LSI Mass Spec Sample Submission Form

Lab Information:

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Name: |  | Lab number: |  | PI Signature: |  |
| Date: |  | Tel: |  | E-mail: |  |

Sample Information:

|  |  |
| --- | --- |
| Sample Numbers： | Species: |
| Sample Names | |

Measurement Required

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Sample Type: | Protein Gel□ | protein solution□ | | cell□ | tissue□ | | | others□ |
|  |  | Buffer composition: | | | | | | |
|  |  | |  | | |  |  |  |
| Measurement Required： | Single protein identification:□ | | ID：\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ | | | | | |
|  | Map Protein Modification：□ | | □Phosphorylation □Acetylation  □Ubiquitination □Methylation  □Lactylation  □others\_\_\_\_\_\_\_\_\_\_ | | | | | |
|  | Protein mixture identification□ | |  | | | | | |
|  | Binding Protein□ | | ID：\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ | | | | | |
|  | Quantitative Proteomics□ | |  | | | | | |
|  | Quantitative Phosphomics□ | |  | | | | | |
| Other Experiments: | | | | | | | | |
| Please contact us for discussion. | | | | | | | | |