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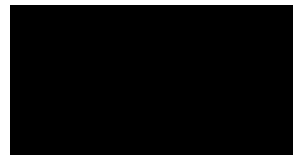
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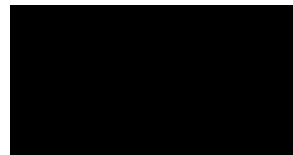


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For further improvement of the spectrum baseline,  **Baseline** leads to the interactive baseline correction. The dataset window will show icons specific for baseline correction.



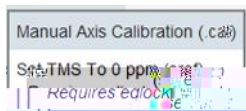
In addition, the Baseline pulldown menu gives access to further baseline correction operations.



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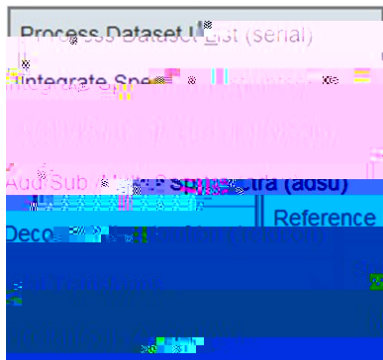
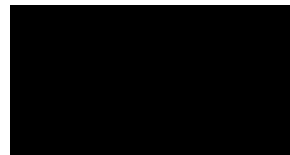
gives access to following pulldown menu with axis calibration options:




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gives access to further options through following pulldown menu:




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To enter the Analyse Process, click , the following workflow appears:



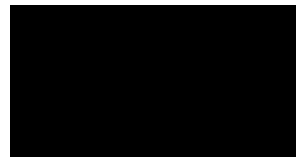
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 gives access to the interactive peak picking. The dataset window shows specific icons for peak picking:

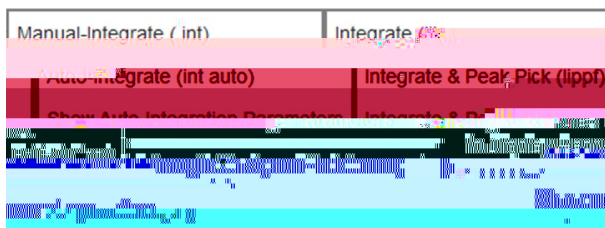


In addition, the Pick Peaks pulldown menu gives access to further peak picking operations and options:



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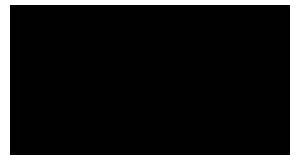
Integrate button gives access to the integration pulldown menu:



Integrate & Peak Pick (lippi)

In addition, more options can be obtained through the command ~ Ł :


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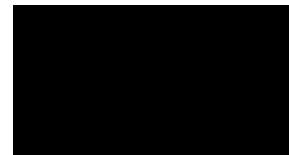


6Z" k" 0fl4 "fl ° " /LZi 5


Printing the spectrum can be done in two different ways:

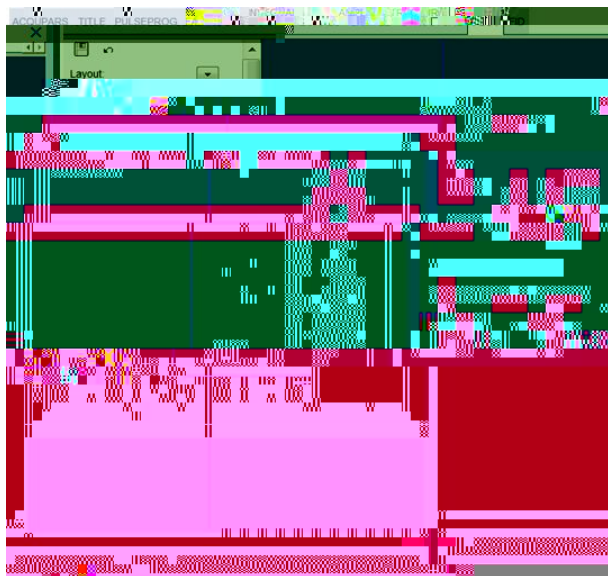
. '% 6Z" kfl ° /Z" " fl7Z" /L*3

Click on the printer icon  , located on the upper right part of the TopSpin window, allows direct printing of the spectrum. Details can be obtained as usual by dragging the mouse on the icon. Clicking the button gives access to the print window:



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To use this TopSpin feature, click the button , or use the ~*~k command in the TopSpin command line. The following TopSpin window appears:



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The left part of the plot window shows the names of the selected layout and printer. They can be changed with the arrow button, as well as other plotting view parameters, like limits or display. New NMR elements (title, parameters, 2D spectrum, etc.) can be inserted.

The right part presents the spectrum like it will be plotted. Changes on the left part will be shown in real time in the layout on the right.

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For detailed information please refer to the TopSpin Manual section under  = fl < t k i t * \$ fl > ! ~ / \$?



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Contact our Bruker BioSpin service centers.

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