



VERSION 2

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**Protocol status:** Working  
We use this protocol and it's working

## Inhibitor-free DNA extraction from soil and sediment samples V.2

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### ABSTRACT

This protocol describes how to extract inhibitor-free DNA from soil and sediment samples. 5 g of soil or up to 10 g of sediment can be processed in one extraction, but there is also a miniaturized version for 250 mg of input material, if less DNA is required. The protocol is based on the DNeasy PowerMax Soil Kit but costs much less. A lot of the buffers can be found in the following patent <https://patents.google.com/patent/US7459548B2/en>

### GUIDELINES

Follow general lab etiquette. Wear gloves to prevent contaminating the samples. Clean the workspace before starting with 80% EtOH.

### MATERIALS

#### Materials required:

Below all materials needed for the protocol are listed. Vendors and part numbers are listed but interchangeable depending on the supply situation.

#### Chemicals:

Sodium phosphate dibasic

Sodium phosphate dibasic Merck MilliporeSigma (Sigma-Aldrich) Catalog #S0876-100G

Guanidinium thiocyanate

Guanidinium thiocyanate Fisher Scientific Catalog #10503345

Sodium phosphate monobasic Sodium phosphate monobasic

Sodium phosphate monobasic Merck MilliporeSigma (Sigma-Aldrich) Catalog #S0751-100G

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91569

SDS ultrapure

✕ Sodium dodecyl sulfate Diagonal Catalog #A1112.0500

Sodium chloride

✕ Sodium chloride Fisher Scientific Catalog #10616082

Tris ultrapure 99.9%

✕ Tris ultrapure 99.9% Diagonal Catalog #A1086.1000

Hydrochloric acid fuming 37%

✕ Hydrochloric acid fuming 37% Merck MilliporeSigma (Sigma-Aldrich) Catalog #1003171011

Ammonium acetate

✕ Ammonium acetate Carl Roth Catalog #7869.2

Aluminium ammonium sulfate dodecahydrate

✕ Aluminium ammonium sulfate dodecahydrate Merck MilliporeSigma (Sigma-Aldrich) Catalog #A2140-500G

Guanidine hydrochloride

✕ Guanidine hydrochloride Fisher Scientific Catalog #10543325

Isopropanol

✕ 2-Propanol Carl Roth Catalog #1HPK.1

Tween 20

✕ Tween 20 Carl Roth Catalog #9127.1

Acetic acid

✕ Acetic acid Carl Roth Catalog #7332.1

Ethanol absolute

✕ Ethanol absolute 99.8% p.a. Carl Roth Catalog #9065.1

#### Labware:

50 mL centrifuge tubes, Ultra-High Performance

✕ Centrifuge tubes Ultra-High Performance VWR International Catalog #525-1098

Garnet Sharp Particles

✕ Garnet Sharp Particles BioSpec Products Catalog #11079103gar

✕ Vortex Adapter for 2 (50 ml) tubes Qiagen Catalog #13000-V1-50

EconoSpin Maxi Spin column

✕ EconoSpin® DNA Only Maxi Spin Column Epoch Life Science Catalog #2040-050

2 mL screwcap tubes

✕ 2 mL screwcap tube Sarstedt Catalog #72.693

The EconoSpin® All-In-One DNA Only Mini Spin Column



### Stock solutions:

🧪 1 L SDS stock solution [M] 10 Mass / % volume

- Add 🧪 100 g SDS ultrapure to a beaker
- Adjust volume to 🧪 1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡️ Room temperature

🧪 1 L sodium chloride stock solution [M] 5 Molarity (M)

- Add 🧪 292.2 g sodium chloride to a beaker
- Adjust volume to 🧪 1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡️ Room temperature

🧪 1 L Tris stock solution [M] 1 Molarity (M) 📏 pH 8

- Add 🧪 121.14 g Tris ultrapure 99.9% to a beaker
- Adjust volume to 🧪 800 mL with ddH<sub>2</sub>O
- Adjust pH to 📏 pH 8 with HCl
- Adjust volume to 🧪 1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡️ Room temperature

🧪 500 mL sodium acetate stock solution [M] 3 Molarity (m) 📏 pH 5

- Add 🧪 123 g sodium acetate to a beaker
- Adjust volume to 🧪 400 mL with ddH<sub>2</sub>O
- Adjust pH to 📏 pH 5 with acetic acid
- Adjust volume to 🧪 500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡️ Room temperature

🧪 1 L Tris stock solution [M] 1 Molarity (m) 📏 pH 7.5

- Add 🧪 121.14 g Tris ultrapure 99.9% to a beaker
- Adjust volume to 🧪 800 mL with ddH<sub>2</sub>O
- Adjust pH to 📏 pH 7.5 with HCl
- Adjust volume to 🧪 1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡️ Room temperature

[M] 1 Molarity (m)  $\text{pH}$  8.5

- Add  $\text{flask}$  121.14 g Tris ultrapure 99.9% to a beaker
- Adjust volume to  $\text{flask}$  800 mL with ddH<sub>2</sub>O
- Adjust pH to  $\text{pH}$  8.5 with HCl
- Adjust volume to  $\text{flask}$  1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at  $\text{flame}$  Room temperature

$\text{flask}$  1 L wash buffer stock solution ( [M] 50 millimolar (mM) Tris )  $\text{pH}$  7.5

- Add  $\text{flask}$  50 mL Tris stock solution  $\text{pH}$  7.5 to a beaker
- Adjust volume to  $\text{flask}$  1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at  $\text{flame}$  Room temperature

### Working solutions:

$\text{flask}$  500 mL bead-beating solution ( [M] 180 millimolar (mM) sodium phosphate ,  
[M] 120 millimolar (mM) guanidinium thiocyanate )  $\text{pH}$  8

- Add  $\text{flask}$  12.8 g sodium phosphate dibasic to a beaker
- Add  $\text{flask}$  7.1 g guanidinium thiocyanate
- Adjust volume to  $\text{flask}$  490 mL with ddH<sub>2</sub>O
- Adjust pH to  $\text{pH}$  8 by adding sodium phosphate monobasic
- Adjust volume to  $\text{flask}$  500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at  $\text{flame}$  Room temperature

$\text{flask}$  500 mL lysis solution ( [M] 150 millimolar (mM) sodium chloride ,  
[M] 4 Mass / % volume SDS , [M] 500 millimolar (mM) Tris )  $\text{pH}$  8

- Add  $\text{flask}$  200 mL of [M] 10 Mass / % volume SDS stock solution to a beaker
- Add  $\text{flask}$  15 mL of [M] 5 Molarity (m) sodium chloride stock solution
- Add  $\text{flask}$  250 mL of [M] 1 Molarity (m) Tris stock solution  $\text{pH}$  8
- Adjust volume to  $\text{flask}$  500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at  $\text{flame}$  Room temperature

$\text{flask}$  500 mL ammonium acetate buffer ( [M] 130 millimolar (mM) ammonium acetate )

- Add  $\text{flask}$  5 g ammonium acetate to a beaker
- Adjust volume to  $\text{flask}$  500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at  $\text{flame}$  Room temperature

🧪 500 mL inhibitor removal solution (

[M] 120 millimolar (mM) aluminum ammonium sulfate dodecahydrate )

- Add 🧪 27.2 g aluminium ammonium sulfate dodecahydrate to a beaker
- Adjust volume to 🧪 500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡 Room temperature

🧪 500 mL DNA binding buffer ( [M] 5 Molarity (M) Guanidine hydrochloride ,

[M] 40 % (v/v) isopropanol , [M] 0.05 % (v/v) Tween 20 ,

[M] 115 millimolar (mM) sodium acetate ) 📏 pH 5

- Add 🧪 238.8 g guanidine hydrochloride to a beaker
- Add 🧪 200 mL isopropanol
- Add 🧪 250 µL Tween 20
- Add 🧪 20 mL [M] 3 Molarity (m) sodium acetate stock solution 📏 pH 5
- Adjust volume to 🧪 500 mL with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡 Room temperature

🧪 1 L wash buffer ( [M] 10 millimolar (mM) Tris , [M] 80 % (v/v) Ethanol )

📏 pH 7.5

- Add 🧪 200 mL was buffer stock solution
- Adjust volume to 🧪 1 L with Ethanol absolute
- Sterilize by filtering and store at 🌡 Room temperature

🧪 1 L elution buffer ( [M] 10 millimolar (mM) Tris ) 📏 pH 8.5

- Add 🧪 10 mL Tris stock solution 📏 pH 8.5 to a beaker
- Adjust volume to 🧪 1 L with ddH<sub>2</sub>O
- Sterilize by filtering and store at 🌡 Room temperature

## SAFETY WARNINGS

- ⚠ Buffers containing guanidine produce highly reactive compounds when mixed with bleach. Don't mix the extraction waste with bleach or solutions that contain bleach.  
Reagents are potentially damaging to the environment. Dispose waste as mandated.

## BEFORE START INSTRUCTIONS

Make sure all buffers are prepared before starting.

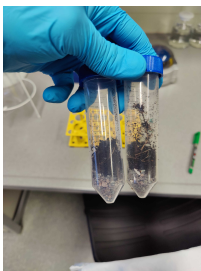
### Protocol for up to 10 g of input material

20m 30s

- 1 Prepare one 50 mL centrifuge tube per sample with 15 g of garnet beads.





- 2 Add up to 10 g of soil to the tube.



### Note



The amount of starting material differs from soil type to soil type. For most soil types 2 g of input material is sufficient. If the output is too low with 2 g it can be increased step by step.




- 3 Add  15 mL bead-beating solution and  1.2 mL lysis solution . Vortex shortly.

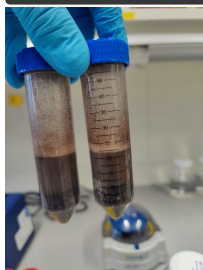
- 4 Place the samples on a Vortex adapter (e.g. Qiagen) and vortex at maximum speed for  00:10:00  10m



### Note




If you want to process more samples, instead of the vortex adapter a Thermoblock can be used. As an alternative, you can incubate the sample for  00:30:00 at  65 °C and at maximum RPM.


- 5  2500 x g, 20°C,  00:03:00 . Transfer the supernatant to a new tube.  3m

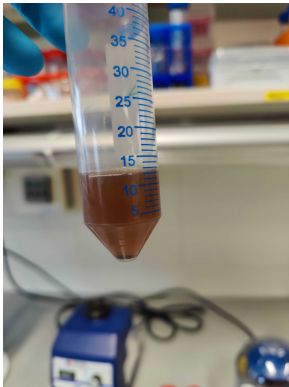


## Note

For the large volume protocol, the samples can be carefully poured instead of being pipetted.


6 Add  5 mL ammonium acetate buffer , vortex shortly, and incubate at  4 °C for  00:10:00 10m

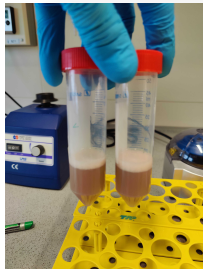
7  2500 x g, 20°C, 00:04:00 . Transfer the supernatant to a new tube avoiding the pellet. The solution may still be colored, depending on the input material. 4m






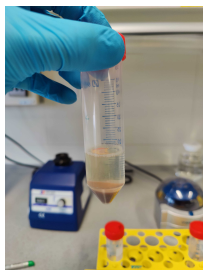
8 Add  4 mL of inhibitor removal buffer . A precipitate may form. Vortex shortly, incubate at  4 °C 10m




for  00:10:00 .

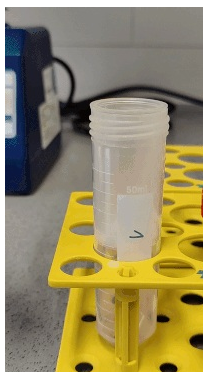









- 9  2500 x g, 20°C, 00:04:00 . The solution will clear up. Avoiding the pellet, transfer up to  15 mL  a new tube.



- 10 Add  30 mL DNA binding buffer . Vortex or invert to mix.





- 11 Add the mixture to a maxi spin column (e.g. Epoch Life Science) in a 50 mL centrifuge tube.





- 12  2500 x g, 20°C, 00:00:30 . Discard the flow-through. Repeat once to bind the complete sample volume. 30s
- 13 Add  10 mL wash buffer .  2500 x g, 20°C, 00:05:00 to wash and dry the column. 5m
- 14 Transfer the column to a new tube. Add  1 mL elution buffer . Incubate for  00:03:00 at  Room temperature . 3m
- 15  2500 x g, 20°C, 00:01:00 to elute the DNA. DNA eluate should be completely colorless and ready go for downstream analysis. 1m

## Protocol for up to 250 mg of input material









50m

- 16 Prepare one 2 mL centrifuge tube per sample with 750 mg of garnet beads.
- 17 Add  250 mg of soil or sediment sample.
- 18 Add  750 µL bead-beating solution and  60 µL lysis solution . Vortex shortly.
- 19 Place the samples on a Vortex adapter (e.g. Qiagen) and vortex at maximum speed for  00:10:00 10m

### Note

If you want to process more samples, instead of the vortex adapter a Thermoblock can be used. As an alternative, you can incubate the sample for  00:30:00 at  65 °C and at maximum RPM.

- 20  10000 x g, 20°C, 00:03:00 . Transfer the supernatant to a new tube. 3m
- 21 Add  250 µL ammonium acetate buffer , vortex shortly, and incubate at  4 °C for  00:05:00 5m
- 22  10000 x g, 20°C, 00:01:00 . Transfer the supernatant to a new tube. 1m
- 23 Add  200 µL of inhibitor removal buffer . A precipitate may form. Vortex shortly, incubate at  4 °C 5m for  00:05:00 .
- 24  10000 x g, 20°C, 00:01:00 . Transfer  600 µL of the supernatant to a new tube. 1m
- 25 Add  1200 µL DNA binding buffer . Vortex to mix.
- 26 Load  650 µL of the mixture to a mini spin column (e.g. Epoch Life Science).

- 27  10000 x g, 20°C, 00:00:30 . Discard the flow-through. Repeat two times to bind the complete sample volume. 30s
- 28 Add  500 µL wash buffer .  10000 x g, 20°C, 00:00:30 to wash the column. Discard the flow-through. 30s
- 29  10000 x g, 20°C, 00:01:00 to dry the column. Transfer the spin column to a clean 1.5 mL microcentrifuge tube. 1m
- 30 Add  50 µL elution buffer . Incubate for  00:03:00 at  Room temperature . 3m
- 31  10000 x g, 20°C, 00:01:00 to elute the DNA. DNA eluate should be completely colorless and ready go for downstream analysis. 1m