Explore nascent RNA with Bru-seq technologies

Profile the nascent transcriptome with **Bru-seq**

Map transcription start sites, enhancer RNA and PROMPTs with **BruUV-seq**

Measure RNA stability with **BruChase-seq**

Calculate RNA PolII elongation rates with **BruDRB-seq**

**Bru-seq lab Core services**

- Total RNA and nascent Bru-RNA isolation
- cDNA library preparation
- Paired-end 150bp Next-Generation Sequencing
- NGS data processing
- Data storage

Learn more: https://umich.corefacilities.org/service_center/show_external/5763