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Version 3

Pollen metabarcoding V.3

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Tomasz Suchan¹

¹Institute of Botany, Polish Academy of Sciences

Molecular Biogeography...



Tomasz Suchan

W. Szafer Institute of Botany, Polish Academy of Sciences

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Protocol status: Working

Corrected the amount of primers in the 2nd PCR reaction

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Guidelines

Perform reactions in small batches until you are confident that there is no cross-contamination among the samples. Including isolation blanks and PCR blanks is crucial for the quality control.

Materials

MATERIALS

🔯 Q5 Hot Start High-Fidelity DNA Polymerase - 100 units New England Biolabs Catalog #M0493S

Water, nuclease free

X dNTP mix (25 mM of each)

Phire Plant Direct PCR Kit Thermo Fisher Scientific Catalog #F130WH

Troubleshooting

Before start

Prepare 5 µM primer solutions:

ITS2 primers used in the 1st PCR:

ITS2-4R GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNTCCTCCGCTTATTGATATGC ITS2-S2F ACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNNATGCGATACTTGGTGTGAAT

Indexing primers used in the 2nd PCR (xxxxxxxx - index):

AATGATACGGCGACCACCGAGATCTACACxxxxxxxxACACTCTTTCCCTACACGACGC CAAGCAGAAGACGGCATACGAGATxxxxxxxxXGTGACTGGAGTTCAGACGTGTGC



Pollen extraction

Vortex the butterfly in 50 µl of water with 0.1% SDS.

Safety information

Add blank sample at this step (= "isolation blank") http://null

- 2 Evaporate water in speedvac.
- 3 Add 5 µl of the Phire Plant Direct sample buffer.
- 4 Spin max speed for 2 min.

1st PCR

- 5 Prepare the mix:
 - 14 μL molecular grade water
 - 25 μL Phire Plant Direct PCR mix
 - Δ 5 μL ITS2-S2F primer
 - Δ 5 μL ITS2-4R primer
- 6 Add 1 µl of the sample to 49 µl of the mix. Use water instead of the sample for the blanks.

Safety information

Add another blank sample at this step (= "PCR blank")

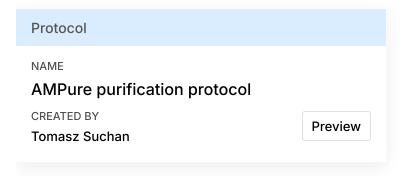
7 Run the PCR program: initial denaturation at 98°C for 5 min; 20 cycles of denaturation at 98°C for 40 s, annealing at 49°C for 40 s and elongation at 72°C for 40 s; followed by a



final extension step at 72°C for 5 min.

Purification

8 Perform AMPure purification with ratio 1x. Elute in 10 μl.



- 8.1 Remove beads from the fridge, equilibrate to the room temperature, mix thoroughly but do not vortex.
- 8.2 Add desired ratio of AMPure beads to the DNA to be purified and mix well by pipetting.
- 8.3 Incubate 5 minutes.
- 8.4 Place on the magnetic rack.
- 8.5 Let it stand for 5 minutes on the rack, aspirate and discard supernatant.
- 8.6 Add 200 μ l of freshly prepared 70% EtOH, incubate 30 seconds, aspirate and discard EtOH.
- 8.7 Repeat the wash: add 200 μ l of freshly prepared 70% EtOH, incubate 30 seconds, aspirate and discard EtOH.
- 8.8 Wait until the pellet dries out completely. DO NOT remove the plate/tubes from the magnetic rack, as the dried AMPure might pop out from the tubes!

- 8.9 Add desired volume of Tris 10 mM or water, wetting the dried AMPure pellets (add 1 µl to the final volume to avoid pipetting out the beads)
- 8.10 Remove from the magnetic rack.
- 8.11 Resuspend by pipetting or vortexing.
- 8.12 Incubate 10 minutes, incubating in 37°C can improve DNA yield.
- 8.13 Place on the magnetic rack.
- 8.14 Let it stand for 5 minutes, pippete out and save supernatant. The eluted DNA is in the supernatant, do not discard it!

2nd PCR

9 Prepare the mix:

4.82 μL molecular grade water

Δ 2 μL Q5 reaction buffer

 \triangle 0.08 µL dNTPs (25 mM each)

Δ 0.1 μL Q5 Hot Start polymerase

- 10 Add 1 μ l of the template to 7 μ l of the mix.
- 11 Add 1 µl of each 5 uM primer (forward and reverse).

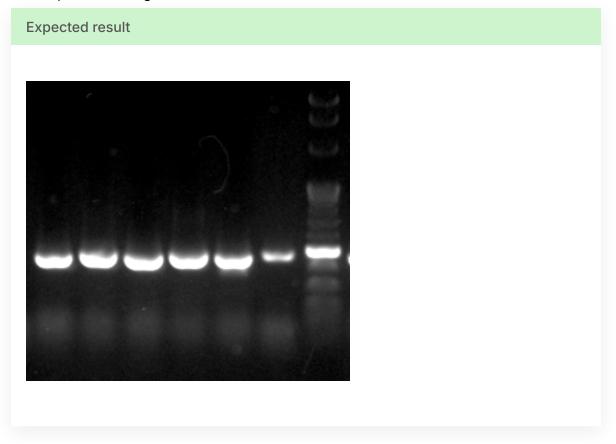
Safety information

Use different pair of indexed primers for each sample.

12 Run the PCR program: 30 s denaturation at 98°C; 12 cycles of denaturation at 98 °C for 10 s, combined annealing and extension at 72°C for 30 s (shuttle PCR); the final extension at 72°C for 5 min.

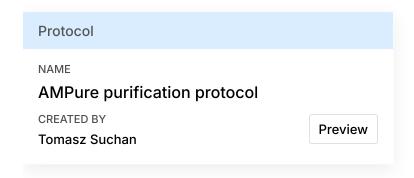


13 Check profiles on a gel.



Pooling

- 14 Pool all the samples together.
- 15 Perform AMPure purification with ratio 1x. Elute in 100 μ l.



15.1 Remove beads from the fridge, equilibrate to the room temperature, mix thoroughly but do not vortex.

- 15.2 Add desired ratio of AMPure beads to the DNA to be purified and mix well by pipetting.
- 15.3 Incubate 5 minutes.
- 15.4 Place on the magnetic rack.
- 15.5 Let it stand for 5 minutes on the rack, aspirate and discard supernatant.
- 15.6 Add 200 µl of freshly prepared 70% EtOH, incubate 30 seconds, aspirate and discard EtOH.
- 15.7 Repeat the wash: add 200 µl of freshly prepared 70% EtOH, incubate 30 seconds, aspirate and discard EtOH.
- 15.8 Wait until the pellet dries out completely. DO NOT remove the plate/tubes from the magnetic rack, as the dried AMPure might pop out from the tubes!
- 15.9 Add desired volume of Tris 10 mM or water, wetting the dried AMPure pellets (add 1 µl to the final volume to avoid pipetting out the beads)
- 15.10 Remove from the magnetic rack.
- 15.11 Resuspend by pipetting or vortexing.
- 15.12 Incubate 10 minutes, incubating in 37°C can improve DNA yield.
- 15.13 Place on the magnetic rack.
- 15.14 Let it stand for 5 minutes, pippete out and save supernatant. The eluted DNA is in the supernatant, do not discard it!



16 Check the concentration using Qubit and the profile using Tapestation/Fragment Analyzer. Calculate molarity from that and proceed to the sequencing. Add 15% PhiX to the sequencing run.