



Non-coding RNA and Epitranscriptomic Solutions



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Accurately profile circular RNAs by highly specific circular junction probe design

LncRNA Arrays

Overcome the limitations of RNA-seq for lncRNAs often at low abundance

Small RNA Arrays

Accurately profile miRNA, pre-miRNA, tRNA, tsRNA, and snoRNA simultaneously

Epitranscriptomic Arrays

Quantify the percentage of m6A modifications at the transcript specific level

m6A Single Nucleotide Arrays

Locate and quantify the exact m6A site at single nucleotide resolution



GENE EDITING PRODUCTS

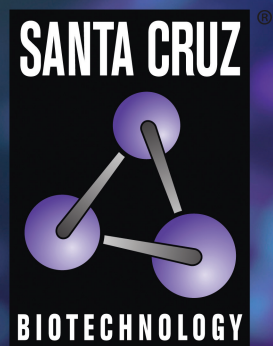
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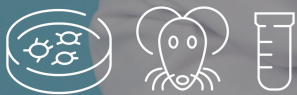


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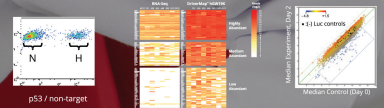


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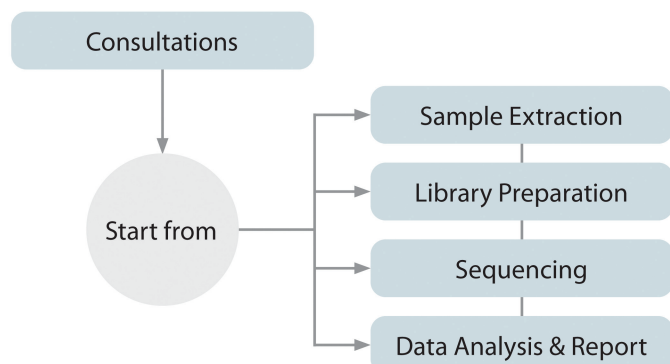
Lexogen NGS Services offers both standard and custom next-generation sequencing services, including a variety of RNA-Seq solutions and DNA-Seq. We also provide Bioinformatics Custom Service, where state-of-the-art bioinformatics is combined with a client-driven approach, all tailored to your needs. To deliver you the best results possible, our services, R&D, and bioinformatics teams work together to adapt our workflows and address your wildest research questions. Let our NGS experts turn your idea into reality!

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Each project starts with a consultation, where we agree on the details of the project and if needed give you advice on how to proceed based on your research questions.

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- Start from Extraction
- Start from Library Prep
- Sequencing only
- Data Analysis only



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- ✓ Shipment Support
- ✓ Various applications, including gene expression profiling, whole transcriptome sequencing, small RNA-Seq, ultra-low RNA-Seq, single-cell RNA-Seq, and DNA-Seq
- ✓ Expertise in handling challenging samples, like FFPE curls, biobank samples, laser microdissections, needle biopsies, etc.
- ✓ Quality Control (QC) at multiple steps along the workflow
- ✓ Spike-in RNA controls to monitor and evaluate RNA-Seq experiments and data quality (optional)
- ✓ Highest standards of data protection

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www.lexogen.com/services/configurator