Mass spectrometry facility-Sample Submission Form
(Use separate form for each sample)
450 Chevron Science center, Department of Chemistry,
University of Pittsburgh, Pittsburgh, PA 15260

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File Number (SAMPLE LOCATION									
Name	Name Advisor									
Email										
Sample ID on Vi	al									
Account No:										
Room Number			bmitted							
Phone No.			/2024							
 Samples should be submitted as <u>neat oils or solids</u> only in 1 dram vials. Approximately 1 mg of the sample is sufficient. Label the vial on the <u>top ½</u> of the area to make the sample visible. 										
Analysis Needed O Nominal Mass /LRMS spectrum (Peaks in integer value) O Fragmentation pattern or MS/MS spectrum (For structural elucidation), provide the mass O Deconvoluted data Monoisotopic mass or Average mass O HRMS/Accurate mass for elemental composition determination (Error in Mass (mDa)/PPM). O Publication (Example: Error 3mDa for JOC, 5PPM for OL); Journal Name/requirement: O Dissertation O Other										
Preferred Ion Source (choose one) O Use appropriate ionization method needed O Electron Ionization: For volatile, low molecular weight compounds O Electrospray Ionization: For polar, non-volatile high m.w compounds O APCI/ASAP: For non-polar and low molecular weight compounds O MALDI: For high M.Wt compounds like Peptides, Proteins, Polymers etc. O LCMS O GCMS						Molecular Formula: Mono Isotopic Nominal Mass: Proposed Structure:				
Solubility (Mu O CH ₂ Cl ₂										
Sample State: O Solid O Liquid O Oil										
Special Handling: O Air Sensitive O Hazardous O Light Sensitive										
Purity checke	d by: O NMR O	TLC O GC O LC	0 0	ther						
Sample back: O Yes O No										
Below For Facility Use Only										
File Name										
Date										
Analyzed by										
5	1									

File Name			
Date			
Analyzed by			
Representative Masses			
Comments			