



Quickly go from RNA sample to RNA-seq insight

Discover QIAGEN's complete RNA-seq workflow, including our new web-based bioinformatics pipeline



Overcome the barriers to a successful RNA-seq experiment with our comprehensive transcriptomics workflow, now including the RNA-seq Analysis Portal. Go from sequence to biological insights in just a few hours with this automated, easy-to-use tool.

You don't need specialized expertise or additional computer resources – just your internet browser.



Learn more about the RNA-seq Analysis Portal at www.qiagen.com/rap

Empower your data analysis with the RNA-seq Analysis Portal

Knowledge is power

Get differential gene expression, plus the top 10 canonical pathways, upstream regulators and diseases.



Save time and maximize efficiency

Cut down analysis time from days to just a few hours and scale up your RNA-seq projects.



Complete autonomy

Take full control of your RNA-seq pathways and biomarker discovery with a fully integrated analysis pipeline.



Build gene expression workflows

Seamlessly move from experimental design to RNA-seq data analysis to qPCR or dCPR based validation using a single interface.



Compatibility

Supports analysis of data generated from most commonly used library prep kits- QIAGEN, Illumina, Takara Bio, New-England Bio and Thermo Fisher Scientific

Access to the RNA-seq Analysis Portal is included as standard with QIAseq Stranded Total RNA Library, miRNA Library and UPX 3' Transcriptome Kits or can be obtained through purchase of GeneGlobe Analyze RNA-seq analysis credits.

RNA-seq Analysis Portal currently supports human, mouse and rat samples.

For up-to-date licensing information and product-specific disclaimers, see the respective QIAGEN kit handbook or user manual. QIAGEN kit handbooks and user manuals are available at www.qiagen.com or can be requested from QIAGEN Technical Services or your local distributor.

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