Plant DNA extraction using the DNAeasy® Plant Mini Kit (Qiagen)

PS: ensure the water bath is set at 65°C before you commence extraction

A. Prepare and label extraction tubes



2. QIAshredder columns (purple/pink)



3. DNA easy mini spin column (colourless)

NB: One tube for each represents one sample.

- a) Label two sets of 2ml microcentrifuge (Eppendorf)tubes (1 for each sample). First set will be used for adding **homogenized sample** and the second set will be used for **binding DNA**.(Not provided with the kit).
- b) Label the QIAshredder columns and the DNAeasy mini spin column
- c) Label 1.5ml Eppendorf tube for final **DNA elution** provided with the kit

B: Procedure

I. Tissue disruption/ homogenization

1. Disrupt Cells Grind plant tissue into a fine powder in liquid nitrogen using a mortar and pestle/genogrinder. Transfer up to 20 mg of the powder to a 2ml microcentrifuge tube. Keep the sample on ice for immediate use or freeze at -70 °C for future use.

II. Lyse Cells

2. Add **400 \muL** of Buffer **AP1** and **4 \muL** of **RNAse A**. Thoroughly mix by vortexing and inverting Incubate the mixture at 65 °C for 10 minutes with occasional inversion (2 to 3 times)

III. Precipitation of DNA and removal of Debris

- 3. Add **130** μ L of **Buffer P3** to the mixture; mix completely by inversion and place the sample on ice for 5 minutes.
- 4. Centrifuge the sample at maximum speed (12,000–16,000 5 g) for 5 minutes to pellet the cellular debris, proteins, and polysaccharides.
- 5. Carefully pipette the supernatant from step 4 and transfer onto the QIAshredder spin column placed in a collection tube (purple/pink). PS: Avoid disrupting the debris settled at the bottom while pipetting. Centrifuge at a maximum speed for 2 minutes. This removes any cellular debris not removed in step 4 above

NB: Discard the filtration column but retain the collection tube since it will now have the flowthrough which contains the DNA.

IV. Binding DNA

- 6. Transfer the (<u>~450 ml</u>) flowthrough (from 5 above) into a new 2ml eppendorf tube without disturbing the pellet if present. Then add 1.5 volumes (<u>~675ml</u>) of Buffer AW1 and mix by inverting 4 -5 times. PS: hold onto the cap tightly to avoid spillage.
- 7. Transfer **650** µL of mixture from step 6 into a DNA Mini spin column(colourless/white) placed in a 2ml collection tube. Centrifuge for 1min at 8000rpm. **Discard the flow-through liquid; retain the collection tube.** Return the column to the collection tube. Apply the remaining mixture from step 6 onto the column. Repeat the centrifugation as above and discard the flow-through liquid and collection tube.

V. First and second column wash

- 8. Place the spin column into a new 2ml collection tube. Add **500 μL Buffer AW2**, and centrifuge for 1min at 8000rpm. Discard the flow through
- 9. Add another 500 μL Buffer AW2. Centrifuge for 2minutes and maximum speed. NB: Remove the spin column carefully from the collection tube so that it does not come into contact with the flow through.

VI. Elution

- 10. Transfer the spin column to a new 1.5ml Eppendorf tube
- 11. Add 70 μ L **Buffer AE** for elution. Incubate for 5 minutes at room temperature. Centrifuge for 1min at 8000rpm.

VII. Quantification of DNA

a. Nanodrop (spectrophotometer)

Materials Needed:

- DNA sample
- Buffer AE (for blanking)
- Lint-free tissue or Kimwipes
- Pipettes and tips

Protocol:

- Turn on the Nanodrop and select the Nucleic Acid mode.
- Blank the instrument:
- Open the pedestal and place 2 µL of buffer AE close the pedestal(gently).
- Lower the arm and press "Blank" to set the baseline.
- Wipe the pedestal clean with a lint-free tissue.

Measure the DNA sample:

- Pipette 2 μL of your DNA sample onto the lower pedestal.
- Lower the arm and press "Measure".

Record the results:

- Note the DNA concentration (ng/μL) and purity ratios (A260/A280, A260/A230).
- Clean the pedestal with a lint-free tissue before measuring the next sample.

Interpretation:

A260/A280 ratio (~1.8): Indicates pure DNA.

A260/A230 ratio (>2.0): Indicates minimal contamination.

Low ratios suggest protein, salt, or phenol contamination.

b. Qubit (fluorometer)

Materials Needed:

- Qubit dsDNA Broad Range (BR) Assay Kit (Includes buffer, dye, and standards)
- DNA sample
- Thin-walled 0.5 mL Qubit tubes
- Pipettes and tips

Protocol:

- Prepare the working solution: Mix 199 μL of Qubit buffer with 1 μL of Qubit BR dye per sample.
- Vortex briefly.
- Prepare standards and samples: Label Qubit tubes for two standards and your DNA samples.
- Pipette 190 μL of working solution into each standard tube.
- Add 10 μL of Standard #1 and Standard #2 to their respective tubes.
- For your samples, mix 198 µL of working solution with 2 µL of DNA sample in a new tube.
- Vortex all tubes briefly and incubate for 2 minutes at room temperature.

Measure on Qubit Fluorometer:

- Select dsDNA Broad Range assay mode.
- Insert Standard #1, press "Read", then insert Standard #2 and repeat.
- Insert your sample tube, press "Read", and record the concentration.

Interpret results:

- Qubit provides DNA concentration in ng/μL.
- If concentration is out of range, dilute the sample and remeasure.

VIII. Electrophoresis Gel Preparation Protocol Prepare 1.8% Agarose Gel in 100 mL

- 1. To prepare a 1.8% agarose gel in 100 mL, you'll need 1.8 grams of agarose powder.
- 2. In a flask, add 1.8 grams of agarose powder to 100 mL of 1X TAE buffer.
- 3. Heat the mixture in a microwave or on a hot plate until the agarose is completely dissolved. Swirl the flask occasionally to ensure thorough mixing.
- 4. Let is cool to about 60°C
- 5. Add 3ul of Gel red (staining dye that enables visualization under UV). **NB: Ensure safety** precautions are followed when handling ethidium bromide.
- 6. Swirl the flask to ensure thorough mixing.
- 7. Pour the agarose solution into the gel tray, ensuring that no bubbles are trapped (Gel tray is assembled prior)
- 8. Let the gel solidify for about 20-30 minutes at room temperature.
- 9. Mix DNA samples with loading dye to make them denser and more visible during electrophoresis.
- 10. Gently mix the samples and centrifuge briefly to ensure proper mixing.
- 11. Carefully remove the combs from the gel tray.
 - -Setup Electrophoresis Apparatus:
 - Fill the electrophoresis tank with 1X TAE buffer until the gel is completely submerged.
 - Carefully place the gel tray into the electrophoresis tank, ensuring that the wells are on the side with the negative (black) electrode.
- 12. Use a micropipette to load DNA samples into the wells of the gel.
- 13. Load a DNA ladder in one of the wells to serve as a size reference.
- 14. Run Electrophoresis: Connect the power supply to the electrophoresis apparatus. Set the voltage to about 100V and run the gel for approximately 45-60 minutes. Visualize the gel under a transilluminator and save photos

PCR protocol

Reagents from the Qiagen kit: AllTaq Master Mix

| Reagent | concentration | n=1 | Master mix n=10 | Mastermix | n= |
|-------------------------|---------------|-------|-------------------------------|-----------|----|
| AllTaq master mix 4X | 1X | 2.5μΙ | 25 | | |
| Primer F (10 μM) | 0.5μΜ | 0.5 | 5 | | |
| Primer R (10 μM) | 0.5μΜ | 0.5 | 5 | | |
| RNAsefree water | | 5.5 | 55 | | |
| DNA | 10ng | 1 μΙ | 1 ul @sample added separately | | |
| aliquot | | 10ul | 9 μl into each tube | | |
| | | | | | |

PCR cycle

| Step | Temperature | Time | |
|--------------|-------------|------------|----------|
| Denaturation | 95 ℃ | 2 minutes | |
| Denaturation | 95 °C | 15 seconds | |
| Annealing | 52 °C | 30 seconds | 35cycles |
| Extension | 72 °C | 1 minute | |
| Extension | 72 °C | 2 minutes | |
| Hold | 12 °C | 10 minutes | |