

# Analysis V3<sub>beta2</sub> shortcuts

## Mouse events

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Context menu	Right click
Pan	Left drag
Zoom into area	Shift right drag
Zoom into area	Middle drag
Zoom in/out around cursor	Scroll wheel in display
Zoom in/out for selected axis	Scroll wheel in axis active area
Select new	Left click
Add/Remove one to/from Selection	Ctrl/Cmd left click
Select in area and add	Ctrl/Cmd left drag
Pick one peak at position	Ctrl/Cmd shift left click
Pick peaks in area	Ctrl/Cmd shift left drag
Move single peak	Ctrl/Cmd middle drag
Move module	Drag&drop purple bar
Move whole window	Alt Left drag
Pop out module (manage as windows)	Double click on Purple bar
Make new object	Double click <New> in Sidebar
View/Edit object	Double click object in Sidebar

## Keyboard shortcuts

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	Shortcut
<i>Graphics :</i>	
Delete selected	Del (Fn+Backspace)
Toggle Grid	GS
Toggle Crosshairs	CH
Toggle Horizontal Trace	TH
Toggle Vertical Trace	TV
Increase Trace Scale	TU
Decrease Trace Scale	TD
Create Mark at cursor position	MK
Clear All Marks	MC
Navigate to Peak position	FP
Navigate to NmrResidue position	FN
Copy PeakList	CL
Copy Peaks	CP
Clear Current Selected Peaks	CZ
Refit Selected Peaks	RP
Snap to Extremum Current Selected Peaks	SE
Select All Peaks	Ctrl/Cmd+A
Propagate Assignments	CA
Toggle Phasing Pivot	PV

Add Integral in 1D plot	I1
Add Phasing Trace	PT
Remove Phasing Trace	TR

*Project :*

New project	Ctrl/Cmd+N
Open project	Ctrl/Cmd+O
Load Data	LD
Save	Ctrl/Cmd+S
Save as	SA
Import Nef File	IN
Export Nef File	EX
Undo	Ctrl/Cmd+Z
Redo	Ctrl/Cmd+Y
Quit	Ctrl/Cmd+Q

*Spectrum :*

Load Spectrum	LS
Show Spectrum Groups	SS
Set Experiment Types	ET
Pick 1D Peaks	P1
Pick ND Peaks	PP
Copy PeakList	CL
Copy Peaks	CP
<i>Make Projection</i>	PJ
Print to File	PR

*Molecules :*

Reference Chemical Shifts	RC
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*Assign :*

Set up NmrResidues	SN
Pick and assign	PA
Backbone assignment	BB
Peak Assigner	AA
Assignment Inspector	AI
Residue information	RI

*View :*

New Blank Display	ND
NmrAtom Chemical Shift Table	CT
NmrResidue Table	NT
Peak Table	LT
Integral Table	IT
Restraint Table	RT

<b>Structure Table</b>	<b>ST</b>
<b>Sequence Graph</b>	<b>SG</b>
<b>Atom Selector</b>	<b>AS</b>
<b>Chemical Shift Mapping</b>	<b>CM</b>
<b>Show Sequence</b>	<b>SQ</b>
<b>Notes Table</b>	<b>NO</b>
<b>Python Console</b>	<b>PY</b>
<i>For current strip</i>	
<b>Show/Hide Toolbar</b>	<b>TB</b>
<b>Show/Hide Phasing Console</b>	<b>PC</b>
<b>Move to Next Spectrum (and Hide Current)</b>	<b>MN</b>
<b>Move to Previous Spectrum (and Hide Current)</b>	<b>MP</b>
<b>Reset Zoom</b>	<b>RZ</b>
<b>Store Zoom</b>	<b>ZS</b>
<b>Restore Zoom</b>	<b>ZR</b>
<b>Cycle through Peak Labelling Types</b>	<b>PL</b>
<b>Flip X-Y Axis</b>	<b>XY</b>
<b>Flip X-Z Axis</b>	<b>XZ</b>
<b>Flip Y-Z Axis</b>	<b>YZ</b>
 <i>Macro :</i>	
<b>Define User Shortcuts</b>	<b>DU</b>
<b>Reserved for user macros</b>	<b>Q0, Q1, ...Q9</b>
<b>Reserved for user macros</b>	<b>W0, W1, ...W9</b>
<b>Reserved for user macros</b>	<b>Y0, Y1, ...Y9</b>