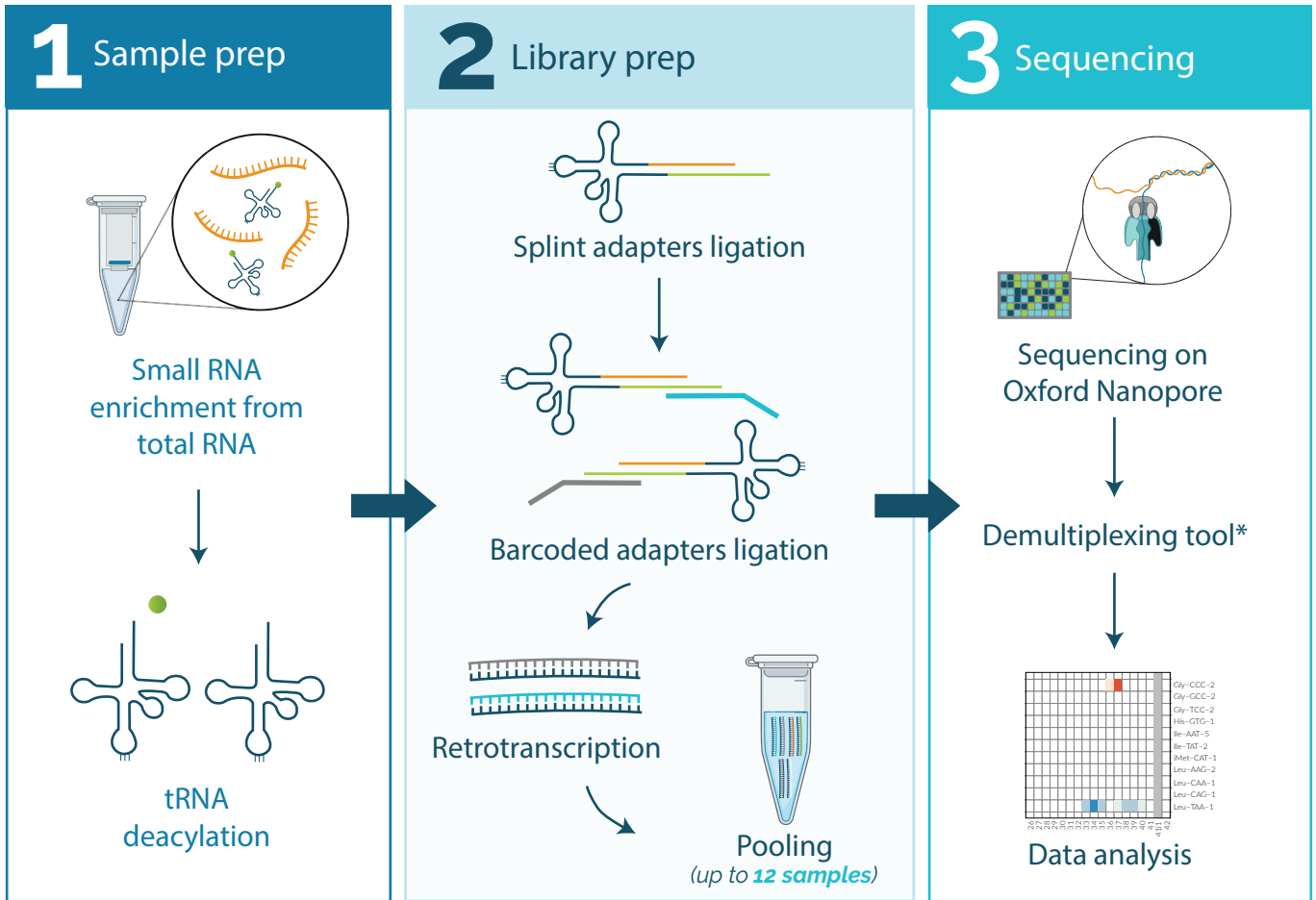


# nano-tRNAseq PRO kit

Built for flexible tRNA analysis across all species

Introducing **nano-tRNAseq PRO (12 samples - 12 barcodes)**, our kit enabling full-length sequencing of native tRNA molecules in a gel-free workflow. Starting from total RNA, it allows reliable quantification of **tRNA abundance** and detection of **chemical modifications**. Building on our previous version, it now supports multiplexing of up to **12 samples (instead of 6)**, enabling more flexible sample pooling while reducing sequencing costs and batch-to-batch variability. Compatible with MinION and PromethION nanopore sequencing platforms.

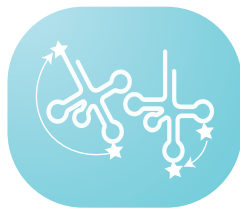


\*License provided with kit

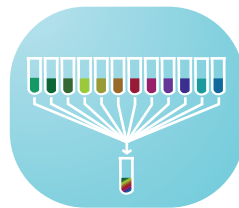
## Highlights



Full-length tRNA sequencing



Chemical modification detection



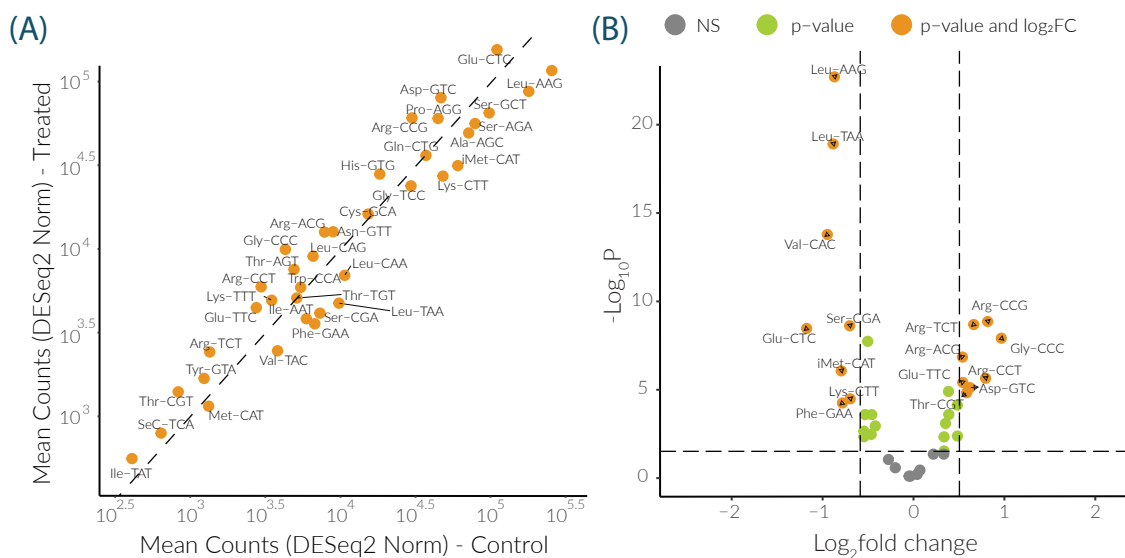
Up to 12 samples multiplexing



Reduced sequencing costs

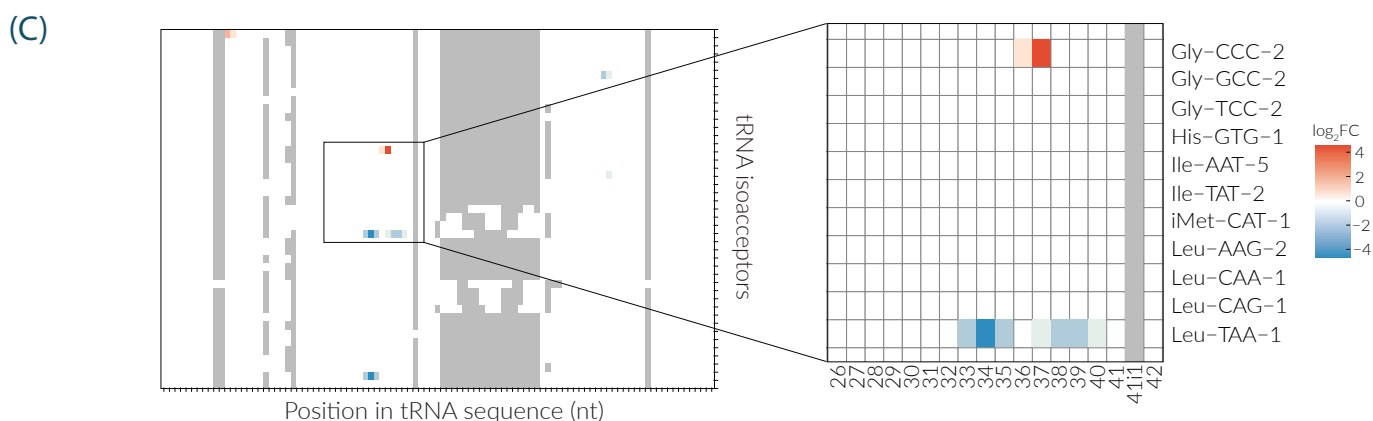
# Explore tRNA biology at isoacceptor resolution

## Uncovering tRNA expression dynamics



Nano-tRNAseq kit coupled with Immagina BioIT pipeline enables detection and analysis of tRNA abundance. **(A)** Scatter plot showing mean counts of tRNA abundances in treated vs control samples. **(B)** Volcano plot highlighting differentially expressed tRNA abundances in treated vs control samples.

## Position-resolved modifications change across tRNA isoacceptors



Chemical modifications are identified based on basecalling errors detected on every position of all tRNA isoacceptors **(C)** Heatmap shows significant changes in modifications observed in treatment vs control. Blue indicates possible loss of modification in treated sample; red indicates insertion of modifications in treated sample.

## Specifications

Species	All species*	Kit size	12 rxns <b>(12 barcodes)</b>
Total RNA	$\geq 3 \mu\text{g}$ of total RNA	Workflow time	2 days

\*with annotated genome

SCAN ME FOR MORE INFO

