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





