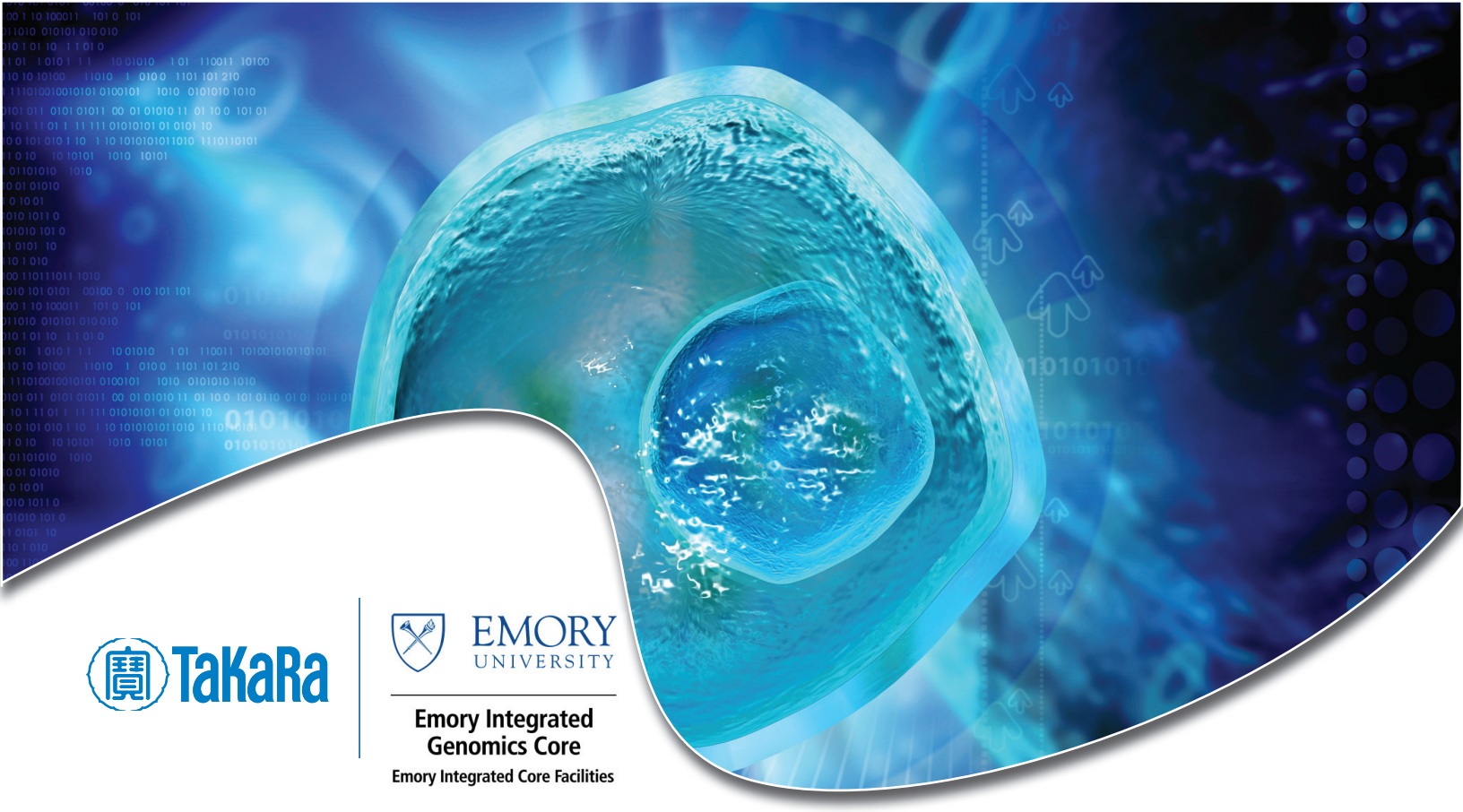

TaKaRa

EMORY UNIVERSITY

Emory Integrated Genomics Core

Emory Integrated Core Facilities

# 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® 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.

**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:**
December 7, 2022
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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