October 2019

# Quick-Start Protocol

# QIAseq<sup>®</sup> FastSelect<sup>™</sup> –rRNA HMR and/or –Globin with the QIAseq Stranded Total RNA Lib Kit

The QIAseq FastSelect Kits for -rRNA HMR (cat. nos. 334386, 334387, 334388), -Globin (cat. nos. 334376, 334377, 334378), and -rRNA/Globin (cat. nos. 335376, 335377, 335378) may be used with the QIAseq Stranded Total RNA Lib Kit (cat. no. 180743 or 180745) to remove human, mouse, or rat rRNA and/or globin.

All components of QIAseq FastSelect should be stored at -30 to  $-15^{\circ}$ C in a constant-temperature freezer.

#### Further information

- QlAseq FastSelect -rRNA HMR and -Globin Handbook: www.qiagen.com/HB-2670
- Safety Data Sheets: www.qiagen.com/safety
- Technical assistance: support.qiagen.com

# Notes before starting

- The QIAseq Stranded Total RNA Lib Kit is required for use with this protocol.
- Refer to the *QlAseq Stranded Total RNA Lib Kit Handbook* available at www.qiagen.com/HB-2465



### Procedure

- 1. Thaw total RNA on ice. Gently mix, briefly centrifuge to collect residual liquid from the sides of the tubes, and return to ice.
- 2. Prepare the reagents required for the RNA fragmentation and QIAseq FastSelect rRNA and/or globin removal.
  - 2a. Thaw 5x RT Buffer, nuclease-free water from the QIAseq Stranded kit, and the tube(s) from the QIAseq FastSelect kit at room temperature.
  - 2b. Mix by vortexing and then briefly centrifuge.
- On ice, prepare the fragmentation/RNA Removal reaction according to Table 1. Briefly centrifuge, mix by pipetting up and down 10 times, and centrifuge briefly again.
  Note: If setting up more than one reaction, prepare a volume of Master Mix that is 10% greater than what is required for the total number of reactions.

Table 1. Setup	of fragmentation	on/RNA Remov	al reactions
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Component	Volume/reaction
Total RNA (100 ng – 1 μg)	Variable
RT Buffer, 5x*	اµ 8
QIAseq FastSelect –rRNA HMR <sup>†</sup>	1 µl
QIAseq FastSelect –Globin <sup>†</sup>	1 µl
ERCC Control <sup>‡</sup>	Optional
Nuclease-free water	Bring total reaction volume to 37 µl
Total volume	37 µl

\* From QIAseq Stranded Total RNA Lib Kit.

 $^{\scriptscriptstyle \dagger}$  Choose QIAseq FastSelect –rRNA HMR and/or QIAseq FastSelect –Globin.

<sup>‡</sup> ERCC Control RNA can be added according to the concentrations specified by the manufacturer. If added, the total fragmentation/RNA Removal reaction volume should remain 37 μl.

4. Incubate as described in Table 2, according to your input RNA quality and desired insert size.

Input RNA quality	Step	Insert size ~150–250 bp	Insert size ~350 bp
High quality (RIN >9)	1*	15 min at 95°C	3 min at 95°C
Moderate quality (RIN 5-6)	1*	10 min at 95°C	3 min at 95°C
FFPE or degraded sample (RIN <3)	1*	No fragmentation <sup>†</sup>	No fragmentation $^{\dagger}$
Steps 2–9 are performed regardless	2	2 min at 75°C	2 min at 75°C
	3	2 min at 70°C	2 min at 70°C
	4	2 min at 65°C	2 min at 65°C
of input RNA quality. They need to	5	2 min at 60°C	2 min at 60°C
be performed whether the RNA is high quality, moderate quality, FFPE, or degraded.	6	2 min at 55°C	2 min at 55°C
	7	2 min at 37°C	2 min at 37°C
	8	2 min at 25°C	2 min at 25°C
	9	Hold at 4°C	Hold at 4°C

Table 2. Combined QIAseq fragmentation and FastSelect hybridization protocol

\* Choose one option for the time on step 1 according to the input RNA quality and desired insert size.

<sup>†</sup> Also suitable for exosomal RNA or RNA of other origin with a size between 80–500 bp.

#### Important: Regardless of time and temperature chosen in step 1, steps 2-9 must be performed.

- 5. Refer to the *QlAseq Stranded Total RNA Lib Kit Handbook* and immediately proceed to "Protocol: First-strand Synthesis".
- 6. Follow the *QlAseq Stranded Total RNA Lib Kit Handbook* to perform all remaining library construction steps.

**Important**: When removing globin, 2 additional cycles of CleanStart<sup>®</sup> Library Amplification need to be performed.

## **Revision History**

Date	Changes
10/2019	Initial release



Scan QR code for handbook.

For up-to-date licensing information and product-specific disclaimers, see the respective QIAGEN kit handbook or user manual.

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