

## Single Cell Seminar

Splice isoforms, gene fusions, and SNVs are not usually found at the 5' or 3' ends of transcripts and can be missed if you use a droplet-based approach for your single-cell NGS studies. See how SMART-Seq<sup>®</sup> technologies can help you uncover novel biomarkers from entire, full-length transcripts.



A complete, end-to-end automation solution with superior sensitivity to detect critical biomarkers such as splice variants, gene fusions, and SNVs.

**Emory Integrated Core Facilities** 

**Bill Warble** Senior Business Development Manager, Automation



An easy, plate-based solution for full-length single-cell sequencing that minimizes sample loss and handling errors.

Kunj Pathak, PhD Territory Manager II

**Date and time:** April 20, 2023 11:00 am – 12:00 pm EST

**Location:** Emory University, HSRB Auditorium, 1760 Haygood Dr. NE, Atlanta, GA 30322 Sponsored by: Emory Integrated Genomics Core



Unable to make it in person? Join us on Zoom: https://zoom.us/j/294261547



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