

Moving from Bulk Next-Gen Sequencing (NGS) to Precision Sequencing with Single-Cell Genomics: Resolving Heterogeneity in Blood and Solid Tumors

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Tuesday,

Feb. 5

2:30 p.m. - 3:30 p.m.

Richard Jones Hall, 4320
3181 S.W. Sam Jackson Park
Rd.
Portland, OR 97239

Current bulk sequencing methods are inadequate to fully characterize cancer. Targeted single-cell DNA analysis resolves tumor heterogeneity, which has critical implications in understanding clonal evolution and the acquisition of therapeutic resistance.

Join us to learn about the applications of the Tapestry Platform.

Please register at:

https://missionbio_ohsu_rjh.eventbrite.com/

Hosted by the OHSU Integrated Genomics Laboratory (IGL) and University Shared Resources (USR) program.

For questions, please contact Aaron Larson
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