**Sample Information - Proteomics**

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| **Customer information** | |
| **Name:** | **Date of submission:** |
| **Phone/Email:** | **Department/Institute/Company:** |
| **Principal investigator (PI):** | **PI email:** |
| **Address:** | |

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| **Sample information** |

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| **Proteomics Services:**  \*Please attach target protein list separately if necessary | Protein identification  Protein identification + label-free quantification  Protein identification + post-translation modifications  PTM type(s):  Quantitative proteomics:  iTRAQ  SWATH  Others:  Targeted proteomics (SRM/MRM):  Relative quantification  Absolute quantification + stable isotope standards  Target protein(s)\*:  Custom analysis  Collaborative project |
| **Experiment requirements,**  **sample species and origin**  (*e.g.* TP53 pulldown in human cell line K562)  Please provide us with any customised protein sequences if applicable. |  |
| **Sample type** | Purified protein  Protein mixture in buffer  Pure peptide  Peptide mixture in buffer  Full gel  Excised gel bands  Others: |
| **Sample buffer composition**  (*e.g.* salts, concentration, pH, etc.) |  |
| **Do you wish to collect the Sample after analysis/testing?**  No  Yes | **Is the Sample subject to any export control / IRB regulations?**  No  Yes, Details: |
| **Are there any safety considerations / Does the Sample contain any hazardous properties?**  No  Yes, Details: | |

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| **Sample details\*** | | | | |
| **Sample number** | **Sample ID** | **Sample volume**  **(μL)** | **Protein concentration**  **(μg/μL)** | **Remarks**  **(if any)** |
| **01** |  |  |  |  |
| **02** |  |  |  |  |
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| **14** |  |  |  |  |
| **15** |  |  |  |  |

\*Please use a separate sheet if you have more samples

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