The Transcriptomics & Deep Sequencing Core

at the

2020 Single Cell Symposium

Monday, April 20, 2020

8:30am - 5:15pm (Registration & Refreshments begin at 8am)

Tilghman Auditorium, Miller Research Building

Keynote Speaker:
Single Cell Sequencing is the latest advancement in Next Generation Sequencing technology, allowing insight into gene expression, chromatin accessibility, antigen/cell surface protein, and CNV at the cellular level. Single Cell research furthers our understanding of how individual cell types function within the larger population. This information is often masked in more traditional bulk sequencing, but can be vital in understanding developmental process, disease progression and various other research fields.

Researcher & Vendor presentations will highlight the most recent advances in Single Cell analysis and provide understanding of current research using these state-of-the-art technologies. Single Cell processing is available through the Transcriptomics & Deep Sequencing Core to ensure all researchers have access to the platforms provided by our vendors.
Space is limited for this symposium (120). Pre-registration is required. Lunch will be provided.

To register:

Please contact Linda Orzolek (ldorsch2@jhmi.edu) in the Transcriptomics & Deep Sequencing Core with any questions.