Bruker Pulse Program "Secret Decoder Ring"

;Pulprog.info

For a Bruker pulse program, the first ~4-6 characters usually specify the type of experiment being run e.g., DEPT. COSY, NOESY. Other information about the pulse program is indicated by a two-letter code, which is listed below in alphabetical order. In case of redundant information some two-character codes may be omitted.

2D NMR experiments always indicate the mode (absolute value, phase-sensitive, echoantiecho). By default, H-decoupling (for heteronuclear observe) or X-decoupling (for 1H observe) is assumed to be on. For homonuclear experiments, H-decoupling is not on by default.

Typical experiment names would be: cosy, dept, dipsi2, hmbc, hmqc, hoesy, hsqc, inad, inept, mlev, noesy, roesy or trosy.

Inverse correlations are denoted as hmbc, hmqc or hsqc. Experiments with a BIRD sequence in the beginning also contain a bi in the name.

1D experiments, which are analogues of 2D experiments by virtue of a selective pulse, start with sel. Semiselective 2D experiments have the same name as the unselective version but with an s at the beginning: scosyph <-> cosyph.

A phase-sensitive (States-TPPI, TPPI etc.) NOESY experiment with presaturation would then be: noesy + ph + pr = noesyphpr.

In the other direction the pulseprogram hmbcgplpndqf = hmbc + gp + lp + nd + qf and therefore an: inverse correlation for long-range couplings (HMBC) with coherence selection using gradients with "gp" syntax, a low-pass J-filter, and no decoupling.

The two-character codes used are the following:

- ac accordion type experiment
- ad using adiabatic spinlock
- ar experiment for aromatic residues
- at adiabatic TOCSY
- bi with bird pulse for homonuclear Jdecoupling
- bp using bipolar gradients
- cc cross correlation experiment

cn C13 and N15 dependent information in different indirect dimensions

- co with COSY transfer
- cp with composite pulse
- ct constant time
- cv convection compensated
- cw decoupling using cw command
- cx using CLEANEX_PM
- dc decoupling using cpd command
- df double quantum filter

di with DIPSI mixing sequence

dh homonuclear decoupling in indirect dimension

- dw decoupling using cpd command only during wet sequence
- dq double quantum coherence
- ea phase sensitive using
- Echo/Antiecho method
- ec with E.COSY transfer
- ed with multiplicity editing
- es excitation sculpting
- et phase sensitive using
- Echo/Antiecho-TPPI method
- fb using f2 and f3 channel
- fd using f1 and f3 channel (for presaturation)
- fr with presaturation using a frequency list

using f1 -, f2 - and f3 - channel (for ft presaturation) F-19 observe with H-1 decoupling fh using a flip-back pulse fp for F-19 ecoupler fl forward directed type experiment fw f2 using f2 - channel (for presaturation) using f3 - instead of f2 - channel f3 using f4 - instead of f2 - channel f4 gated decoupling using cpd gd command gradient echo experiment ge using gradients with ":gp" syntax gp using gradients gr using shaped gradients gs hydrogen bond experiment hb homodecoupling of a region using hc a cpd-sequence hd homodecoupling H-1 observe with F-19 decoupling hf with homospoil pulse hs InPhase-AntiPhase (IPAP) ia experiment IDIS - isotopically discriminated id spectroscopy inverse gated ig using inverse (invi/HSQC) sequence ii with incremented mixing time im with INEPT transfer in in phase ip using inverse (inv4/HMQC) i4 sequence for determination of J coupling ic constant id homonuclear J-decoupled with jump-return pulse jr jump symmetrized (roesy) js ld low power cpd decoupling with low-pass J-filter lp. with Q-switching (low Q) lq for long-range couplings lr with two-fold low-pass J-filter 12 with three-fold low-pass J-filter 13 mf multiple quantum filter with MLEV mixing sequence ml

mq using multiple quantum

nc N15 and C13 dependent information in different indirect dimensions

- nd no decoupling
- no with NOESY mixing sequence

pc with presaturation and composite pulse

- pg power-gated
- ph phase sensitive using States-
- TPPI, TPPI, States or QSEQ
- pl preparing a frequency list
- pn with presaturation using a 1D

NOESY sequence

- pp using purge pulses
- pr with presaturation

ps with presaturation using a shaped pulse

- qf absolute value mode
- qn for QNP-operation
- qs phase sensitive using qseq-mode
- rc for determination of residual dipolar couplings (RDC)/ J couplings
- rd refocussed
- re relaxation optimised (H-flip)
- rl with relay transfer
- ro with ROESY mixing sequence
- rs with radiation damping suppression using gradients
- rt real time
- ru using radiation damping
- compensation unit
- rv with random variation
- r2 with 2 step relay transfer
- r3 with 3 step relay transfer
- se spin echo experiment
- sh phase sensitive using States et al. method
- si sensitivity improved

sm simultaneous evolution of X and Y chemical shift

- sp using a shaped pulse
- sq using single quantum
- ss spin-state selective experiment
- st phase sensitive using States-TPPI method

- sy symmetric sequence
- s3 S3E experiment
- tc temperature compensation
- tf triple quantum filter
- tp phase sensitive using TPPI
- tr using TROSY sequence
- tz zeroquantum (ZQ) TROSY
- ul using a frequency list
- us updating shapes
- wg watergate using a soft-hard-soft sequence
- wt with WET watersuppression
- w5 watergate using W5 pulse
- xf x-filter experiments
- xy with XY CPMG sequence
- x1 x-filter in F1
- x2 x-filter in F2
- x3 x-filter in F3
- zf with z-filter
- zq zero quantum coherence

- zs using a gradient/rf spoil pulse
- 1d 1D version
- 1s using 1 spoil gradients
- 11 using 1-1 pulse
- 19 using 3-9-19 pulse
- 19f for F19
- 2h using 2H lockswitch unit
- 2s using 2 spoil gradients
- 3d 3D sequence

3n for E.COSY (3 spins, negative correlation)

3p for E.COSY (3 spins, positive correlation)

- 3s using 3 spoil gradients
- 30 using a 30 degree flip angle
- 45 using a 45 degree flip angle
- 90 using a 90 degree flip angle
- 135 using a 135 degree flip angle
- 180 using a 180 degree pulse