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| MasSpecLab**Mass Spectrometry Facility** | C:\Users\Stan\AppData\Local\Temp\gabarit-uvsq-ufr-sc-sante-s-veil-9x3x150-rvb.jpg |
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| **Managers: Jean-Claude Alvarez – Stanislas Grassin Delyle****Contact:**Stanislas Grassin DelylePlateforme de Spectrométrie de MasseUFR Sciences de la Santé Simone Veil2, avenue de la source de la Bièvre78180 Montigny le Bretonneux, FranceTel : +33 (0)1 70 42 94 22stanislas.grassin-delyle@uvsq.fr |
| Project Identification |
| Project reference : | Internal reference |
| Applicant laboratory: | Click here to enter text. | Laboratory Head: | Click here to enter text. |
| Project Manager: |
|  Last name: Click here to enter text. First name: Click here to enter text. | Phone number: Click here to enter text.Email address: Click here to enter text. |
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| Nature of the project: | [ ]  Elementary analysis[ ]  Drug analysis[ ]  Proteomics[ ]  Other: Click here to enter text. | Clinical trial: [ ]  Yes [ ]  NoClinicaltrials.gov or EudraCT ref: Enter reference. |
| Type of service desired: | [ ]  Scientific collaboration[ ]  Service delivery |
| Project Information |
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| Overall objective of the research: | Please specify here the goal of your research project. |
| Detailed objective of the analyzes requested: | Please specify here the results you expect for the analyzes requested |
| Project funding: | [ ]  Public : [ ]  Laboratory’ own funds[ ]  Grant. Reference: Enter the reference[ ]  Private |
| Provisional date for submission of samples: Click here to enter a date.Desired date for reporting results (indicative only): Click here to enter a date. |
| Samples |
| * Matrices:
 | [ ]  Blood[ ]  Plasma or serum : Enter here the nature of the anticoagulant[ ]  Cell culture medium: Describe the composition of the medium | [ ]  Urine[ ]  CSF[ ]  Other: Nature and composition |
| * Origin/species of matrices:
 | Species |
| * Total number of samples:
 | Number of samples |
| * Number of biological groups:
 | Number of groups |
| * Presence of organic solvents:
 | [ ]  No [ ]  Yes: Nature. |
| * Samples storage:
 | Choose an item. |
| * Samples pH (if known):
 | Samples pH. |
| * Presence of pH buffer:
 | [ ]  No [ ]  Yes: Nature. |
| * Stability of samples:
 | [ ]  Sensitive to light[ ]  Sensitive to heat: Critical temperature[ ]  Sensitive to pH change: Specify |
| * Potential toxicity of the sample to the handler:
 | [ ]  No [ ]  Yes: Nature. |
| * Other notes:
 | Specify any other relevant information. |
| Elements or molecules to be studied(Join Excel spreadsheet if more molecules) |
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| 1. Total number of elements or molecules to analyze: Number of analytes.
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| 1. For elementary analysis (ICP-MS):
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| * + List of elements to be determined:

 List of the elements of the periodic table. Specify the isotopes of interest. |
| 1. For analysis of drugs, toxics, proteomics or others:
 |
| Molecule 1 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
|  |
| Molecule 2 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Molecule 3 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Molecule 4 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Molecule 5 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Molecule 6 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Molecule 7 | Name: Molecule name.Range of expected concentrations in the different matrices: Specify the extremes of expected concentrations.Presence of other similar molecules in the matrix: [ ]  No [ ]  Yes: Nature.For experimental molecules:Structure or PubChem link: Structure, PubChem link. |
|  | Monoisotopic mass: MM. | Log P: Log P. | pKa: pKa. |
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| Analysis report |
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| [ ]  Minimal (only qualitative and/or quantitative results for each sample)[ ]  Standard (+ description of the methodology used)[ ]  Expanded (+ individual chromatograms and results of quality controls)[ ]  Integral (+ results of method validation) | **Language:**[ ]  French[ ]  English |
| List of samples provided(Join Excel spreadsheet if more samples) |
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| Sample Name | Patient / Animal / Tissue ID | Nature(Blood, urine…) | Volume (mL) | Biological condition tested | Specific information |
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[ ]  I declare to have read and accepted the policy for use of the facility **MasSpecLab**

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| Name of applicant: Last Name.Date: Click here to enter a date. | Signature: Handwritten signature or paste image below. |