

CORALL RNA-Seq V2!

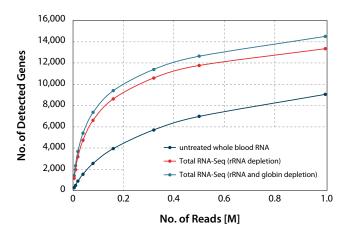
For whole blood sequencing experiments that are not in vein!

Increase Gene Detection from Blood Samples

Get the best sequencing results with convenient CORALL V2 workflows for whole blood samples!

Whole Transcriptome Blood RNA-Seq

CORALL RNA-Seq V2 combined with simultaneous rRNA and globin depletion provides a highly convenient workflow for processing whole blood samples and significantly increases gene detection (Fig 1).



90 80 70 60 40 30 20 10 0 untreated whole mRNA-Seq mRNA/globin depletion and depletion and

Figure 2 | CORALL mRNA-Seq V2 with an additional rRNA and globin depletion step (Cat. No. 186 + Cat. No. 157) efficiently removes globin mRNA from human whole blood RNA for mRNA-Seq applications.

Figure 1 | Increased gene detection for CORALL Total RNA-Seq with rRNA and globin depletion (Cat. No. 186). Number of detected genes as a function of the number of reads uniquely mapping to exons.



Interested to learn more?

Check the product page for more information!

Benefits:

- Unlock the potential of whole blood RNA-Seq.
- Optimize your workflow, increase gene detection.
- Save time with a fully automatable workflow.

mRNA-Seq Workflow from Whole Blood

RNA of interest

100

Sequence what matters most by combining CORALL mRNA-Seq with rRNA and globin depletion to free up sequencing space for your mRNAs of interest (Fig. 2).

Globin

rRNA

mRNA-Se