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KAPA RNA HyperPrep Kits

**Roche Sequencing partners with Genialis
data analysis to offer complete RNA solutions**

G Genialis

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Specifications

The table below describes the specifications for the KAPA RNA HyperPrep Kits, including all four workflows:

		KAPA RNA HyperPrep Kit	KAPA mRNA HyperPrep Kit	KAPA RNA HyperPrep Kit with RiboErase (HMR)	KAPA RNA HyperPrep Kit with RiboErase (HMR) Goblin (for blood samples)
	Input	1 ng - 100 ng	50 ng – 1 µg	25 ng – 1 µg	25 ng – 1 µg
Input Type	High-quality total RNA	Recommended			
	Low-quality total RNA (e.g. FFPE RNA)	Compatible	Not Recommended	Not Recommended	Compatible
	Poly(A)-selected and/or depleted RNA	Recommended	N/A	N/A	N/A
Sample Type	Species	Eukaryotic, microbe, plant, etc.	Eukaryotic,plant, etc.,	Human, mouse, rat	
Workflow	Fragmentation	Tunable			
	Library Prep Time	4 hrs	5.5 hrs	6.5 hrs	6.5 hrs
	Automation Friendly	Yes			

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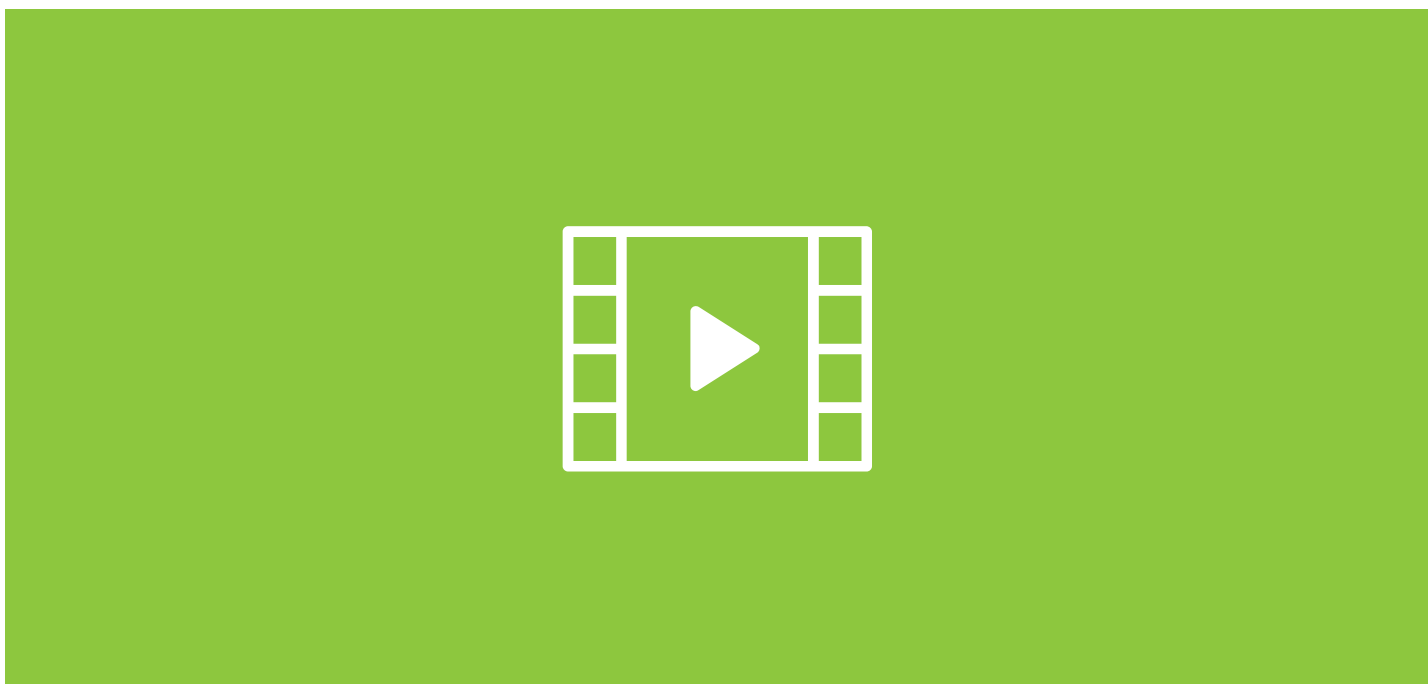
Featured Resources

(/content/dam/rochesequence/worldwide/resources/applicationnote-whole-transcriptome-sequencing-from-ffpe-derived-rna-SEQ100281.pdf)

Whole Transcriptome Sequencing from FFPE-derived RNA.
(/content/dam/rochesequence/worldwide/resources/applicationnote-whole-transcriptome-sequencing-from-ffpe-derived-rna-SEQ100281.pdf)

(/content/dam/rochesequence/worldwide/resources/roche-and-genialis-collaboration.mp4)
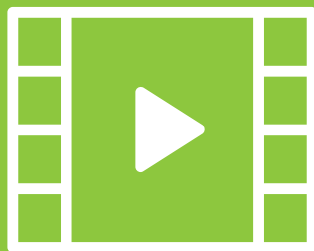
Roche and Genialis collaboration (/content/dam/rochesequence/worldwide/resources/roche-and-genialis-collaboration.mp4)

(/content/dam/rochesequence/worldwide/resources/brochure-sample-prep-solutions-SEQ100215.pdf)

Roche Sample Prep Solutions for RNA Sequencing.
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(/content/dam/rochesequence/worldwide/resources/KAPA-RNA-HyperPrep-Webinar-Improved-performance-with-degraded-inputs-tumor-profiling-applications.mp4)

Working with degraded inputs and tumor profiling applications

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Selective Transcript Depletion. (/content/dam/rochesequence/worldwide/resources/applicationnote-selective-transcript-deletion-SEQ100319.pdf)

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