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***New User or New Project Introductory Questionnaire***

***\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_***

Thank you for your interest in our core. For new users or new project requests, we ask that you please create a Power Point slide or two addressing the following questions and send the attachment to SingleCell@bwh.harvard.edu:

**Customer Information**

* **User/Principle Investigator Name**: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
* **Goal of Project**: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
* **Funding**: [ ]  Internal MGB Peoplesoft Fund [ ] External non-MGB Source

**Project Information**

1. Please state basic experimental design, questions, and project goals\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
2. What is the species of origin & the tissue type you wish to submit? \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
3. How many different condition(s) are you interested in comparing? \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
4. When were your samples collected and fixed?
5. Were the samples fixed and the FFPE blocks made under [standard protocol](https://kb.10xgenomics.com/hc/en-us/articles/20341795998349-Do-you-have-recommendations-for-FFPE-tissue-fixation-for-Xenium)? Please list reagents used for tissue fixation here: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
6. Have you done H&E staining or any other staining for the blocks? (*Note: We require H&E staining or other stains for all our spatial services.)* \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
7. Which spatial platform are you interested in? Are you aware of the specific gene panels of the platform of your interest?

[ ]  Visium [ ]  Visium HD [ ]  Xenium [ ]  Xenium 5K

1. If using Xenium or Xenium 5K, do you want to use any customized panel? If so, please provide the information to us. \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_
2. How many slides do you plan to submit (*Note: must be a multiple of 2*)? Please refer to our [Slide Preparation Guidelines](https://ccp.bwh.harvard.edu/spatial-transcriptomics-services/) for each platform.
	1. Visium: \_\_\_\_\_\_\_\_\_

*Samples from separate FFPE blocks can be cut and placed onto a singular slide, each slide allows for an 6.5mm x 6.5mm or 11mm x 11mm imageable area.*

* 1. Visium HD: \_\_\_\_\_\_\_\_\_

*Samples from separate FFPE blocks can be cut and placed onto a singular slide, each slide allows for an 6.5mm x 6.5mm imageable area.*

* 1. Xenium (*Note: Xenium slides must be requested from us prior to sectioning*): \_\_\_\_\_\_\_\_\_

*Samples from separate FFPE blocks can be cut and placed onto a singular slide, each slide allows for an 12mm x 24mm imageable area.*

* 1. Xenium 5K (*Note: Xenium slides must be requested from us prior to sectioning*): \_\_\_\_\_\_\_\_\_

*Samples from separate FFPE blocks can be cut and placed onto a singular slide, each slide allows for an 12mm x 24mm imageable area.*

1. What is your bioinformatic capacity? \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_