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| **The MRC Weatherall Institute of Molecular Medicine**  **Advanced Single Cell Omics Facility (WASCOF)**  **Project Submission Form**  **Project Code:** |
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| **Date** |  |
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| **Project Name** |  |
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| **Investigator Name** |  |
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| **Investigator email** |  |
| **Lab Details (PI name)** |  |
| **Address (if not WIMM)** |  |
| **LAB to charge to (email, address)** |  |
| **Describe the service you need in details** |  |
| **Please, add the link of the protocol you would like to be used (e.g. 10x 3’)** |  |
| **Chemistry (e.g. 10x\_3’ GEX and CITEseq, Smart-seq3, Target-seq)** |  |
| **Technology/Platform to use (e.g. Chromium X; BD Rhapsody; Tape Station)** |  |
| **No. of samples (run) planned for the project**  **(Describe how you will submit the cells)** |  |
| **Descive nature of Samples: Cells/Nuclei Type & Size** |  |
| **Are you hashing your samples. If yes which chemistry has been used?** |  |
| **Cell Numbers/ Viability requested by the facility (YES/NO)**  **If No, please report Cells/Nuclei and viability to the facility before the loading stage** |  |
| **Cell Recovery Number Requested**  **(after GEMs generation, e.g. 5000/6000 per samples)** |  |
| **Expected Date for Sample Submission** |  |
| **Reagent supplied YES/NO (describe what, amount and when)** |  |
| **Type of Libraries Requested** |  |
| **QC Library requested YES/NO** |  |
| **Any special run requirements/machine type** |  |

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| **Project Summary** | |
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| ***Indicative Costing (WIMM Use):*** | |
| **Signed (researcher/PI):** | |
| **Project Code (WIMM Use):** | |
| **PO number (Researcher/PI use):** | |
| **Samples Submitted (date):** |  |
| **Samples Processed (date):** |  |
| **Result (WIMM Use) –**  **key results, QC and traces, concentration and index details will be uploaded at the end of the run/project.** | |