

| For ATGC Facility use only: |  |
|-----------------------------|--|
| Project ID:                 |  |
| Submission Date:            |  |
| Time:                       |  |

## Advanced Technology Genomics Core

### 10X Genomics Single Cell Service Request Form

Instructions: Fill out both pages of the service request form. Please submit information about your samples in the Sample Information table below. Please be sure to include an active account and authorized signature from your Financial Administrator or Designee and send an e-copy of the completed form to [dpollock@mdanderson.org](mailto:dpollock@mdanderson.org). For additional questions on filling out this form, please contact David Pollock at the above-mentioned email or via telephone 713-834-6199.

#### Contact Information

|                                 |  |
|---------------------------------|--|
| <b>PI Name:</b>                 |  |
| <b>User Name:</b>               |  |
| <b>User Email:</b>              |  |
| <b>Immediate Contact Phone#</b> |  |
| <b>Department:</b>              |  |
| <b>Project Title:</b>           |  |

#### Sample Information

| Sample Name  | Cell and Tissue Type<br>(cell line, tumor cells,<br>tissue type, etc...) | # Cells to<br>capture<br>per sample<br>suspension | Genome<br>(Human,<br>Mouse, Rat,<br>etc..) | Volume<br>(uL) | Cell Density<br>(cells/uL) | Media<br>Type |
|--|--|---|--|----------------|----------------------------|---------------|
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
|  |  |   |  |                |                            |               |
| <b>Additional information<br/>regarding project:</b> |  |   |  |                |                            |               |

| Sequencing Application |  |
|------------------------|--|
|                        | Single Cell RNA: 3' RNA                      |
|                        | Single Cell RNA: 5' GE only                  |
|                        | Single Cell RNA: 5' GE+ TCR Enrichment       |
|                        | Single Cell RNA: 5' GE+ BCR (Ig) Enrichment  |
|                        | Single Cell RNA: 5' GE+ TCR + BCR Enrichment |
|                        | Single Cell ATACseq                          |
|                        | Single Cell Multiome (ATAC + 3' scRNA)       |

| Sequencing Instrument |                        |
|-----------------------|------------------------|
|                       | NextSeq500 High Output |
|                       | NextSeq500 Mid Output  |
|                       | NovaSeq6000 SP         |
|                       | NovaSeq6000 S1         |
|                       | NovaSeq6000 S2         |
|                       | NovaSeq6000 S4         |
|                       |                        |

### Billing Information

| Dept ID<br>(6 digits) | Fund Group<br>(2 digits) | Fund<br>(6 digits) | Fund Type<br>(2 digits) | *PCBU<br>(5 letters) | *Project<br>(6 digits) | *Activity<br>(4 digits) | Account<br>Expiration<br>Date | Percent<br>Distribution<br>(%) |
|-----------------------|--------------------------|--------------------|-------------------------|----------------------|------------------------|-------------------------|-------------------------------|--------------------------------|
|                       |                          |                    |                         |                      |                        |                         |                               |                                |
|                       |                          |                    |                         |                      |                        |                         |                               |                                |
|                       |                          |                    |                         |                      |                        |                         |                               |                                |

### Billing Authorization (usually not PI signature)

|  |  |
|--|--|
| <b>Financial Administrator (or Designee) Authorized Signature:</b>   |  |
| <b>Financial Administrator (or Designee) Printed Name and Title:</b> |  |

### Bioinformatics Analysis

|   |  |
|---|--|
| <b>Bioinformatician Name:</b>           |  |
| <b>Bioinformatician Email:</b>          |  |
| <b>Bioinformatician Phone:</b>          |  |
| <b>Project Description Design/Goal:</b> |  |

### Data Output

ATGC will provide sequencing data in bcl, fastq, QC metrics html and analysis file formats. 10X Genomics Sequencing software (specific to application- e.g. Cell Ranger) demultiplexes the samples and provides QC metrics output and raw data files. Raw data can be viewed using 10X Genomics desktop analysis software (specific to application-e.g. Loupe Cell Browser). The software can be downloaded from 10X Genomics website and is freeware. Please be sure to check computer requirements before downloading software.