



Feb 05, 2019

Addition of the adaptor to RNA substrates for 5' RACE

DOI

<https://dx.doi.org/10.17504/protocols.io.xsrfnd6>

Matus Valach¹

¹Université de Montréal, Montreal, Quebec, Canada



Matus Valach

Université de Montréal, Montreal, Quebec, Canada

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.xsrfnd6>

Protocol Citation: Matus Valach 2019. Addition of the adaptor to RNA substrates for 5' RACE. **protocols.io**
<https://dx.doi.org/10.17504/protocols.io.xsrfnd6>

License: This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: February 05, 2019

Last Modified: February 05, 2019

Protocol Integer ID: 20017

Keywords: RNA, RNA-Seq, 5' RACE, using t4 rna, rna termini by rt, t4 rna, rna termini, rna substrate, rna, pcr after the addition, pcr, addition of the adaptor

Abstract

Simple protocol for mapping 5'-P RNA termini by RT-PCR after the addition of a 5' adaptor using T4 RNA ligase 1.

Materials

MATERIALS


 T4 RNA Ligase 1 (ssRNA Ligase) - 1,000 units
 New England Biolabs
 Catalog #M0204S

Before start

Prepare samples, including controls, according to the aim of the experiment. Use the chart below to decide which enzyme is appropriate for pre-treatments of the RNA. For example, if only interested in mapping 5' RNA ends, use T4 polynucleotide kinase with ATP and without ATP to infer the original phosphorylation state of the end of interest.

Setup of the phosphorylation reaction can be found here: [dx.doi.org/10.17504/protocols.io.cpdvi5](https://doi.org/10.17504/protocols.io.cpdvi5) However, avoid heat denaturation of the enzyme and rather purify the RNA using a trizol extraction (e.g., [http://dx.doi.org/10.17504/protocols.io.eiebcbe](https://doi.org/10.17504/protocols.io.eiebcbe)) or a column clean-up (e.g., Monarch RNA Cleanup Kits from NEB is optimal when interested in small RNA molecules <200 nt, which is the usual exclusion limit in other products).

	Original	Original	T4PNK (+ATP)	T4PNK (+ATP)	T4PNK (-ATP)	T4PNK (-ATP)	T4PNK 3'Pase (-) (+ATP)	T4PNK 3'Pase (-) (+ATP)	T4PNK 3'Pase (-) (-ATP)	T4PNK 3'Pase (-) (-ATP)
	5'	3'	5'	3'	5'	3'	5'	3'	5'	3'
	P	P	P	OH	P	OH	P	P	P	P
	P	OH	P	OH	P	OH	P	OH	P	OH
	OH	P	P	OH	OH	OH	P	P	OH	P
	OH	OH	P	OH	OH	OH	P	OH	OH	OH

- 1 Mix the following components (10 μ L):

	Component	Amount [μ L]	Final concentration
	RNA [1 μ g]	4	100 ng/ μ L
	5' RACE RNA oligo [100 μ M]	2	20 μ M
	RNase-free water (ddH ₂ O)	4	

- 5' and 3' ends of the 5' RACE oligo are hydroxylated.

- 2 Denature for 2 min at 70 °C, place on ice.

- 3 Mix the following components (20 μ L):

	Component	Amount [μ L]	Final concentration
	RNA + oligo mix (step 2)	10	
	10 $\times$ RNA ligase buffer	2	1 $\times$
	50% PEG-8000	6	15%
	ATP [10 mM]	2	1 mM
	T4 RNA ligase 1 [10 U/ μ L]	2	1 U/ μ L

- 4 Incubate using the following program: 60 min at 25 °C, then 60 min at 16 °C, and then for 120 min at 4 °C.
- 5 Purify the RNA from the RNA ligase reaction (e.g., trizol extraction or column clean-up).
- 6 Proceed to RT-PCR. For the RT reaction, use 250-500 ng of the purified adaptor-ligated RNA and random hexamers (or a primer of choice). For the PCR, use an upstream



(forward) primer binding to the 5' adaptor and a downstream (reverse) primer binding to the RNA of interest.