

## V. VNMR to Bruker-AM/AC Parameter Conversion Table

[comments apply to VNMR unless specifically mentioned otherwise]

Parameter	AM/AC	VNMR	Comments
<b>Experiments</b>			
standard 1d $^1H$	ZG	go / s2pul	go zeros memory, and starts acquisition; seqfil='s2pul'
homonuclear decoupling	HOMODEC.AU	s2pul homo='y'	set the cursor on the peak and use <b>sd</b> to get the decoupler frequency
1d NOE difference	NOEDIFF.AU	s2pul homo='y' dm='yyn'	array can be used to run multiple decoupler frequencies in one exp.
standard 1d decoupled $X/^{13}C$	CPD ZG	s2pul dm='yyy'	
NOE-enhanced coupled $^{13}C$	GATEDEC.AU	s2pul dm='yyn'	DEPT is preferred unless you need quat's
quantitative decoupled $^{13}C$	INVGATE.AU	s2pul dm='nny'	X nucleus $T_1$ 's can be quite long, so this experiment can be arduous
DEPT $^{13}C$ editing	DEPT.AU	dept	mult=0.5; 1; 1.5 produces dept45; 90; 135 respectively
homonuclear correlation 2d	COSY.AU	cosy	gcosy is the better experiment; sw/(ni*2) gives usable digital resolution; usually need $\leq 6$ Hz/pt
long-range cosy	COSYLR.AU	cosy tau $\neq$ 0	gcosy is the better experiment
double quantum cosy	DQCOSY.AU	dqfcosy	DQCOSY; complete phase cycling is crucial for the dq filter; nt = multiple of 8
phase sensitive noesy	NOESYPH.AU	noesy	NOESY; flat baselines are important for observing small noe's; use <b>calfa</b>
phase sensitive roesy	ROESYPH.AU	roesy	ROESY; flat baselines are important for observing small noe's; use <b>calfa</b>
total correlation/WOHAHA	???	tocsy	TOCSY or TOCSY1D; useful for mixtures or separated spin systems
heteronuclear correlation	XHRCORR.AU	hetcor	use only if need very high $^{13}C$ resolution
inverse hetero correlation	???	hmqc	gHMBCAD preferred
multiple bond hetero correl.	???	hmqc bond $\neq$ 0	HSQCAD; important experiment for observing through linkage bonding

<b>Read and Save Commands</b>			
read data file	<b>RE</b> <i>filen.ame</i>	<b>rt</b> ('filename')	in VNMR, use also MAIN MENU FILE click on filename and LOAD
save data file	<b>WR</b> <i>filen.ame</i>	<b>svf</b> ('filename')	in VNMR, no menu selections for this
read parameter file	<b>RJ</b> <i>filen.ame</i>	<b>rtp</b> ('filename')	in VNMR, use also MAIN MENU SETUP NUC,SOLV
save parameter file	<b>WJ</b> <i>filen.ame</i>	<b>svp</b> ('filename')	
read shim file	<b>RSH</b> <i>filen.ame</i>	<b>rts</b> ('filename')	in VNMR, will search ~/vnmrsys/shims and /vnmr/shims paths
save shim file	<b>WSH</b> <i>filen.ame</i>	<b>svs</b> ('filename')	in VNMR, will save to ~/vnmrsys/shims
load shim file	none needed	<b>loadshims</b>	loadshims is UW-written macro having <b>load='y'</b> <b>su load='n'</b> <b>su</b>

<b>1d Acquisition Commands</b>			
tune <sup>1</sup> H observe	<b>RJ H1.SET</b>	<b>UWmacro</b> <b>TuneH1</b>	tuneh is a UW-written macro
tune <sup>13</sup> C channel	<b>RJ C13.SET</b>	<b>UWmacro</b> <b>TuneC13</b>	tunec is a UW-written macro
setup 1H for CDCl <sub>3</sub>	<b>RJ CDCL3.1DJ</b>	use menus	in VNMR, use MAIN MENU SETUP NUC,SOLV
zero and go	<b>ZG</b>	<b>go</b> or <b>ga</b>	<b>ga</b> will automatically apply a <b>wft</b> after acquisition
automation run	<b>AU</b> <i>autom.nam</i>	<b>au</b>	all VNMR programs run from compiled routines
halt acquisition with data	<b>^H</b>	<b>sa</b>	in VNMR, <b>svf</b> can be issued during acquisition to save data
resume acquisition	<b>GO</b>	<b>ra</b>	seems to work only for 1d (unless the 2d or 3d has been modified); vnmr's <b>ra</b> follows an <b>sa</b> that stops acq.
abort acquisition	<b>^E</b> or <b>^K</b>	<b>aa</b>	in VNMR, data is retained and ok
automation setup	<b>AS</b> <i>auton.ame</i>	none (try <b>dps</b> )	in VNMR, only parameters used in experiment will be shown
<b>1d Acquisition Parameters</b>			
sweep width	<b>SW</b>	<b>sw</b>	
center or offset frequency	<b>O1</b>	<b>tof</b>	
center decoupler freq	<b>O2</b>	<b>dof</b>	
solvent	none (change O1 thru jobfile)	<b>solvent='cdcl3'</b>	with solvent set correctly in VNMR, <b>tof=0</b> will center spectrum for normal organic compounds

set spectrum window	<b>EP</b> set window <b>^O</b>	set cursors <b>movesw</b>	changes <b>SW, O1</b> → <b>sw, tof</b> ; on Brukers, <b>AQ</b> will change, whereas in VNMR, <b>np</b> will change (leaving <b>at</b> unchanged)
set offset frequency	<b>EP</b> set cursor <b>O1</b>	set cursor <b>movetof</b>	changes <b>O1</b> → <b>tof</b>
relaxation delay	<b>RD</b> or <b>D1</b>	<b>d1</b>	AM/AC delay differs depending on ZG or AU to run experiment
common pulse width	<b>PW</b>	<b>pw</b>	90° length fixed by probe on AM/AC's; depends also on <b>pw90</b> and <b>tpwr</b> in VNMR
acquisition time	<b>AQ</b> (=TD*DW)	<b>at</b> (=np/sw)	
dwelt time	<b>DW</b> (=1/(2*SW))	= 1/sw	Bruker acquires complex pairs sequentially, vnmr simultaneously
# of transients to acquire	<b>NS</b>	<b>nt</b>	Bruker NS -1 which goes continuously → vnmr nt=1e6
# transients before acquis.	<b>DS</b>	<b>ss</b>	
receiver gain	<b>RG</b> (larger # → larger gain)	<b>gain</b> (larger # → larger gain)	on Unity, if gain=0 still clips (get ADC OVERFLOW message), insert attenuator at preamp output
observe transmitter power	none	<b>tpwr=52</b> (higher # → higher power)	AM/AC's observe power is fixed; Unity's have linear amplifiers on both observe and decouple
number of points acquired	<b>TD</b> (usually = SI)	<b>np</b>	
temperature	<b>TE</b>	<b>temp=24</b>	see <b>temp24</b> and similar macros (written at UW)
decoupler transmitter power	<b>DP&lt;ret&gt;20H</b> (lower # → higher power)	<b>dpwr=40</b> (lower # → lower power)	vnmr parameters are logical

### 1d Processing Commands and Parameters

fourier transform	<b>FT</b>	<b>ft</b>	
number of points FT'd	<b>SI</b>	<b>fn</b>	zero-filling occurs here (e.g., np=1024, fn=2048 will zero-fill once)
line broadening parameter	<b>LB</b>	<b>lb</b>	
interactively set weighting parameters	none	<b>wtia</b> or <b>wti</b>	in vnmr, middle button still control intensity in all windows; left button sets parameter
apply exponential line broad.	<b>EF</b>	<b>wft</b> or <b>dsx</b>	in vnmr, <b>wft</b> applies whatever weighting function is setup; <b>dsx</b> is a UW macro, and simply adds a <b>dscale(0)</b> after the FT

set reference	<b>EP</b> set cursor <b>G</b>	set cursor <b>nl</b> <b>rl(0p)</b>	vnmr gives example for TMS
automatic phasing	<b>AZPK??</b>	<b>aph</b>	in vnmr, <b>lp</b> should ~ 0, otherwise advise <b>calfa</b> command and/or back-linear prediction
baseline correction	<b>EP K</b>	<b>dc</b> or <b>bs(5)</b>	in vnmr, <b>bs</b> is <i>not</i> recommended for 1d (default spline fit), but is very good for 2d work-ups (recommended there)
normalized intensities	<b>AI&lt;ret&gt;0</b>	<b>nm</b>	in vnmr, vs=100 will fill screen
absolute intensities	<b>AI&lt;ret&gt;1</b>	<b>ai</b>	
<b>1d Plotting Commands</b>			
plot spectrum	<b>PX</b>	<b>pl</b>	
plot parameters	in <b>DPO</b> setup	<b>ppa</b> or <b>pap</b>	
plot axis	in <b>DPO</b> setup	<b>pscale</b>	in vnmr, axis='p' sets axis to ppm
plot coordinates	<b>X0, Y0</b>	<b>sc, vp</b>	sc is mm from right side, vp is mm vertically up from bottom
plot size	<b>CX, CY</b>	<b>wc, vs</b>	wc is mm width of chart, vs is vertical scale
plot integrals	<b>PXD???</b>	<b>pirn</b>	
plot peak picks	in <b>DPO</b> setup	<b>dll</b> or <b>ppf</b>	<b>printon dll printoff</b> prints table to separate page (do before any plotting commands); <b>ppf</b> plots on spectrum
new page	<b>NP</b>	<b>page</b>	

<b>2d Acquisition Parameters</b>			
sweep width for F1	<b>SW1</b>	<b>sw1</b>	AC/AM SW1 is 1/2 of observed sweep width; vnmr sw1 = observed sweep width
# increments/experiments	<b>IN</b>	<b>ni</b>	in vnmr, phase determines total # experiment = 1× or 2×ni
type of 2d acquisition	<b>MC2</b>	<b>phase</b>	AC/AM: only absolute value and TPPI are available in software vnmr: absolute value → phase=0 States-Habercorn → phase=1,2 TPPI → phase=3
2d setup	<b>ST2D</b>	none	in vnmr, type in sequence macro then <b>dps</b>
total time of experiment	none	<b>time</b>	
interleaved acquisition	depends on routine	<b>il='y'</b>	in vnmr, acquire <b>bs</b> scans per increment, loop until <b>nt</b> completed
<b>2d Processing Commands and Parameters</b>			
FT size in F1	<b>SI1</b>	<b>fn1</b>	AC/AM square requires SI1=SI/2=SI2/2
reference in F1	<b>SR1</b>	set cursor, <b>r11(0p)</b>	
FT and weight full set	<b>XFB</b>	<b>wft2da</b>	in vnmr, for absolute value sets use <b>do2d</b> or <b>wft2d</b>
FT and weight $t_2$ dimension	<b>XF1</b>	<b>wft1da</b>	counter-intuitive commands, but mean 1st transform
interactive weighting	none	<b>wtia</b>	can be done on $t_2$ fid, e.g., <b>wft(1) wtia</b> , and on $t_1$ fid, e.g., <b>wft1da TRACE wtia</b>
display color map	<b>EP2D</b> or <b>AP2D</b>	<b>dconi</b>	
<b>2d Plotting Commands</b>			
plot contours	<b>CPL</b>	<b>pcon</b>	in vnmr, see also <b>pconpos</b> and <b>pconneg</b> , UW written macros
plot size	<b>CX, CY</b>	<b>wc, wc2</b>	in vnmr, sc still controls distance in mm from right-hand side
Plot 2d using high res 1d		<b>plot2dhr</b>	have 1d high res already worked up in separate exp, follow prompts
peak picking and volumes	none	<b>ll2d</b>	