

# SMARTer® Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian Components

Catalog Nos.	Amount	Lot Number
634489 (Not sold separately; sold as a part of 634485)	24 rxns	Specified on product label.
634490 (Not sold separately; sold as a part of 634486)	48 rxns	Specified on product label.
634491 (Not sold separately; sold as a part of 634487)	96 rxns	Specified on product label.
634492 (Not sold separately; sold as a part of 634488)	192 rxns	Specified on product label.

## **Description**

The SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian Components are used with a SMARTer RNA Unique Dual Index Kit - 24U as part of SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian to generate strand-specific RNA-seq libraries for Illumina® sequencing from 10–1,000 cells or 250 pg–10 ng inputs of purified total RNA. The Components incorporate Takara Bio's proprietary SMART® (Switching Mechanism at the 5' end of RNA Template) technology and include refinements to the SMARTer method for stranded RNA-seq that simplifies the library preparation workflow and improves sequencing performance.

The SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian was specifically designed to deliver highly sensitive and reproducible data from low-input RNA while keeping the workflow short and user friendly. The incorporated UMIs (Unique Molecule Identifiers) help customers perform UMI-based correction to obtain more reliable results. The kit does not require additional rRNA removal methods and produces sequencing libraries that retain strand-of-origin information. The integrated removal of cDNAs derived from rRNA—typically present in high abundance following cDNA synthesis from total RNA inputs—makes the workflow extremely sensitive, yielding data that is highly reproducible with low mapping to rRNA.

634492

## Package Contents

634489

634490

Pacl	kage	1	:
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	(24 rxns)	(48 rxns)	(96 rxns)	(192  rxns)	
	110 µl	225 μ1	450 μl	900 μl	SMART UMI-TSO Mix v3*
	40 μ1	80 µl	160 µl	320 μ1	R-Probes v3*
	5 μl	5 μ1	5 μl	5 μ1	Control Total RNA (1 µg/µl)
Package 2:					
_	634489	634490	634491	634492	
	(24 rxns)	(48 rxns)	( <u>96 rxns)</u>	(192 rxns)	
	40 μ1	80 µl	160 μ1	320 μ1	ZapR® v3
	30 μ1	55 µl	110 μl	220 μ1	SMART Pico Oligos Mix v3
	100 μ1	200 μ1	400 μ1	800 μ1	5X First-Strand Buffer
	50 μ1	100 μ1	200 μ1	400 μ1	SMARTScribe™ II Reverse Transcriptase
	100 μ1	200 μ1	400 μ1	800 μ1	ZapR Buffer (10X)
	30 μ1	60 μ1	100 μ1	200 μ1	RNase Inhibitor (40 U/µl)
	100 μ1	200 μ1	2 x 200 µl	4 x 200 μl	SeqAmp <sup>™</sup> DNA Polymerase
2	x 1.25 ml	4 x 1.25 ml	10 ml	2 x 10 ml	SeqAmp CB PCR Buffer (2X)

634491

Takara Bio USA, Inc.

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634489 (24 rxns)	634490 (48 rxns)	634491 ( <u>96 rxns)</u>	634492 (192 rxns)	
100 μ1	200 μ1	400 μ1	800 μ1	PCR2 Primers v3
2 x 1.25 ml	4 x 1.25 ml	10 ml	2 x 10 ml	Nuclease-Free Water
1.25 ml	2 x 1.25 ml	5 ml	2 x 5 ml	Tris Buffer (5 mM)
0.5 ml	1 ml	1 ml	2 x 1 ml	10X Lysis Buffer

<sup>\*</sup> Takara Bio proprietary sequences

## **Storage Conditions**

- Store Package 1 components at -70°C.
- Store Package 2 components at -20°C.
- Do not freeze R-Probes v3 more than 3 times. We recommend aliquoting into multiple vials to avoid repeated freeze/thaw cycles

### **Expiration Date**

• Specified on product label.

## **Shipping Conditions**

Dry ice

### **Product Documents**

Documents for our products are available for download at <u>takarabio.com/manuals</u> The following documents apply to this product:

SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian User Manual

# **Quality Control Data**

To assess the quality of the reagents in the SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian Components, 250 pg of Control Total RNA (Human Brain) was converted to an Illumina-ready cDNA sequencing library with the SMARTer Stranded Total RNA-Seq Kit v3 - Pico Input Mammalian Components as described in the user manual, using 5 cycles in PCR1 and 16 cycles in PCR2 steps. Purified libraries were quantified using the Qubit 2.0 Fluorometer (Life Technologies) with a dsDNA HS Assay Kit, resulting in an output of at least 4 ng/ $\mu$ l. Libraries were diluted to 2 ng/ $\mu$ l, and 1  $\mu$ l of the dilution was analyzed with an Agilent Bioanalyzer and High Sensitivity DNA Kit (Agilent, Cat. No. 5067-4626). The Bioanalyzer electropherogram showed a broad distribution from 200–1,000 bp with an average size in the 350–510 bp range. The libraries were sequenced on an Illumina platform and confirmed to contain  $\leq$ 20% of the reads mapping to rRNA (5S, 5.8S, 18S and 28S) and  $\leq$ 7% of the reads mapping to mitochondrial ribosomal RNA (m12S and m16S).

It is certified that this product meets the above specifications, as reviewed and approved by the Quality Department.

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# SMARTer® Stranded Total RNA-Seq Kit v3 - Pico Input **Mammalian Components**

### CATALOG NOS.

634489, 634490, 634491 & 634492

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