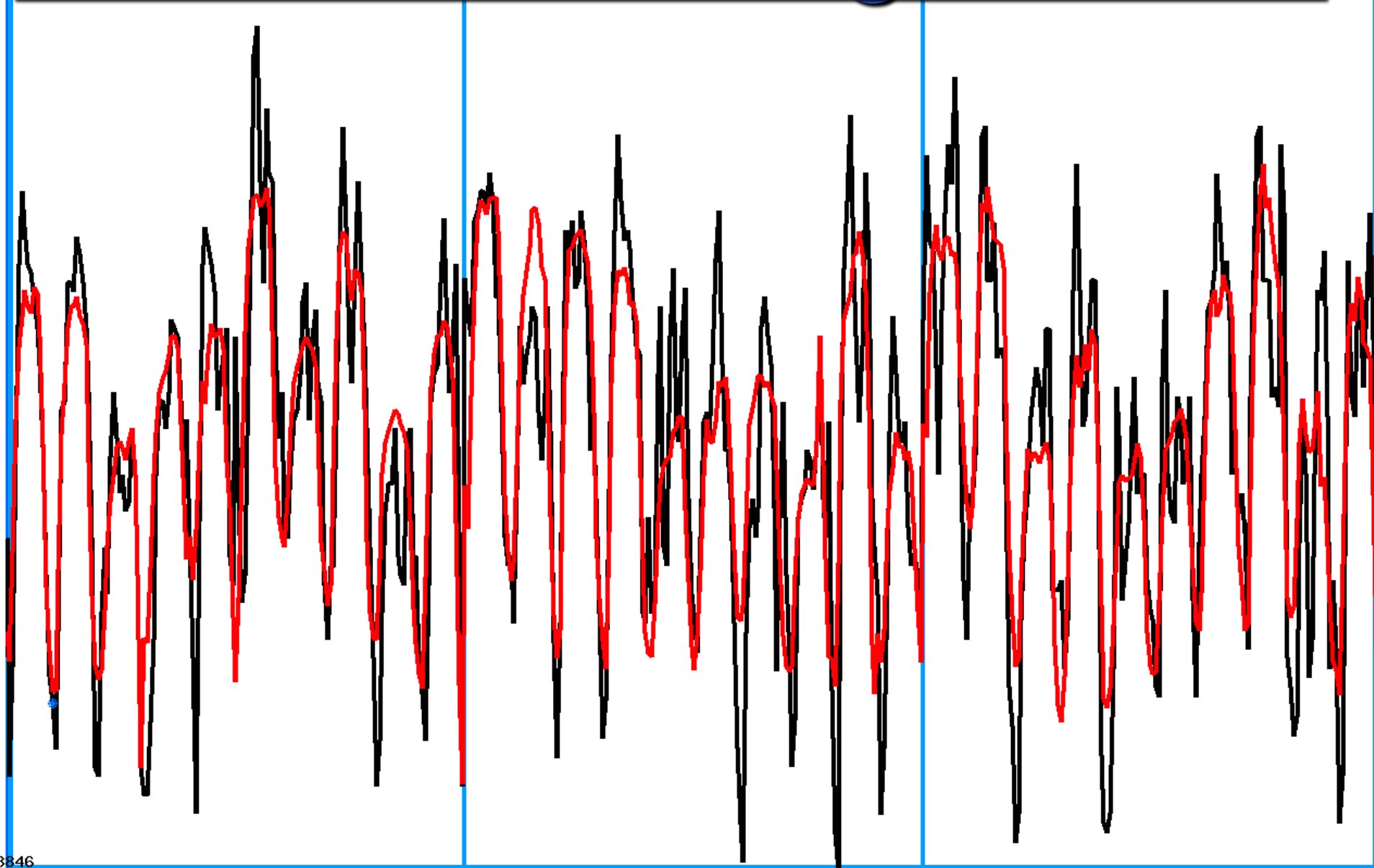
92.78846



I: 57 | **indx=135 val=96.38354 @t=270**
J: 45 | Grid: 150 | Scale: 55 pix/datum | Mean: 100. | Tran OD = -none-
K: 28 | # 0:449 | Base: separate | Sigma: 2.8749 | Tran 1D = L1_Fit

108.715
[+15.927 71]

Fit Now Includes Signal Model



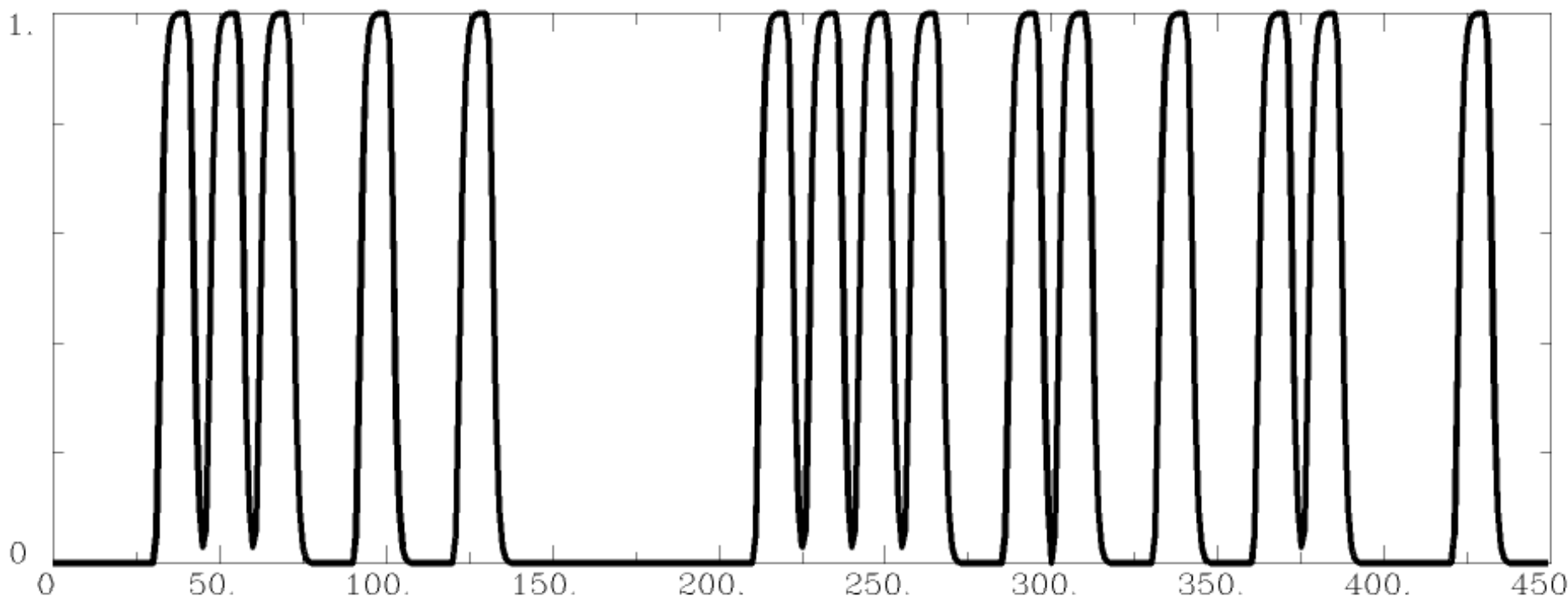
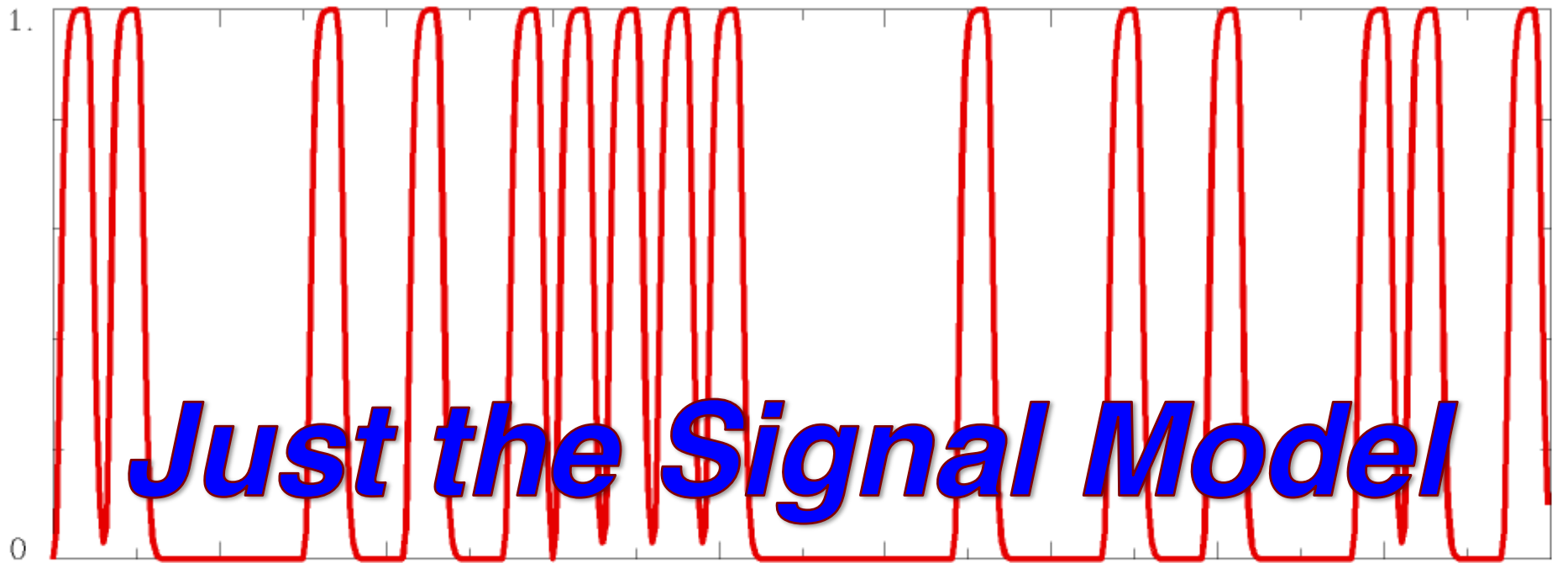
92.78846

AXIAL I: 57 | **indx=15 val=95.62058 @t=30**
J: 45 | Grid: 150 | Scale: 55 pix/datum | Mean: 100. | Tran OD = -none-
AFNI K: 28 | # 0:449 | Base: separate | Sigma: 2.8749 | Tran 1D = Dataset#N



**All The Regressors
= The Full Model**

Just the Signal Model



Analysis by Super-Script – by GUI

The screenshot shows the Super-Script GUI with the following sections:

- general subject info**: subject ID group ID
- input data and options**:
 - anatomical dataset
 - Buttons: browse anat, clear anat, ? help: anat
 - Text field:
 - include copy of anat+tlrc
 - EPI datasets
 - Buttons: browse EPI, clear EPI, ? help: EPI
 - Table:

	scan index ▼	EPI dataset
1	1	FT_epi_r1+orig.HEAD
2	2	FT_epi_r2+orig.HEAD
3	3	FT_epi_r3+orig.HEAD
 - EPI directory
 - wildcard form

