CORALL RNA-Seq V2 -

For all your sequencing experiments!

RNA-Seq from FFPE Samples with Ease

Get the best sequencing results for all sample types, incl. FFPE, with CORALL RNA-Seq V2 – Lexogen's fast and flexible whole transcriptome RNA-Seq Library Prep Kit.

Exceptional Performance on FFPE Samples

The robust and fragmentation-free workflow including RiboCop rRNA depletion is the ideal choice for degraded and FFPE RNA (Fig. 1). The sensitive protocol with built-in unique molecular identifiers (UMIs) delivers high-quality RNA-Seq data for all RNA input amounts and qualities.
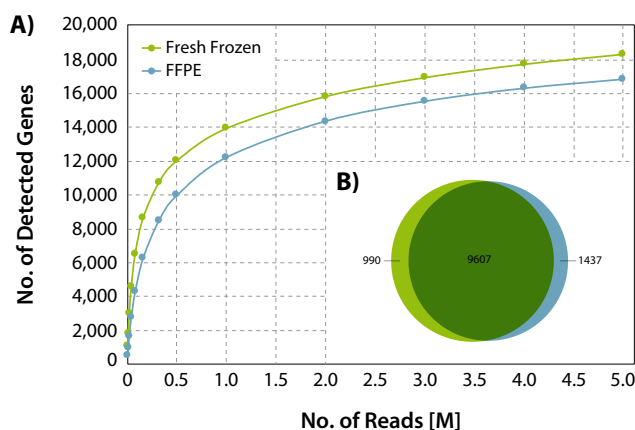

Figure 1 | Gene discovery (A) and overlap of detected genes (B) for Fresh Frozen and FFPE derived human liver RNA. The overlap of detected genes is measured at normalized expression levels >5 CPM (for uniquely mapping reads).

Benefits:

- **NEW!** Fragmentation-free generation of longer library inserts for isoform and fusion detection, alternative splicing and more.

- Only 6 steps and 4.5 hours processing time: Sample to sequencing in only one day.

- Fully automatable protocol.

- Any input (1 ng - 1000 ng) and any sample type, including degraded and FFPE RNA.

Interested to learn more?

Check the product page for more information on our website!