## **Customer Developed Protocol**

# CTAB Protocol for Isolation DNA from Plant Tissue

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#### CUSTOMER DEVELOPED PROTOCOL

CTAB Protocol for Isolation DNA from Plant Tissue

## Contributor Research Profile

The Van Deynze lab at UC, Davis is a Plant Molecular Biology, Biochemistry and Genomics lab that works on elucidating the basis of capsaicin synthesis and resistance to phytophthora in pepper.

Read more about the Van Deyzne Lab: <u>http://biosci3.ucdavis.edu/Faculty/Profile/View/14262</u> **Contact:** avandeyzne@ucdavis.edu

## Reference

This protocol was used in:

Amanda M. Hulse-Kemp, *et al.*, Reference Quality Assembly of the 3.5 Gb genome of *Capsicum annuum* from a Single Linked-Read Library, *Biorxiv*. 2017 June 20. https://www.biorxiv.org/content/early/2017/06/20/152777

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## **Required Buffers and Reagents**

- 1. AMPure XP Beads (Beckman Coulter)
- 2. DNA Extraction Buffer:

100 mM Tris (pH 8.0) 20 mM EDTA 2% CTAB 1.2 M NaCl 0.1% ß- mercaptoethanol (Add before use. After adding ß- mercaptoethanol, the Extraction Buffer must be used within a day)

- 2. Chloroform-Isoamylalcohol (CIA) 24:1 (v/v)
- 3. Dilution Buffer:

100 mM Tris Buffer (pH 8) 20 mM EDTA 2% CTAB

#### 4. High Salt TE:

10 mM Tris (pH 8) 2 mM EDTA 1 M NaCl (add RNase A at 50 μg/ml)

6. Wash Buffer:

Add 300 ml TE to 700 ml ethanol

#### 7. TE:

10 mM Tris 1 mM EDTA (pH 8)

## **DNA Extraction Protocol**

- For freeze-dried tissue, grinding is done using a disposable microcentrifuge pestle directly in a 2 ml microcentrifuge tube (~200 mg tissue). Keep samples cold until the DNA Extraction Buffer is added. For seeds, grinding is done using a Geno/Grinder 2010 by placing one 5 mm tungsten ball into a rounded bottom 2 ml microcentrifuge tube.
- 2. Add 750  $\mu$ l Extraction Buffer to the ground tissue using a multichannel pipette.
- 3. Cap the tubes and mix well. Incubate at 60°C for 1 hour. Mix the tubes gently every 10-15 minutes during the incubation period.
- 4. Cool the tubes to room temperature for 5 min, then add 750 μl Chloroform:Isoamylalcohol (24:1, v/v) to each tube, mix well and centrifuge at 3000 g for 15 minutes.
- 5. Transfer the aqueous layer (~500  $\mu$ l) to a new set of labeled tubes.

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- 6. Add 1 ml Dilution Buffer to the aqueous phase.
- Mix well and incubate at 60°C for ~30 minutes. (Note: Significant amounts of precipitated DNA-CTAB complex should be observed at the end of incubation).
- 8. Centrifuge at 4500 g for 15 minutes and discard the supernatant.
- 9. Add 1 ml Wash Buffer to the pellet and soak it in at room temperature for 30 minutes to remove any excess CTAB.
- 10. Centrifuge at 4500 g for 15 minutes and discard the supernatant.
- 11. Re-suspend the DNA pellet in 100  $\mu$ l high salt TE with RNase A and incubate at 60°C for 30 minutes.
- 12. Transfer the high salt TE DNA solution to a 96-well microtiter plate, one sample per well.
- 13. Add 1X volume of AMPure XP beads (i.e. 100 µl sample + 100 µl AMPure XP beads).
- 14. Mix well by pipetting up and down.
- 15. Incubate the plate for 10-15 minutes.
- 16. Place on a magnetic stand for 2-5 minutes (or until clear) to collect beads.
- 17. Gently remove the solution without disturbing the bead pellet.
- 18. Add 150 µl 75% ethanol solution (keep plate on the magnet).
- 19. Wait 30 seconds and remove ethanol without disturbing the beads.
- 20. Repeat the ethanol wash and then let the beads air dry for 2-5 min (make sure you do not over-dry the beads as it will make re-suspension harder).
- 21. Add 100 µl Tris/H2O to each sample to re-suspend the DNA, mix well by pipetting up and down.
- 22. Incubate plate for 10-15 minutes.
- 23. Place on a magnetic stand for 2-5 minutes (or until clear) to collect beads.
- 24. Transfer the solution into a new 96-well plate (stock DNA).

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