

**OKLAHOMA STATE UNIVERSITY**

*Proteomics / Mass Spectrometry Core Facility*

**Proteomics/Mass Spectrometry Work Order**

Name: \_\_\_\_\_ Department: \_\_\_\_\_  
 Email: \_\_\_\_\_ Campus account #: \_\_\_\_\_  
 Phone: \_\_\_\_\_ Principal Investigator: \_\_\_\_\_  
 Date: \_\_\_\_\_

**Sample Descriptions:**

sample/set identifier <sup>a</sup>	# of samples	analysis desired <sup>b</sup>	est. amount and volume <sup>c</sup>	purity, buffer, contaminants	other notes <sup>d</sup>

(Please use continuation pages as necessary.)

**Notes:**

- (a) Please note that your residual samples will be discarded after analysis, unless you make other arrangements.
- (b) "ID" = protein identification. "DE" = differential protein expression. All other analyses should be discussed with Dr. Hartson prior to sample preparation and submission.
- (c) E.g., "strong/medium/weak" x "Coomassie/silver" x "gel plug/gel band," or "est. solution conc."
- (d) Include species, and any other details or instructions that might increase the quality of our work.
- (e) Incomplete work orders will not be processed.

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**DO NOT WRITE IN SHADED AREA**

Analysis #:	_____	_____ 1 sample @ \$100 per sample _____	_____
		_____ sets of 10 @ \$700 per set	_____
Date Received:	_____	_____ hrs @ \$40 per hr	_____
Date Completed:	_____	_____ technical replicates @ \$10 each	_____
Operator:	_____	additional consumables :	_____
cc: file			
customer			
		<b>TOTAL DUE:</b>	_____