**Analysis order sheet Single-Cell Center Würzburg**

|  |  |
| --- | --- |
| Contact Person |  |
| Project Leader |  |
| Project Title |  |
| Institute |  |
| Street |  |
| Postal Code, City |  |
| Telefone |  |
| Email |  |

Number of samples:

Sample type:

Number of cells/samples:

Type of multiplexing:

Organism:

**Planned Investigation**

|  |  |
| --- | --- |
| [ ]  10x Genomics 3‘ Gene Expression | [ ]  10x Genomics 5‘ Gene Expression |
| [ ]  10x Genomics TCR V(D)J Enrichment | [ ]  10x Genomics BCR V(D)J Enrichment |
| [ ]  10x Genomics ATAC-Seq | [ ]  other application:       |
| [ ]  Sequencing only  | [ ]  Libraries only |

**Additional experimental details**

Further details regarding sample preparation (sorting, frozen or fresh etc):

TotalSeq antibodies/hashtags will be used (yes/no), number and type (A, B or C):

CiteSeq will be used (yes / no):

Requested sequencing depth per cell:

Comments: