

Orders are final after submission. \* All Required Fields must be filled out before orders can be processed. Please complete the form and email to [support@appliedbiomics.com](mailto:support@appliedbiomics.com), or send the completed form with your samples.

<u><a href="#">ABO Internal Use</a></u>
PC: _____ PL: _____
Order: _____

Customer Information

<b>*Category</b>	<input type="checkbox"/> Same as previous project(s). Project Number _____ <input type="checkbox"/> Academic / Governmental Lab <input type="checkbox"/> Industrial / Commercial Lab	
<b>*Date</b>		
	<b>*Contact Person</b>	<b>*PI ( <input type="checkbox"/>Same as Contact Person)</b>
<b>*Name (First Last)</b>		
<b>*Company/Institution</b>		
<b>*Department</b>		
<b>*Phone number</b>		
<b>*Email Address</b>		
<b>*Mailing Address</b>		

**\* Billing Information (Check One and fill out all corresponding fields)**

- Purchase Order: Please send a hardcopy by email: [ar@appliedbiomics.com](mailto:ar@appliedbiomics.com)
- Credit card: Please provide the following information

<b>Contact person</b>	
<b>Phone number</b>	
<b>Email Address</b>	

How did you find us: Email   Search Engine   Conference   Referred by colleague \_\_\_\_\_

**\* Mass Spectrometry Services**
Number of Samples

<input type="checkbox"/> Protein ID by MALDI-TOF/TOF for 2D DIGE customers	
<input type="checkbox"/> Protein ID by MALDI-TOF/TOF for external customers	
<input type="checkbox"/> LC-MS/MS on Low complexity samples (1D Gel band)	
<input type="checkbox"/> LC-MS/MS on Moderate complexity samples	
<input type="checkbox"/> LC-MS/MS on Moderate to High complexity samples	

**Mass Spectrometry Order Form**

- LC-MS/MS on High complexity samples
- Phospho-peptide enrichment
- Identification of Phosphorylation-site \*
- \* Please provide accession number or protein sequence
- De Novo Peptide Sequencing
- Customized mass spectrometry services
- Customized sample preparation


**\* iTRAQ® Services**
**# of samples**

- iTRAQ® Experiment – 1
- iTRAQ® Experiment – 2
- Customized sample preparation

**\* iTRAQ® Services**
**# of samples**

- iTRAQ® Experiment – 3
- iTRAQ® Experiment – 4

Sample description for each experiment:**\*Biohazardous Material (we will not process samples if the following information is not provided)**

- The samples do NOT contain any biohazardous material or radioisotopes (e.g. <sup>14</sup>C, <sup>32</sup>P, <sup>35</sup>S, etc.) of any kind.
- The samples contain \_\_\_\_\_, which is a Level \_\_\_\_ Biohazardous material. I have completely deactivated the biohazardous material using the appropriate procedure.

**\*Sample information (required)**

Sample species:  Human  Mouse  Plant  Bacteria  Others \_\_\_\_\_

Gel slices:  Coomassie  Silver  Sypro ruby  Other \_\_\_\_\_

In-solution sample: buffer and estimated protein amount or concentration (mg/mL) \_\_\_\_\_

Comments:

\* Order Submitted and Agreed to by : \_\_\_\_\_

**Print Name and Title**

\_\_\_\_\_  
**Signature**

By signing this form, I certify that I have read the "Terms and Conditions" at <http://www.appliedbiomics.com/terms.html>, and will fully comply with those terms as a condition to the services provided by Applied Biomics.