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| **To be completed by the Researcher** |
| **Project title1** |  |
| **Reference number/network2** |  |
| **ApplicantInstitutionContact (email; phone)** |  |
| **Research field** |  |
| **Project summary3** |  |
| **NGS application(s)4** |  |
| **Type & number of samples5** |  |
| **Sample delivery dates(in batches of)6** |  |
| **Data analysis support required7** |  |

1 Please provide the title of your research project (e.g. if available, the title of your DFG grant application)

2 Please provide a reference number for your project (e.g. DFG XY1234) if available and add any affiliations with research networks or collaborative grants (SFB, FOR initiatives, etc.)

3 Please provide a concise summary of the project

4 Please list all NGS applications that will be required for the project (e.g. bulkRNASeq, ATAC, scRNA)

5 Please provide information about the type (cells, or processes material like chromatin, DNA, RNA) and the number of sample you wish to analyse

6 Please indicate, when your samples are delivered, and in which batches

7 Please indicate, which level of bioinformatics support will be required.

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| **To be completed by the NGS Core** |
| **Project ID (internal)** |  |
| **Library preparation protocols** |  |
| **Limitations of sample input**  |  |
| **Sequencer** |  |
| **Data output per sample** |  |
| **Study design (controls/replicates)** |  |
| **Comments on design** |  |
| **Comments on experiments (e.g. pilot needed)** |  |
| **Costs of library preparation** |  |
| **Costs of sequencing** |  |
| **Comments on Data Analysis** |  |

Date of Consultation Signature Applicant Signature NGS Core