



Jun 24,
2019

RNA extractions from de-etiolated Arabidopsis seedlings using CTAB

Akila Wijerathna-Yapa¹, Andrew Bowerman², Diep Ganguly²

¹School of Molecular Sciences - The University of Western Australia, ²The Australian National University

Working

[dx.doi.org/10.17504/protocols.io.3f6gjre](https://doi.org/10.17504/protocols.io.3f6gjre)

Pogson Genomics Group



Diep R. Ganguly
The Australian National University



ABSTRACT

A CTAB based method for extracting RNA, which is particularly useful for tough tissues. This has been adapted from a protocol that was originally used for pine tree tissue, which is difficult due to the high concentrations of polysaccharides, phenolics, and RNase (Chang, S., Puryear, J. & Cairney, J. Plant Mol Biol Rep (1993) 11: 113. <https://doi.org/10.1007/BF02670468>). The protocol described herein was an effective alternative to TRIzol based extractions for recovering RNA from juvenile de-etiolated tissues.

GUIDELINES

Ensure that you use RNA friendly practices (e.g. clean surfaces with RNasey or 80% ethanol, use RNase-free filter tips, use DEPC-treated water for buffers, adjust RNA buffer pH to be slightly acidic [e.g. ~6] in which RNA is more stable).

MATERIALS

NAME ▾	CATALOG # ▾	VENDOR ▾
Isoamylalcohol		
Beta-mercaptoethanol		
Chloroform		
DEPC		
Hexadecyltrimethylammonium bromide (CTAB)	H9151	Sigma Aldrich
80% Ethanol		
Lithium chloride	793620	Sigma Aldrich
Polyvinylpyrrolidone K 30	81420	Sigma Aldrich

MATERIALS TEXT

Extraction buffer

1. 2% CTAB
2. 2% PVP K30
3. 100 mM Tris-HCl (pH 8.0)
4. 25 mM EDTA (pH 8.0)
5. 2 M NaCl
6. Autoclave
7. 2% 2-mercaptoethanol (add before use = 100 uL per 5 mL buffer)

SAFETY WARNINGS

Perform steps with 2-mercaptoethanol in a fume hood.

- 1 Harvest tissues into liquid nitrogen and grind (under liquid nitrogen) using mortar and pestle to obtain a fine powder. Ground tissue should be kept in safe-lock tubes in liquid nitrogen (or returned to -80°C storage) until all samples are processed. This step is critical for efficient extractions so take your time here.
- 2 Add 1 mL of prewarmed (65°C) extraction buffer to each tube, mix well and incubate for 5 mins at 65°C (can be longer if needed e.g. 10-15 mins)
- 3 Add 200 µl of chloroform:IAA (24:1), mix well and spin @ 14,000 rcf for 10 mins. Remove upper aqueous phase to a new tube. Make sure not to disturb or pipette any material from the interface. Repeat chloroform:IAA step twice, being increasingly conservative when recovering the aqueous phase.
- 4 Add equal volume of 5 M LiCl to aqueous layer, mix well and incubate overnight @ -20°C.
- 5 Spin tubes @ 14,000 rcf for 20 mins @ 4°C.
- 6 Remove supernatant with pipette, add 1 mL 80% ethanol and invert tube ~10X. Centrifuge @ 7,500 rcf for 5 minutes @ RT. Remove ethanol and repeat.
- 7 Remove as much ethanol as possible using a pipette. Air-dry tubes with lid open for 1 min. Resuspend in 30 - 50 µl (depending on expected concentration and purpose, e.g. we aim for ~ 500 ng / µl) DEPC-treated water or low EDTA TE buffer (0.1 mM EDTA, 10 mM Tris base, pH 6.5).
- 8 Check quality of RNA e.g. visualize ~50 ng RNA on a 1% agarose gel or use BioAnalyser / LabChip GXII.



This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited