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Extracellular DNA extraction from sediment using phosphate buffer and NucleoSpin® Soil kit (MACHEREY NAGEL)

 Forked from [DNA extraction from environmental biofilm using the NucleoSpin® Soil kit \(MACHEREY-NAGEL\)](#)

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Manuscript citation:

Taberlet, P., Prud'Homme, S. M., Campione, E., Roy, J., Miquel, C., Shehzad, W., ... Coissac, E. (2012). Soil sampling and isolation of extracellular DNA from large amount of starting material suitable for metabarcoding studies. *Molecular Ecology*, 21, 1816– 1820.

<https://doi.org/10.1111/j.1365-294X.2011.05317.x>

Giguët-Covex, C., Ficetola, G.F., Walsh, K. et al. New insights on lake sediment DNA from the catchment: importance of taphonomic and analytical issues on the record quality. *Sci Rep* 9, 14676 (2019). <https://doi.org/10.1038/s41598-019-50339-1>

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Protocol status: Working

We use this protocol and it's working

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Protocol Integer ID: 35410

Guidelines

- Sample preparation
- Desorption of DNA from mineral and organic particles
- Contaminants elimination
- DNA fixation and washing
- DNA elution

Materials

- Samples
 - sediment \approx 0.75g
- Reagents
 - NucleoSpin® Soil kit (MACHEREY-NAGEL)
 - Ethanol (96 - 100%), molecular grade to prepare buffer SW2
 - Saturated phosphate buffer (0.12 M Na₂HPO₄; pH \approx 8), stored less than one week at +4°C
 - Ethanol to sterilize spatula
- Materials
 - specific DNA-work station (sterile area equipped with air filtration and UV systems)
 - microcentrifuge for 1.5 to 2 mL tubes (relative centrifugal force needed: 11,000 to 14,000 x g)
 - agitator for rotation with 2mL tubes holder
 - water bath
 - spatula
 - metallic pincer
 - digital burner
 - metal or glass support
 - clean chisel wash with DNA off or ethanol
 - precision weighting scale, precision : 0.0001g
 - pipettes : 1000 μ L - 100 μ L
 - 2 trash cans : 1 for liquid and 1 for solid
- Consumables
 - tips with filter :
 - > 1000 μ L > 12 tips per sample
 - > 100 μ L > 1 tip per sample
 - 2 mL sterile microcentrifuge tube > 4 per sample
(1 to transfert sediment - step 1; 1 to collect supernatant - step 3; 1 to collect solution after filter lysate - step 5; and 1 to elute DNA - step 10)
 - parafilm
 - gloves

Safety warnings

- ❗ The manufacturer advise to wear gloves and goggles and to flow the safety instructions for 2 reagents :
 - SB contains Guanidinium thiocyanate 45 - 60%,
CAS number : CAS 593-84-0
Signal word : Irritant
Hazard phrases : 302, 412
Precaution phrases : 264W, 273, 301+312, 330
 - SW1 contains Guanidine hydrochloride 36 - 50% and 2 - propanol 20 - 35%
CAS number : CAS 50-01-1, 67-63-0
Signal word : Irritant and Flammable
Hazard phrases : 226, 302, 319, 336
Precaution phrases : 210, 260D, 264W, 280sh, 301+312, 330

Before start

- The following precautions must be applied :
 - If possible, it is best to work in a room dedicated to rare and ancient DNA
 - Wear gloves throughout the extraction process
 - Clean the bench with DNA off
 - Use tips with filters to avoid contaminations
 - Include negative controls
 - All steps have to be performed under a specific DNA-work station (sterile area equipped with air filtration and UV systems).

- Materials preparation :
 - Clean a specific DNA - work station and apply the UV for 15min
 - Sterilize spatulas, one spatula per sample. To limit risk of burns, during this step, you must wear a cotton blouse and do not use gloves
 - * put a clean spatula in an ethanol solution
 - * with metal pincer, take the spatula
 - * pass the spatula through the flame, warning of the risk of burns
 - * place sterilize spatula in a clean metal or glass support

- Samples preparation :
 - thaw sediment used for extraction, between 30min - 1h at room temperature or between 1h - 2h at +4°C (the time depends of the quantity of sediment and freezing temperature)

- Solutions preparation :
 - Prepare saturated phosphate buffer (0.12 M Na₂HPO₄; pH ≈ 8)(according to Taberlet et al 2012)
 - * weigh the 2 compounds - see table below
 - * transfer these compounds in a becher
 - * dissolve them in ultra pure water
 - * check pH, it will be ≈8
 - * put solution into 1L graduated flask, make up to the mark with ultra pure water and mix
 - * filter this solution on 0.2μM filter
 - * store this buffer at +4°C, maximum one week

	Name	Linear Formula	Molecular Weight	Final concentration	Weight for 1L of buffer
	Sodium phosphate monobasic	NaH ₂ PO ₄	119.98 g/mol	16.4mM	1.97g
	Sodium phosphate dibasic	Na ₂ HPO ₄	141.96 g/mol	103.6mM	14.7g



Informations on the compounds of saturated phosphate buffer

- Check buffer SW2 - before the first utilisation, you need to add the indicate volume of ethanol (96 - 100%) to buffer SW2 concentrate and mark the label of the bottle to indicate that ethanol was added. This solution is stable at room temperature (18 - 25°C) for at least one year
- Optional : Incubate the elution buffer SE at +50°C

Prepare the sample

1

- Annotate 2mL tubes, one 2mL tube per sample
- Weigh annotated tubes, use the weigh scale (0.0001g of precision)
 - On your notebook, note the name of tubes and their weights
- With a sterilized spatula, homogenize a sediment and check that it is completely thawed (if not, wait for total defrost)
- Transfer $\approx 0.5\text{g}$ of sediment in the associated tube, 2 methods :
 - if the sediment is "compact", use a sterilized spatula to transfer the sediment
 - if the sediment contains a lot of water, use a 1000 μL tip cutted with clean chisel and transfer sediment by pipetting few times.

→ On your notebook, note the used method and your comments about the sediment particularity

- Weigh the tube containing sediment
 - On your notebook, note the weight
- On the storage tube, make a cross to show that this tube contains thawed sediment thaw once : this is highly

recommended to do not subject sediment to freeze-thaw cycles in particular if the aim is to perform downstream analyses on sed-DNA extracts.

Keep this sediment at -40°C or -80°C for long term storage.

→ On your notebook, note where the sediments are stored



- Repeat this process for all the sediments that will be extracted during this session
- Add one volume of saturated phosphate buffer (0.12 M Na_2HPO_4 ; $\text{pH} \approx 8$; stored less than one week at $+4^{\circ}\text{C}$) for one volume of sediment

e.g if you have weighed 0.5g of sediment, add 0.5mL of saturated phosphate buffer

- Close the cap and you can add parafilm to secure the closure


Desorb DNA from mineral and organic particlesAdjust lysis conditions

2





- Attach 2mL tubes (containing sediment and P buffer) to agitator for rotation
- Agitate tubes at slow speed at  Room temperature for  00:15:00

Precipitate contaminants

3

- Centrifuge  10000 rpm, 00:10:00
- Transfer the supernatant (without taking the pellet) to a new 2mL tube



4

- Add  150 μL of **buffer SL3** and vortex for  00:00:05
- Incubate at  4°C in a fridge for  00:05:00



- Centrifuge  11000 x g, 00:01:00

Filter lysate



- 5
 - Place a NucleoSpin® Inhibitor Removal Column (red ring) in a Collection Tube (2mL, lid)
 - Load up to  650 µL of clear supernatant (obtained at the step 4) onto the filter
 - Centrifuge  11000 x g, 00:01:00
 - Repeat the load and the centrifuge step as many time as there is still some supernatant from step 4 to be filtered

After each centrifugation, collect the filtered liquid in a clean tube : 1 single tube for all the filtration




- Discard the NucleoSpin® Inhibitor Removal Column

Note : if a pellet is visible after the centrifugation, transfer the clear supernatant to a new collection tube (not provided in the kit) to get ride of this pellet, and continue with the clear supernatant

Adjust binding conditions

- 6
 - Add  250 µL of **buffer SB**
 - Close the lid
 - Vortex for  00:00:05 , make a brief centrifugation

Bind DNA

- 7
 - Place a NucleoSpin® Soil Column (green ring) in a collection Tube (2mL)
 - Load  550 µL of sample onto the column
 - Centrifuge  11000 x g, 00:01:00
 - Discard the flow through and place the column back into the collection tube
 - Load the remaining sample onto the column
 - Centrifuge  11000 x g, 00:01:00
 - Discard the flow through and place the column back into the collection tube


Wash and dry silica membrane

- 8
 - *Note : the same collection tube is used throughout the entire washing procedure to reduce plastic waste*



8.1 1st wash :

- Add  500 µL of **buffer SB** to the NucleoSpin® Soil Column



- Centrifuge  11000 x g, 00:00:30
- Discard the flow through and place the column back into the collection tube


8.2 2nd wash :

- Add  550 µL of **buffer SW1** to the NucleoSpin® Soil Column
- Centrifuge  11000 x g, 00:00:30
- Discard the flow through and place the column back into the collection tube



8.3 3rd wash :

- Add  650 µL of **buffer SW2** to the NucleoSpin® Soil Column

Note : verify that ethanol was added to buffer SW2 during the first utilisation

- Centrifuge  11000 x g, 00:00:30
- Discard the flow through and place the column back into the collection tube

8.4 4th wash :

- Add  650 µL of **buffer SW2** to the NucleoSpin® Soil Column
- Centrifuge  11000 x g, 00:00:30
- Discard the flow through and place the column back into the collection tube

Dry silica membrane


- 9 Centrifuge  11000 x g, 00:02:00




Note : if for any reason, the liquid in the collection tube has touched the NucleoSpin® Soil Column after drying step, discard flow through and centrifuge again

Elute DNA

- 10
- Place the NucleoSpin® Soil Column into a new microcentrifuge tube (not provided in the kit)

- Add  30 µL of **buffer SE** to the column

Optional : to increase yield you can heat buffer SE at  50 °C

- Do not close the lid and incubate at  Room temperature for  00:01:30
- Close the lid and centrifuge  11000 x g, 00:00:30
- Discard the NucleoSpin® Soil Column and keep the tube containing the DNA
- We recommend storing DNA frozen at -20°C until preparation of DNA library for HTS (or at -40°C to -80°C for longer storage)
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