## **Laboratory for Biological Mass Spectrometry**

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General Information	on			http://mass-spec.tamu.edu		
C1				DI.		
Submitter				PI: Email of PI:		
e-mail						
Phone Number				Account #		
Sample ID				Department/Institution		
Type of Service						
Intact mass				Digestion with PMM		
				(MALDI)		
High Accuracy Protein				Digestion with PMM and		
Mass (sample must be salt				sequencing (MALDI)		
free, no Na, K, buffer,						
Top- down sequencing	5			MALDI-TOF peptide mass		
			]	mapping		
Protein Complex			1	MALDI-TOF/TOF peptide		
Trotein Complex				mass mapping with		
				sequencing		
PTM Analysis				LC-MS/MS (Protein		
1 Tivi / Midiysis				Identification)		
Other (describe):			<u> </u>	,	•	
I DMC DEDOMNEI		ECLMC		LC MS	4700 4800	
LBMS PERONNEL ONLY		ESI MS SynaptG2 QStar		SynaptG2 QStar	4700 4800	
ONLI		Synapioz QS	Stai	SyllapiO2 QStal		
Sample Informatio	10					
Sample Informatio		1,	0.37			
1. Are any buffers, det	_	=				
2. Was sample purifie	d? No	☐ Yes ☐ List	t	Solvent use	ed	
3. Solution concentrat	ion			Solvent is		
4. Radioactive? □	Irritant	? □ O	Other k	known hazards		
5. Chemical Formula:						
7. Additional Comme						