DPX 300 Selective 1D experiments. (SELNOESY and SELROESY does not work on DPX 300.) SELCOSY

- 1. edc define name user expno and so forth
- 2. rpar proton all
- 3. getprosol
- 4. rga
- 5. zg
- 6. efp, apk, abs
- 7. Note down the O1 value of peak to be irradiated
- 8. edc increase expon with 1
- 9. rpar uioselcosy all (do NOT type getprosol)
- 10. Enter the O1 value of the peak to irradiate
- 11. rga
- 12. ns and TD0 to be set (total number of scans = $NS \times TD0$) usually NS = 16.
- 13. zg
- 14. The experiment can be stopped by typing STOP (not HALT)
- 15. Do not type tr while doing the experiment
- 16. efp (NB efp antiphase correlation peaks)
- 17. Alternatively: FT, abs, PS, /8 several times (all peaks positive).
- 18.

SELTOCSY (better quality than selcosy)

- 1. edc define name user expno and so forth
- 2. rpar proton all
- 3. getprosol
- 4. rga
- 5. zg
- 6. efp, apk, abs
- 7. Note down the O1 value of peak to be irradiated
- 8. edc increase expno with 1
- 9. rpar uioseltocsy5 all (short range correlations) (do NOT type getprosol)
- 10. or rpar uioseltocsy80 all (medium range correlations)(do NOT type getprosol)
- 11. or rpar uioseltocsy240 all (long range correlations) (do NOT type getprosol) This the recommended one!
- 12. Enter the O1 value of the peak to irradiate
- 13. ns and TD0 to be set (total number of scans = $NS \times TD0$) usually NS = 16
- 14. rga
- 15. zg
- 16. The experiment can be stopped by typing STOP (not HALT)
- 17. Do not type tr while doing the experiment
- 18. For the 240 and 80 versions: efp (NB all peaks should be positive and phased as for a standard 1H spectrum)
- 19. For the 5 version

Alternatively: If some peaks show anti phase distortion (with shorter mixing times) power mode processing can be used to generate positive peaks: type: FT, PS, abs, /8 several times (all peaks positive)

- 19. To get an NMR spectrum with much higher intensity do as follows:
- 20. For resolution enhancement set LB = -1.5 (or -2), GB = 0.33
- 21. GFP
- 22. PS
- 23. abs
- 24. /8 several times (all peaks are positive with resolution enhancement processing).
- 25. LB to 0.1 (or other normal value)
- 26. GB = 0 BEFORE using EFP to reprocess the spectrum as a conventional 1H spectrum (otherwise EFP with GFP parameters gives rubbish/noise.