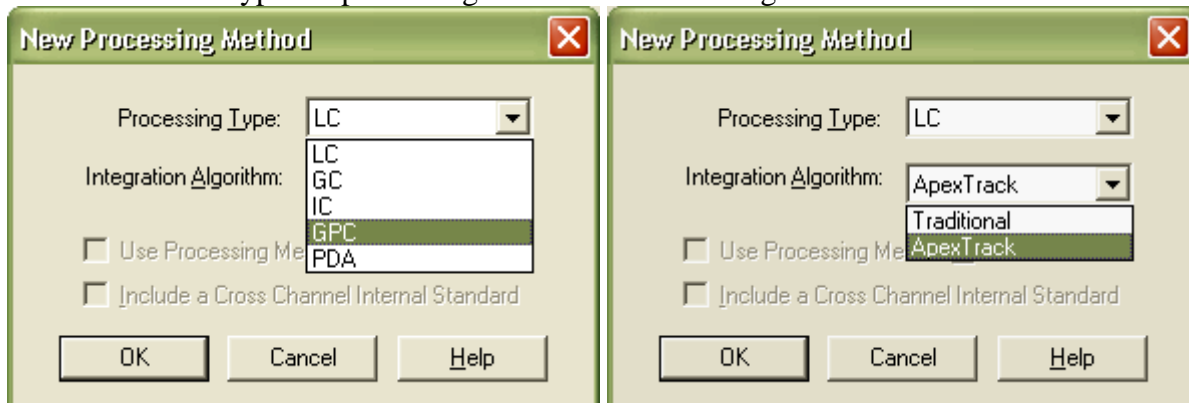


## The Processing Method in Empower 2

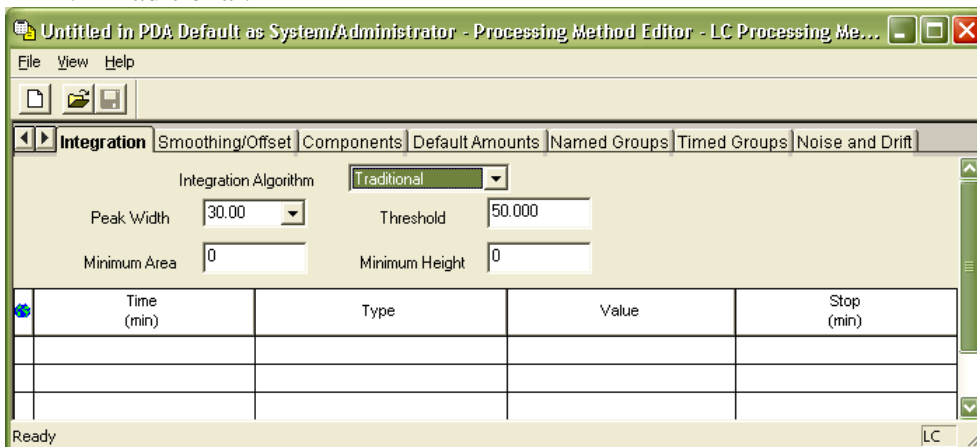
There are various types of processing methods and two integration modes:



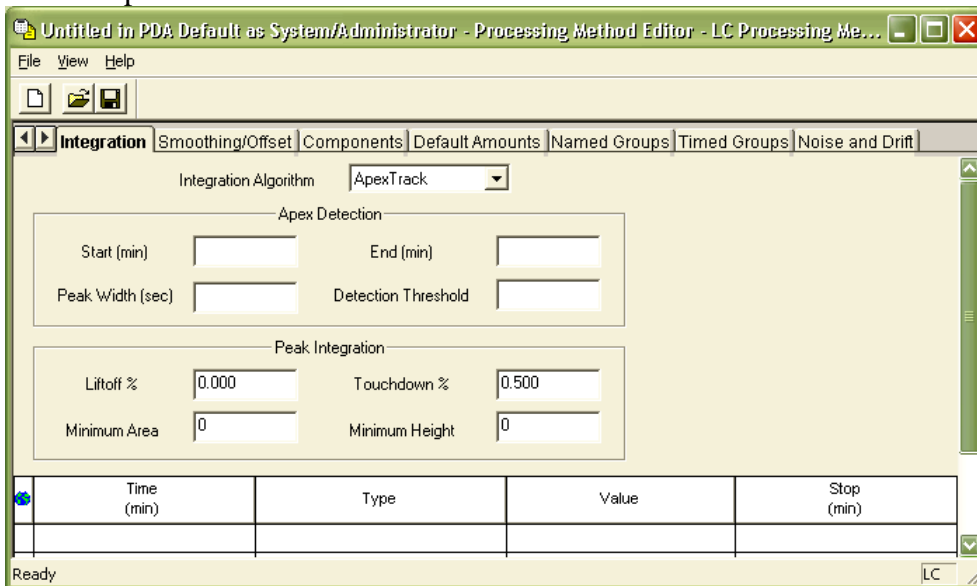
### Screens in the LC Processing Method:

#### Integration:

1. Traditional:



2. Apex Track:

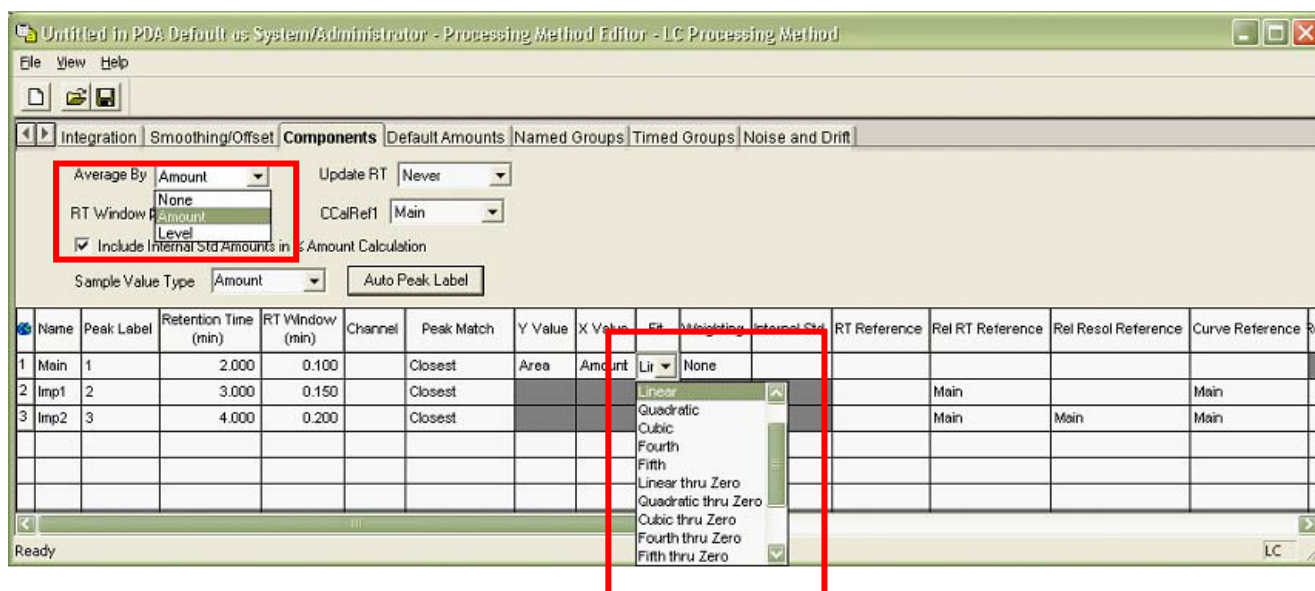


For more details on integration events: [http://www.forumsci.co.il/HPLC/Integration\\_events.pdf](http://www.forumsci.co.il/HPLC/Integration_events.pdf)

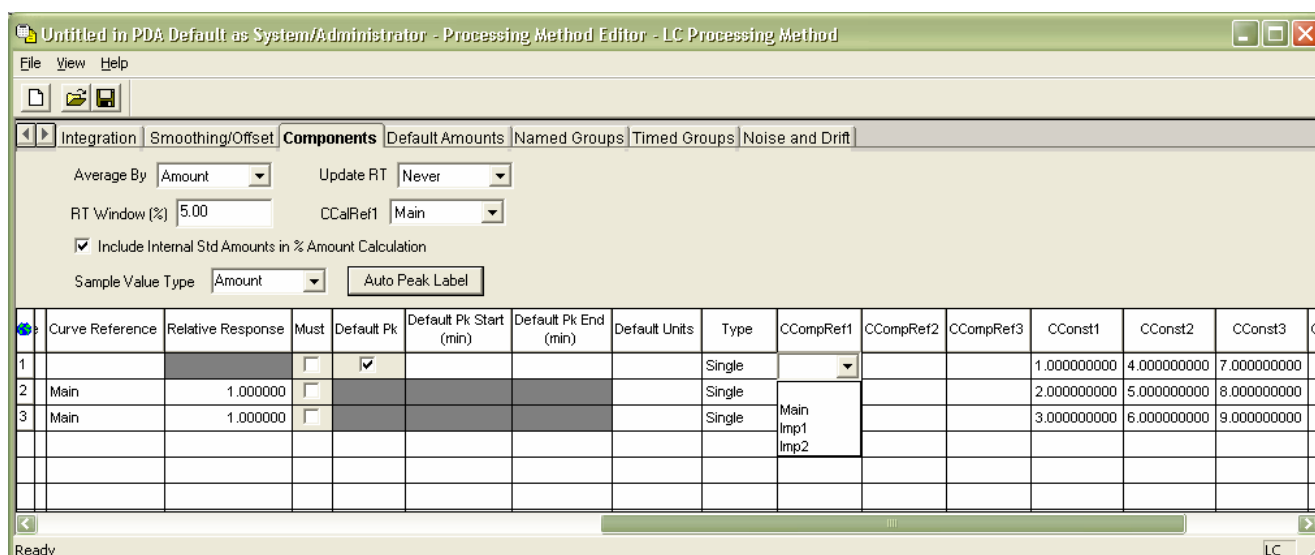


## Average by

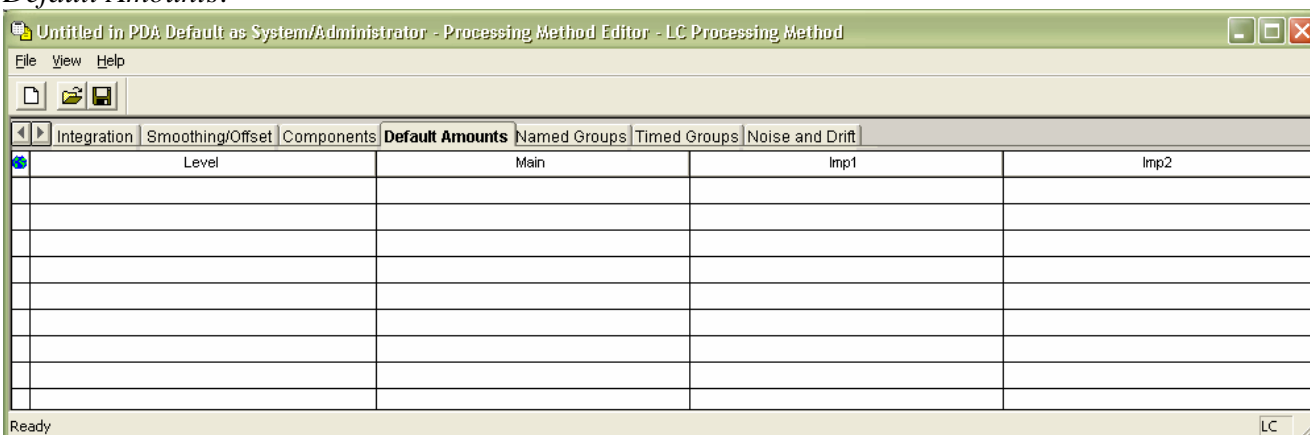
Set it to *Amount* or *Level* or *None* to average calibration standards that generate a calibration curve data in the Calibration Curve window. For example, if your protocol states that the standards should be averaged: **you must use the *Average by* option.**



The *Components Table* include parameters such as fit type (*Linear* or *Linear thru Zero*), RRT (*Rel RT Reference*), RRF (*Curve Reference*), defining one of the peaks as the Default Peak for calculations of *Amount* and RRT of unknown peaks, inter-peaks calculations pointers (*Ccompref1-3*) and values for custom calculations (*Cconst1-7*).



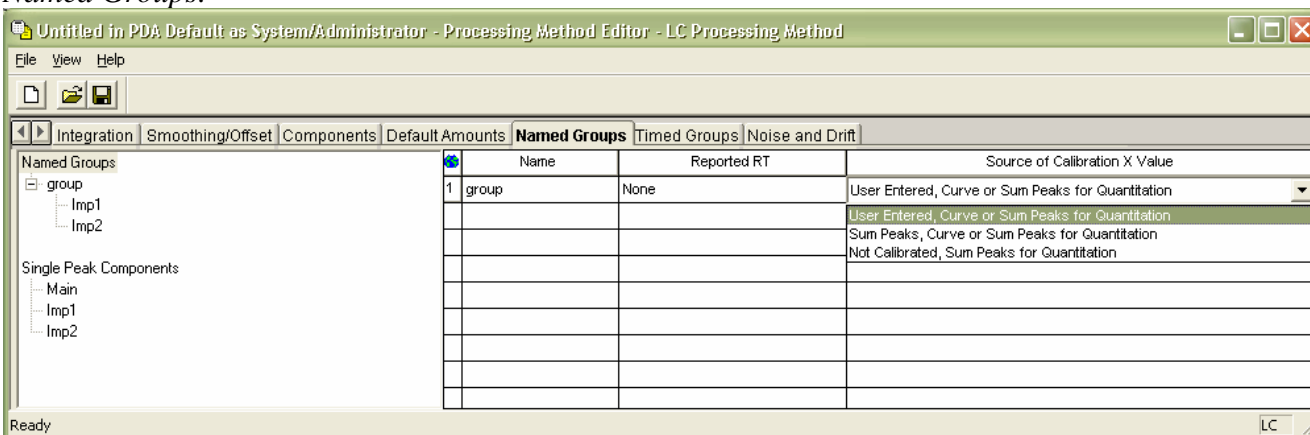
**Default Amounts:**



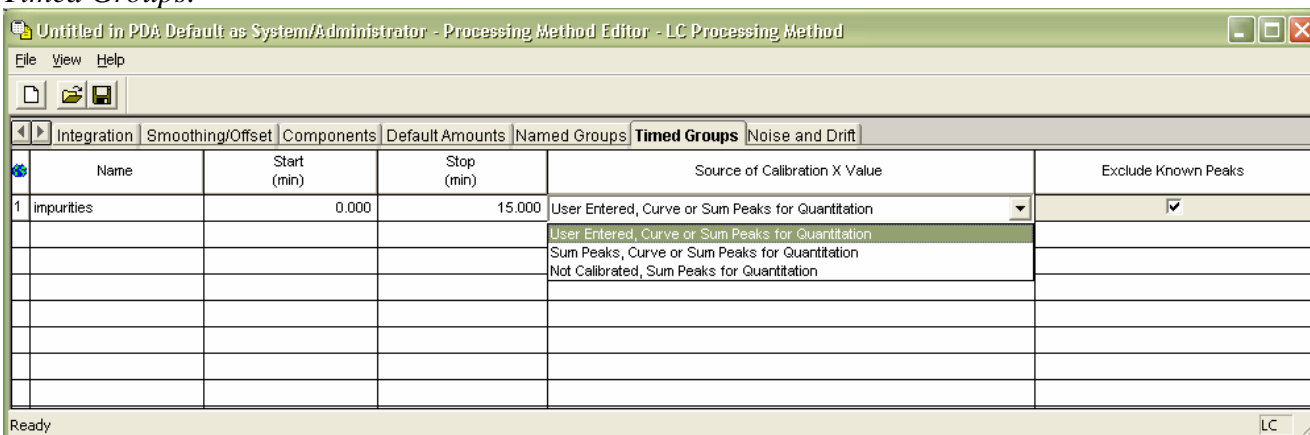
This is an optional way to enter the standards' amounts and levels.

Peaks can be grouped by time or by Peaks' names:

**Named Groups:**

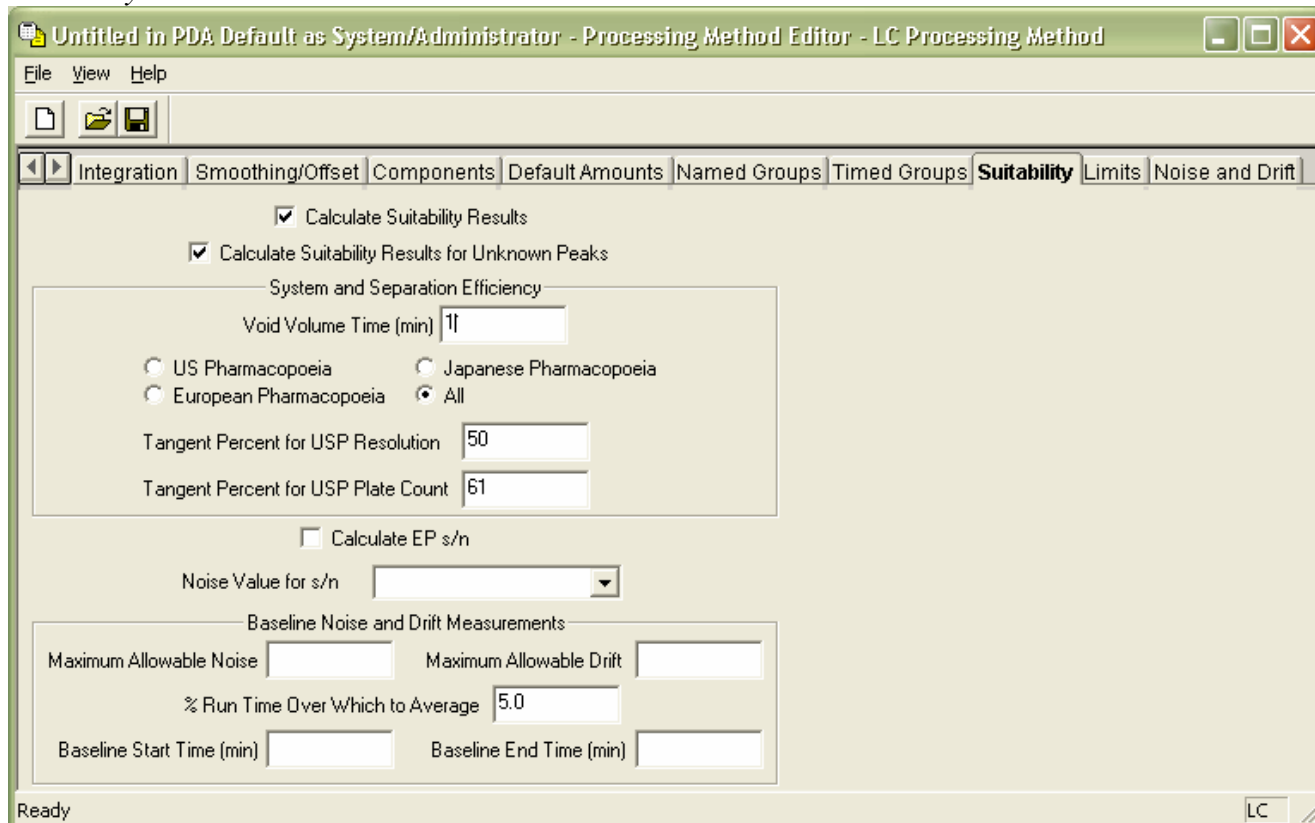


**Timed Groups:**



If you have **System Suitability** Option:

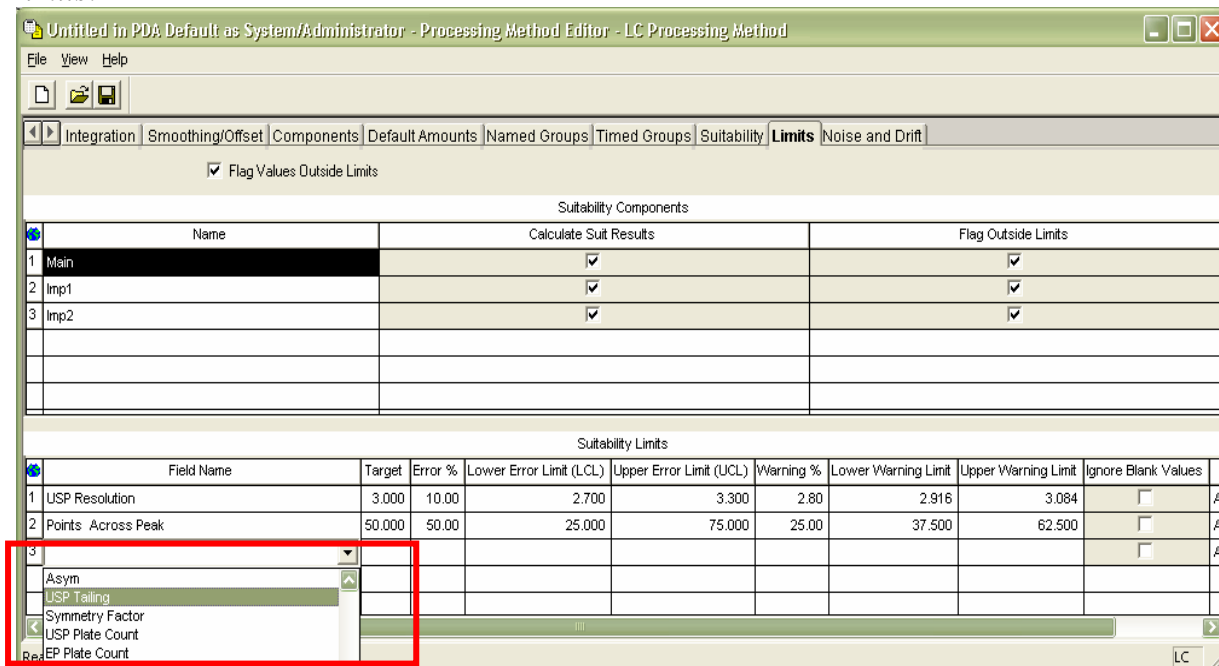
**Suitability:**

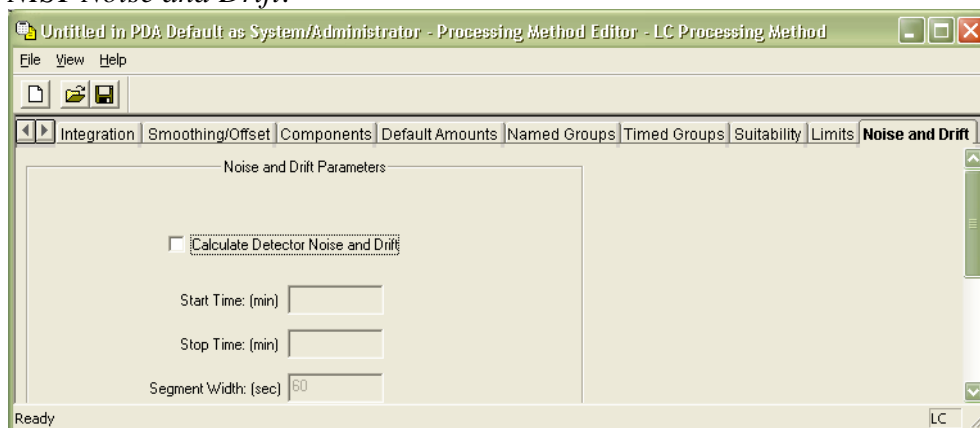


Baseline noise and drift are calculated here according to the pharmacopoeia. For more details: [http://www.forumsci.co.il/HPLC/SST\\_Emp2.pdf](http://www.forumsci.co.il/HPLC/SST_Emp2.pdf)

Setting limits for System Suitability: Select the component and fill the limits according to the protocols:

**Limits:**



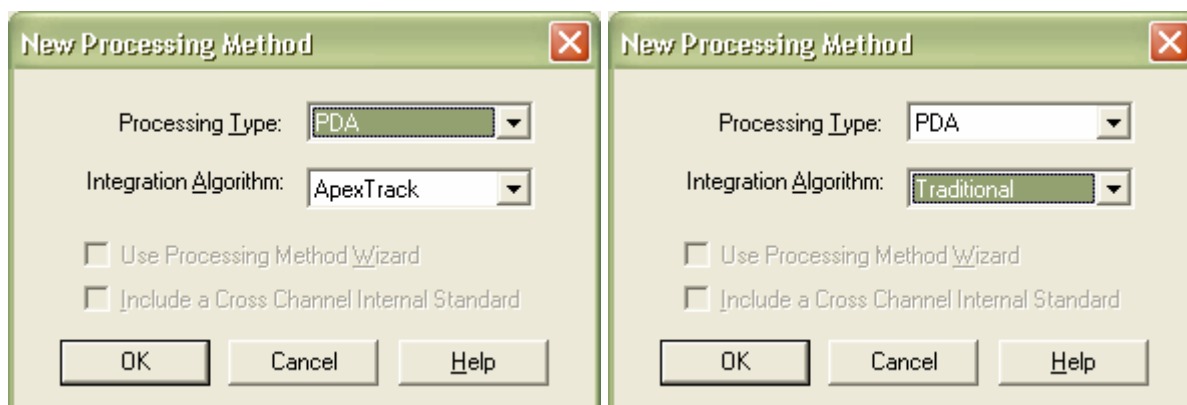
**NIST Noise and Drift:**

For more details on quantitative processing in Empower:

[http://www.forumsci.co.il/HPLC/Quantitative\\_Processing.pdf](http://www.forumsci.co.il/HPLC/Quantitative_Processing.pdf)

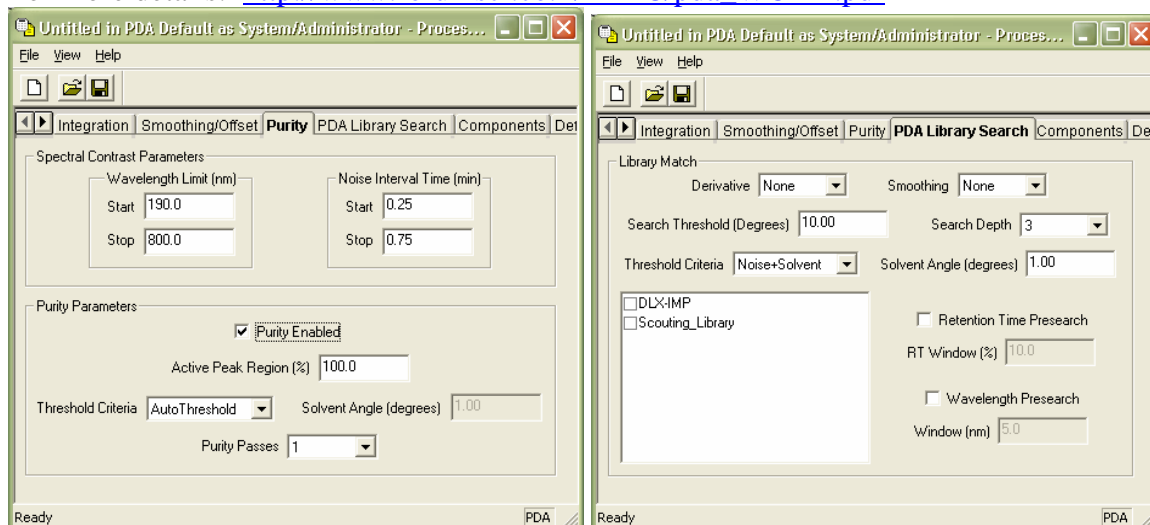
Acquisition and Processing Theory (register first):

<http://www.waters.com/webassets/cms/support/docs/71500031209rb.pdf>

**The PDA Processing Method:**

It contains 2 additional screens for measurement of Peak Purity and PDA Library Search of 3D data:

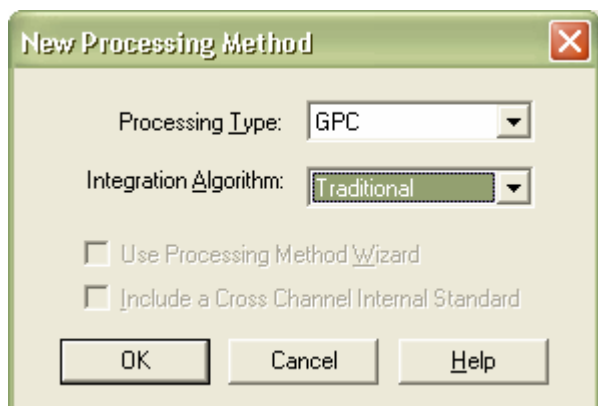
For more details: [http://www.forumsci.co.il/HPLC/pda\\_WORK.pdf](http://www.forumsci.co.il/HPLC/pda_WORK.pdf)



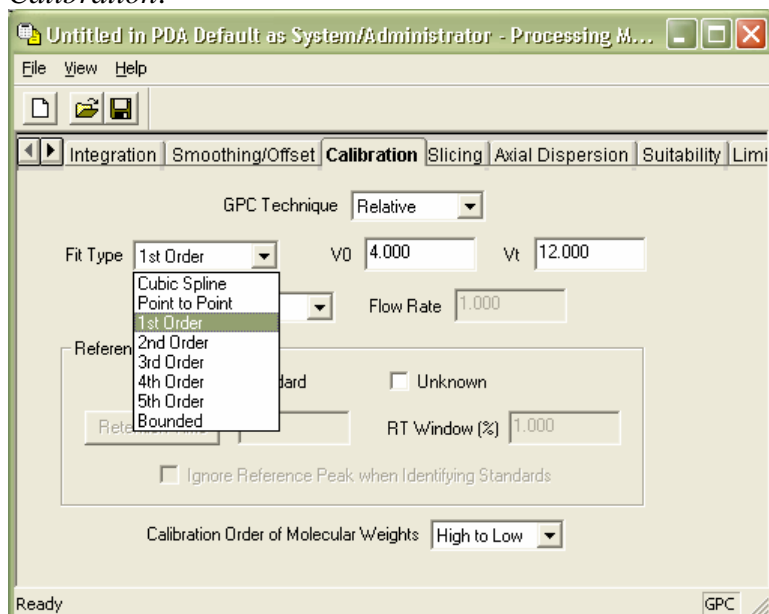
and getting started (register first):

<http://www.waters.com/webassets/cms/support/docs/71500031503ra.pdf>

The GPC (Gel Permeation Chromatography) Method:

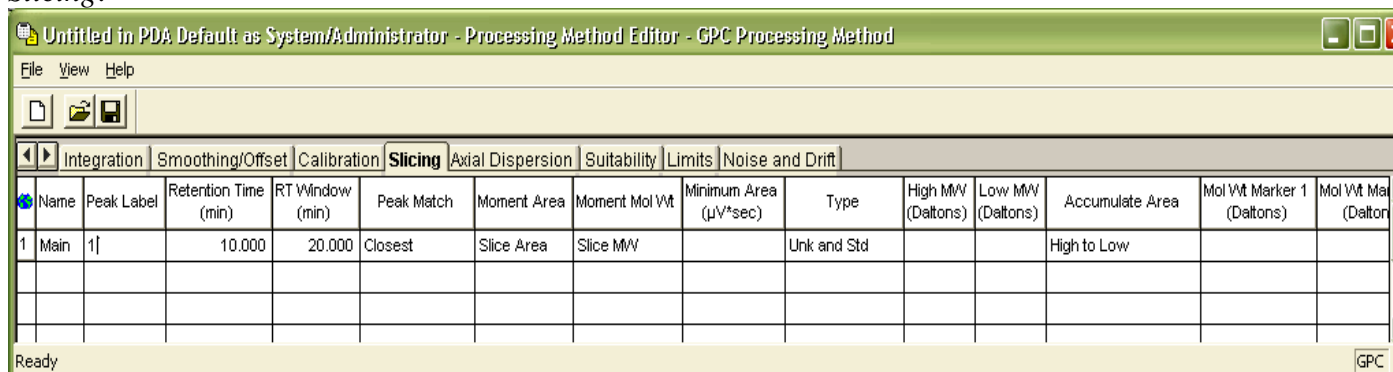


There are 3 additional screens specific to GPC work, *Calibration*, *Slicing* and *Axial Dispersion*:  
**Calibration:**

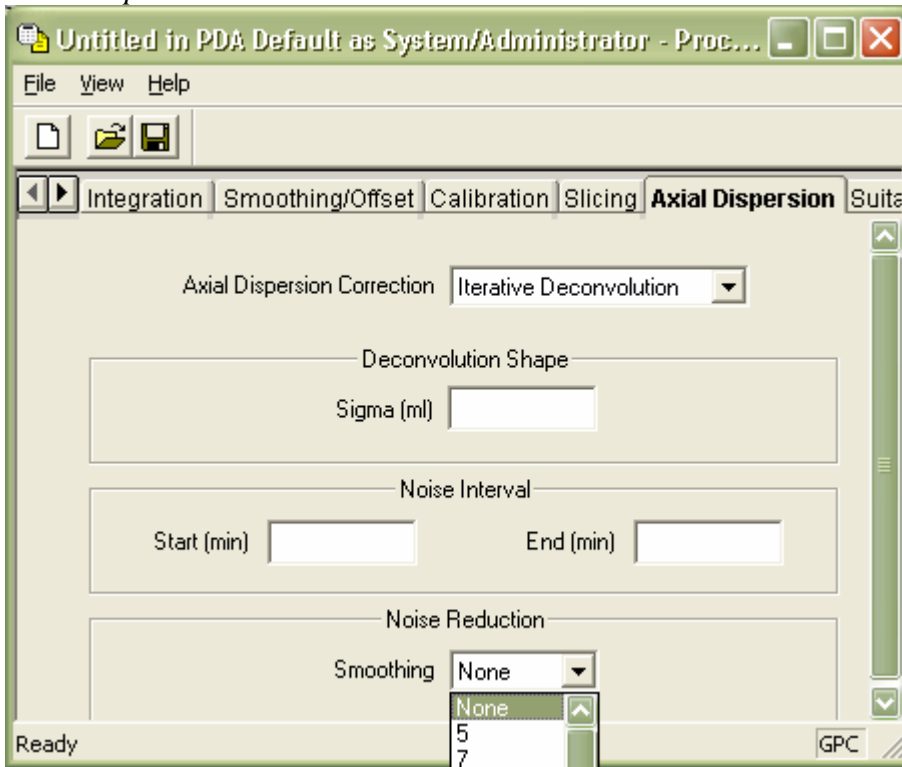


$V_0$  and  $V_t$  enable extrapolation outside the existing calibration curve, fit type above 1<sup>st</sup> order requires 5 points and above.

Where the Broad Unknown peaks need to be defined properly for the calculations of MW distribution:  
**Slicing:**

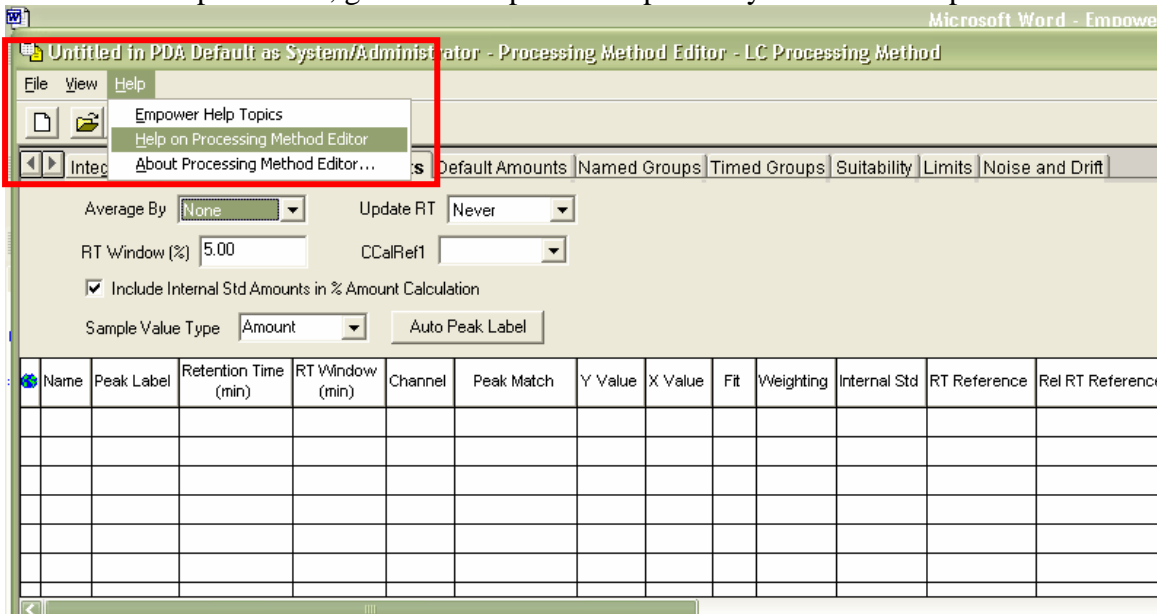


Deconvolution of the system's diffusional dispersion from the broad unknown peaks:  
*Axial Dispersion:*



For GPC processing: [http://www.forumsci.co.il/HPLC/gpc\\_steps.doc](http://www.forumsci.co.il/HPLC/gpc_steps.doc)  
 and <http://www.waters.com/webassets/cms/support/docs/71500031303ra.pdf>

For detailed explanations, go to the Empower Help while you are on the specific screen of question.



For Acquisition and Processing details:  
<http://www.waters.com/webassets/cms/support/docs/71500031209rb.pdf>