



Long non-coding RNAs (IncRNAs) are defined as ≥ 200 nucleotide long RNAs, which regulate gene expression at epigenetic, transcriptional, and post-transcriptional levels through binding to DNA, RNA, or protein.

In this post-genomic era, IncRNAs are gaining increasing attention in areas of medical and agricultural research.

Our Key Features & Advantages



Directional Library

We offer directional library after rRNA removal, adding information on the originating strand of transcripts.



Comprehensive Integrated Analysis

Comprehensive integrated analysis of LncRNA and mRNA with one library construction.



Reliable & Faster Turnaround Time

The strategic locations of our labs across Asia and our extensive NGS system can provide faster turnaround time.



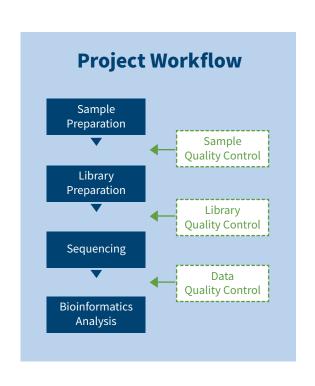
Extensive Experience

We have extensive records of sequencing projects that have been published in journals.



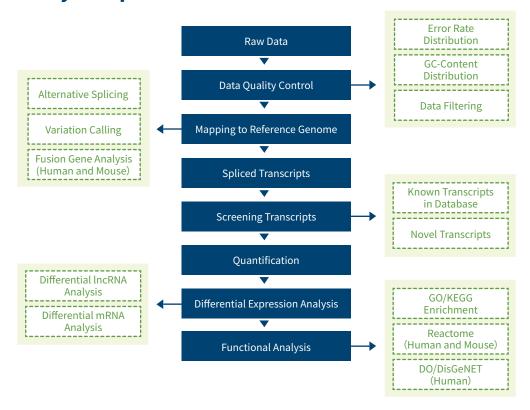
Unsurpassed Data Quality

We guarantee that \geq 80% of bases have a sequencing quality score \geq Q30, exceeding Illumina's official guarantee of \geq 75%.





Standard Analysis Pipeline



Sample Requirements

Library Type	Sample Type	Amount	Volume	Concentration	RIN	Purity (Nanodrop/ Agarose Gel)
IncRNA Library (250-300bp insert directional library with rRNA removal)	Total RNA	≥ 0.5 μg	≥ 20 μL	≥ 50 μL	Animal ≥ 6.5 Plant/Fungi ≥ 6, with the smooth baseline	OD260/280 ≥ 2.0
Exosomal IncRNA Library	Exosome Total RNA	≥ 5 ng	≥ 10 μL	-	Fragments distributing between 25-200nt (by high sensitive Agilent Bioanalyzer 2100), FU>10, with no peak>2000nt	OD260/230 ≥ 2.0; no degradation, no contamination

Publications

Journal	IF	Title
Molecular Cancer	10.68	A novel IFNa-induced long non-coding RNA negatively regulates immunosuppression by interrupting H3K27 acetylation in head and neck squamous cell carcinoma (2020.01)
Nature Genetics	25.46	Analysis of the androgen receptor-regulated lncRNA landscape identifies a role for ARLNC1 in prostate cancer progression (2018.06)

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