

PCR and Sequencing Protocols - *matK*

I. PCR protocol for *matK* marker

Note: Phusion® High-Fidelity DNA polymerase was tested on a broad range of plant taxonomic groups and selected as the enzyme with the highest performance for PCR amplification of the chloroplast barcode markers.

First choice matK primers

Forward: matK-xf	TAATTTACGATCAATTCATTC	Ford et al. 2009
Reverse: matK-MALP	ACAAGAAAGTCGAAGTAT	Dunning & Savolainen, 2010

Second choice matK primers

MatK-1RKIM-f	ACCCAGTCCATCTGGAAATCTTGTTTC	Ki-Joong Kim, pers. comm.
MatK-3FKIM-r	CGTACAGTACTTTTGTGTTTACGAG	Ki-Joong Kim, pers. comm.

PCR reagents per 10 µL reaction

# of reactions	1	100
5X HF Buffer (with MgCl ₂)	2 µL	200 µL
100% DMSO	0.3 µL	30 µL
10 mM dNTPs	0.2 µL	20 µL
10 µM Primer Forward	0.5 µL	50 µL
10 µM Primer Reverse	0.5 µL	50 µL
ddH ₂ O	5.375 µL	537.5 µL
Phusion High Fidelity Fisher Scientific #-530 (5U/ µL)	0.125 µL	12.5 µL
Total	9 µL	900 µL
DNA template	1 µL per reaction	

Recommendation: taking into account pipette error, aliquot approximately 73 µL of the PCR cocktail in each of the upper 12 wells of the plate. Using a 12-channel pipette, transfer approximately 8.7 µL of the PCR cocktail into each well. Add 1 µL of DNA. Centrifuge the plate before thermocycling.

PCR Thermocycling Program for *matK* marker

First choice matK primers:

98°C for 45 seconds;

35 cycles of 98°C for 10 seconds, 54°C for 30 seconds, 72°C for 40 seconds;

final extension 72°C for 10 minutes.



Second choice matK primers:

98°C for 45 seconds;

35 cycles of 98°C for 10 seconds, 52°C for 30 seconds, 72°C for 40 seconds;

final extension 72°C for 10 minutes.

II. Sequencing protocol for *matK* marker

Dilute PCR product adding 40 µL of water in each well. Centrifuge the plate.

For the first choice PCR use primers:

MatK-1RKIM-f	ACCCAGTCCATCTGGAAATCTTGTTTC	Ki-Joong Kim, pers. comm.
Reverse: matK-MALP	ACAAGAAAGTCGAAGTAT	Dunning & Savolainen, 2010

For the second choice use the same primers those were used for PCR

Sequencing reagents per 10 µL reaction

# of reactions	1	104
5X Sequencing Buffer	1.875 µL	195 µL
100% DMSO	0.355 µL	37 µL
10 µM primer	1 µL	104 µL
BigDye	0.250 µL	26 µL
ddH ₂ O	5.520 µL	574 µL
Total	9 µL	936 µL
Diluted DNA	2 µL per reaction	

Recommendation: taking into account pipette error, aliquot approximately 78 µL of the sequencing mix in each of the upper 12 wells of the plate. Using a 12-channeled pipette, transfer approximately 9.5 µL of the PCR cocktail in each well. Add 2 µL of DNA. Centrifuge the plate before thermocycling.

Sequencing Thermocycling Program for *matK* marker

matK Forward (matK-KIM-1R-f)

94°C for 10 seconds;

35 cycles of 94°C for 20 seconds, 48°C for 20 seconds, 60°C for 4 minutes;

hold at 4°C.

matK Reverse (matK-MALP-R)

94°C for 10 seconds;

35 cycles of 94°C for 20 seconds, 50°C for 20 seconds, 60°C for 4 minutes;

hold at 4°C.



Sequencing Cleanup

- Add Sephadex powder to the Acroprep 96 filter plate. The standard amount of powder is measured by a column loader.
- Add 300 µL of dH₂O. Let Sephadex hydrate for 2 hours at room temperature or overnight at 4°C.
- Assemble the Sephadex plate onto collection plate and centrifuge at 2100 rpm for 5 minutes.
- Immediately proceed to loading sequencing product onto the Sephadex columns to avoid drying. Use a fresh plate as a collecting plate.
- Centrifuge at 2100 rpm for 5 minutes.
- Dry the cleanup product at 88°C for 20 minutes, then cover the plate with a rubber lid and place in the freezer at -20°C until it is placed in ABI capillary sequencer.
- * The front panel of the plate should be labeled by the standard order number generated by Smithsonian GeneSifter (<https://smithsonian.genesifter.net/login>).

References

Dunning LT, Savolainen V (2010) Broad-scale amplification of matK for DNA barcoding plants, a technical note. *Botanical Journal of the Linnean Society* 164: 1–9.

Ford CS, Ayres KL, Haider N, Toomey N, van-Alpen-Stohl J, et al. (2009) Selection of candidate DNA barcoding regions for use on land plants. *Botanical Journal of the Linnean Society* 159: 1–11.