



Dec 08, 2023

Version 1

HMW DNA extraction protocol for ferns V.1

DOI

<https://dx.doi.org/10.17504/protocols.io.bp2l69kb1lqe/v1>



Geferson Fernando Metz¹, Rafael Plá Matielo Lemos², Cristiane Barbosa D'Oliveira Matielo¹, Tiego Ferreira¹, Filipe Victoria¹

¹Núcleo de Estudos da Vegetação Antártica - NEVA; ²Universidade Federal do Pampa



Rafael Plá Matielo Lemos

Universidade Federal do Pampa, Núcleo de Estudos da Vegetaçã...

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.bp2l69kb1lqe/v1>

Protocol Citation: Geferson Fernando Metz, Rafael Plá Matielo Lemos, Cristiane Barbosa D'Oliveira Matielo, Tiego Ferreira, Filipe Victoria 2023. HMW DNA extraction protocol for ferns. **protocols.io**

<https://dx.doi.org/10.17504/protocols.io.bp2l69kb1lqe/v1>

Manuscript citation:

Geferson Fernando Metz, Tiego Ferreira, Cristiane Barbosa D'Oliveira Matielo, Rafael Plá Matielo Lemos, Filipe de Carvalho Victoria. HMW DNA extraction protocol for ferns. protocols.io

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

Protocol status: Working

We use this protocol and it's working

Created: August 25, 2022

Last Modified: December 08, 2023

Protocol Integer ID: 69196

Keywords: Ferns, Nanopore, HMW, long reads, Plant DNA, hmw dna extraction protocol for fern, disinfecting fern spore, fern spore, disinfection protocol for spore, hmw dna extraction protocol, dna extraction, fern, spore, read sequencing, germinated plant, disinfection protocol, large amounts of dna, sequencing, dna, dna in quantity, plant, extraction, growth free of contamination, extracting large amount, contamination, plants in the axenic state

Funders Acknowledgements:

Auxílio a Pesquisa GENO-ISLAND: Adaptações moleculares das plantas aos ambientes insulares

Grant ID: CNPQ 443237/2019-0

Coordenação de Aperfeiçoamento de Pessoal de Nível Superior - Brasil

Grant ID: CAPES 001

Abstract

Among the difficulties encountered in a laboratory, simple situations such as efficiently disinfecting fern spores and extracting large amounts of DNA with low tissue input in the protocol are one of the major impediments in carrying out sequencing of these plants in the axenic state. The objective of this work is to present a replicable, scalable and easy-to-execute protocol for work with ferns following the current generation of long-read sequencing. Our method is based on providing a disinfection protocol for spores and sporangia that guarantees growth free of contamination and, after this growth, DNA extraction using a low amount of material in order to obtain a good yield. The results are promising since, in up to 21 days, we obtained germinated plants. After their growth (in average 180 days) we were able to extract DNA in quantity and quality and perform the sequencing, emphasizing that our best N50 is 24 Kb.

Image Attribution

Illustration by Crix D'Oliveira.

Materials

-  Liquid nitrogen
-  Chloroform: Isoamyl Alcohol (24:1)
-  Isopropanol
-  TE Buffer
-  3M sodium acetate
-  Ethanol 100%
-  nuclease free water
-  Sodium Hypochlorite Solution
-  70% Ethanol

β -mercaptoethanol

PVP 40

TRIS 1 M pH 8.0

EDTA 0.5 M pH 8

Sodium Metabisulfite

NaCl 2.5M

Sodium dodecyl sulfate (SDS) 20%

2 ml LoBind tubes

1.5 ml LoBind tubes

Mortar & pestle

Water Bath

Centrifuge

Protocol materials

-  Sodium Hypochlorite Solution
-  Liquid nitrogen
-  Chloroform: Isoamyl Alcohol (24:1)
-  TE Buffer
-  Ethanol 100%
-  nuclease free water
-  Isopropanol
-  3M sodium acetate
-  70% Ethanol

Safety warnings



Safety information

Work under fume hood when add β -mercaptoethanol and Chloroform.

Safety information

Be careful when handling liquid nitrogen.



Before start

- Prepare the SDS Lysis Buffer on the day of the experiment;

	A	B	C
	Reagent	Reagent Stock Concentration	FINAL
	PVP 40	100%	1%
	Sodium Metabisulfite	1%	1%
	NaCl	5 M	0,5 M
	TRIS HCL pH 8	1 M	100 mM
	EDTA pH 8	0.5 M	50 mM
	DDH ₂ O	-	~
	Sodium dodecyl sulfate (SDS)	20%	1,5%
	β-MERCAPTOETANOL	-	2% (v/v)

- When preparing the SDS Lysis Buffer, add SDS at last this will avoiding bubble formation.
- Preheat the water bath; Keep the SDS Lysis Buffer at 65°C until the tissue powder is added.
- Washing solution (EtOH 70%), fresh
35 ml Ethanol 100% + 15 ml H₂O
- Potassium Acetate 5M
Dissolve 4.9 gr KAc in 10 ml ddH₂O (4.9 gr KAc + ≈ 7.5 ml H₂O).
Adjust the pH with glacial acetic acid.
- Prepare TE buffer (10 mM Tris pH 8 and 1 mM EDTA pH 8)

Plant material sterilization

3d 0h 45m

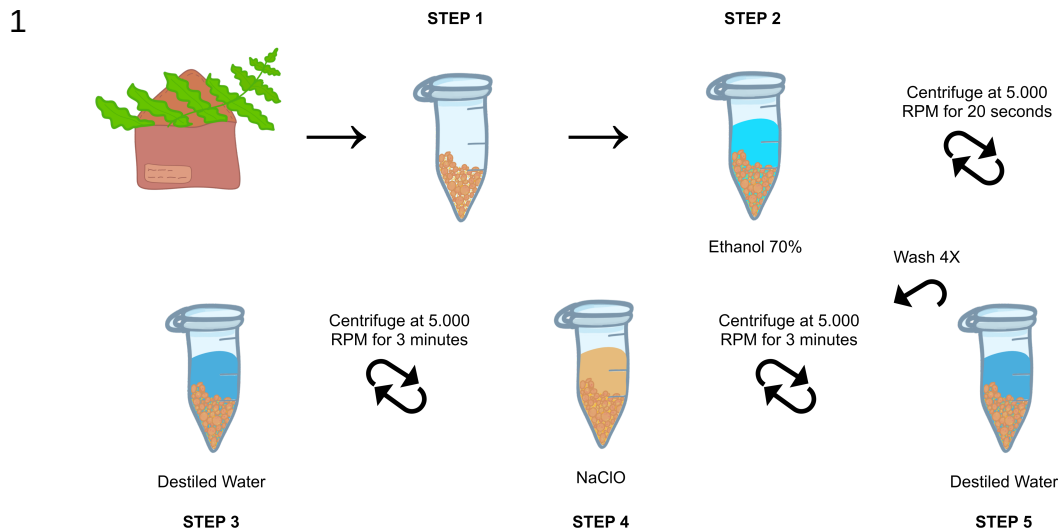


Figure 1. Step by step plant spore and sporangia sterilization. Illustration by Crix D'Oliveira.

- 2 Sample and store the leaf tissue (fronds) in paper envelopes for three days to induce dehiscence.
- 3 Recover and store the spores together with sporangia in 1.5 mL microtubes until one-third of the tube was filled and then stored at $-20\text{ }^{\circ}\text{C}$ until disinfection.
- 4 Add 1 mL $\text{70\% Ethanol}$, homogenize by inversion for $00:00:20$ and briefly centrifuge them. 20s
- 5 Discard the supernatant and wash by inversion with 1 mL of autoclaved purified water, briefly centrifuge and discard the supernatant. 2m
- 6 Add 1 mL of an $\text{Sodium Hypochlorite Solution}$ (active chlorine $\sim 2\%$) at $10\text{ \% (v/v)}$, homogenize by inversion for $00:20:00$, centrifuge at 5.000 rpm $00:03:00$. 23m

- 7 Discard the supernatant and wash by inversion with 1 mL of autoclaved purified water, centrifuge at 5 5.000 rpm 00:03:00 . Repeat this step 4 times. 3m
- 8 Add 1 mL autoclaved purified water, homogenize and pipete into a Petri Dish with BCD medium (MgSO₄·7H₂O - 0.1 mM, KH₂PO₄ - 1.84 mM, KNO₃ - 1M, FeSO₄·7H₂O - 4.5mM). 1m

Extraction of high-molecular-weight DNA

2h 31m

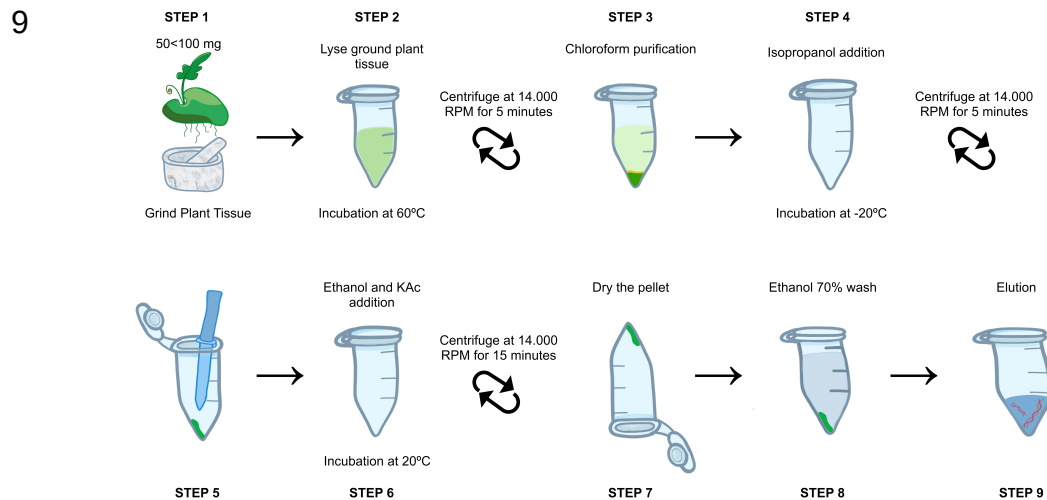


Figure 2. Step by step extraction of high-molecular-weight DNA illustrated protocol. Illustration by Crix D'Oliveira.

- 10 Weigh between 50 mg and 100 mg of leaf tissue. Grind must be done with a crucible and pestle (previously exposed to Liquid nitrogen) until a very fine powder is obtained. 7m
- 11 Pre-heat the Lysis Buffer and aliquot 600 µL for each sample individually in 1.5 mL microtube. With the aid of a spoon or spatula, transfer the macerate to the microtube. 1m
- 12 Homogenize by inversion 10 times and incubate at 60 °C for 00:10:00 . 10m
- 13 Add 4 µL of RNase A (100 mg/mL); add 4 µL of proteinase K (> 40 U/mg);



- 14 Homogenize by inversion 10 times and incubate at 60 °C for 00:20:00 gently homogenizing by inversion every 00:05:00 . 25m
- 15 Add 600 µL of Chloroform: Isoamyl Alcohol (24:1) and homogenize by inversion for at least 00:03:00 until an off-white emulsion forms. 3m
- 16 Centrifuge at 14.000 rpm for 00:05:00 at room temperature. 5m
- 17 Carefully aspirate the upper phase of the tube and transfer to a new 1.5 mL microtube. 1m
- 18 Add 400 µL of Isopropanol -20 °C and incubate it for at least 01:00:00 at -20 °C . (Can be stored overnight) 1h
- 19 Centrifuge at 14.000 rpm for 00:05:00 at room temperature and discard the supernatant. 5m
- 20 Add 50 µL of TE Buffer . Add 5 µL of 3M sodium acetate . 1m
- 21 Add 120 µL of iced Ethanol 100% (-20 °C) and mix well by flicking the tube.
- 22 Store tubes at -20 °C for at least 00:20:00 . 20m
- 23 Centrifuge at 14.000 rpm for 00:15:00 at room temperature and carefully discard the supernatant. 15m
- 24 Add 1 mL of freshly prepared 70% Ethanol , mix gently by inversion, centrifuge briefly and carefully discard the supernatant. 1m



- 25  50 μL Dry the pellet for approximately  00:10:00 at Room Temperature on the bench with the tube upside down. 10m
- 26 Elute in  50 μL volume in  nuclease free water . (Elution volume can be adjusted for your needs) 1m

Quality Assessment

- 27 Quantify the DNA on a Qubit® fluorometer (dsDNA high sensitivity assay). DNA yield can be 500 - 1500 ng.
- 28 Check DNA integrity though electrophoresis in a 1% Agarose Gel.

DNA Size Selection

- 29 Remove short DNA fragments with Circulomics® Short-Read Eliminator Kit.