

List of common experiments on Ag500

Experiment entries	Description	Recommended setting for ~10mg material, please modify "NS" accordingly based on your sample concentration	
PROTON_icon	routine 1D proton spectrum	<p>Common parameters for all experiments:</p> <p>D1: relaxation delay, 1 to 5 times of T1, 2-5 seconds SW: spectral width in ppm for F2 (direct dimension) AQ: Acquisition time in seconds NS: number of scans O1P: offset freq for channels 1 in ppm, usually 1H O2P: offset for channel 2, usually 13C 1TD: Time domain data points for F1 (indirect dimension), aka number of increments 1SW: spectral width in ppm for F1</p>	
C13CPD_icon	1D ¹ H-decoupled ¹³ C spectrum		
C13DEPTQ135_icon	DEPTQ 135 experiment detect all Carbons - CH3/CH positive CH2/C negative		
C13IG_icon	13C with inverse gated 1H decoupling no NOE for quantitative NMR		
C13DEPT90_icon	DEPT 90 experiment only CH		
C13DEPT135_icon	DEPT 135 experiment CH3/CH positive CH2 negative - ¹³ C 1-bond correlations, all peaks positive (dept-45 analog) DEPT-135 experiment		
gCOSY_icon	Gradient selected COSY		
COSYDQF_icon	COSY with double quantum filter		
HSQC_EDIT_icon	1H-13C multiplicity edited HSQC with gradient selection		
HSQC_icon	1H-13C 1-bond correlations, all peaks positive, HSQC with gradient selection		
HSQC_EDIT_NUS_icon	1H-13C multiplicity edited HSQC with gradient selection Non Uniform Sampling w/ 25% sampling density		
HMBC_icon	1H-13C HMBC with gradient selection using 3-fold low pass filter for better 1J suppression		CNST13 = 3-12 Hz (default 8Hz for J _{2/3} CH)
HMBC_NUS_icon	1H-13C HMBC with gradient selection using 3-fold low pass filter for better 1J suppression Non Uniform Sampling w/ 50% sampling density		
TOCSY_icon	Phase sensitive 2D TOCSY experiment using MLEV-17 mixing	d9 = 30 to 120 ms	
NOESY_icon	Phase sensitive NOESY 1H-1H correlations based on proximity also for exchange	d8 = 0.1 to 1 second	
ROESY_icon	1H-1H correlations based on proximity for intermediate MW around 1600 Da	p15 = 0.1 to 0.5 second	
WATER_SUPP_icon	Solvent suppression with noesygprr1d sequence	p15 = 0.1 to 0.5 second	