## Texas A&M University Laboratory for Biological Mass Spectrometry

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## Sample Submission Form

– Contact Information ——					
Name:					
Phone:					
Email:					
Principal Investigator:		Address:			
- Sample Information —					
Sample ID:			Monoisotopic Mas	s:	
Attach sample vial within box.	Estimated Q		Solubility	Purity	
Provide chem. structure.	☐ In Solution Solvent _		☐ Methanol ☐ Acetonitrile	☐ Crude ☐ Semi Pure ☐ Pure	
	☐ Neat Liquid	l:ml mg	☐ Dichloromethane ☐ Water ☐ Tetrahydrofuran ☐ Other		
	(NOTE: Sai	(NOTE: Samples are NOT accepted in DMF or DMSO)			
	Confidence ☐ Tentative ☐ Confident ☐ Confirmed	Requir	rement ☐ Safe gerate ☐ Toxic Description  □ Biohazard	Sensitive to  ☐ Acid ☐ Base ☐ Air ☐ Light	
Please provide chemical structure	in the space provided abo	ove or attach drawing/im	age on the back of this for	<i>m</i>	
—Analysis Requested ———	Ionization	$\Box$ LC/MS			
Mass Analysis	<i>Tonization</i> □ + ESI	Provide the followi	☐ GC/MS ing info. Provide the	following info.	
☐ Unit Mass (Low-res)	□ - ESI	Column:	Temperature		
☐ Accurate Mass (Hi-Res)	□ + MALDI	ID:	· r	7 1 1 - 0	
☐ Fragmentation Pattern	□ - MALDI	Length:			
	□ + APCI	Flow rate:			
Mass Range Desired:	□ - APCI	Solvent A: Solvent B:			
Fromto	□ EI, CI □ Not Sure	Gradient:			
Acknowledgment Policy. Research acknowledgements of facility and st	carried out in part or in fu	ull using LBMS facilities		ibutions requires	
LBMS Facility Use Only Log in #			Operator:		
			-		
File Name:	Theoretic	al Mass:	Accuracy:	ppm	
Ionization Method: □ +ESI □ -ES Separation: □ LC □ GC Instrument: □ Qstar □ DECA □ S Matrix Used: □ DHB □ THAP □ C Time for Analysis:	Fragmentation: ☐ M FR ☐ DSQ Other CHCA ☐ SA ☐ HABA ☐ I	IS/MS □ Zip-Tip			
Comments:					
Billed Date:					