

Sep 28, 2021

DNA Extraction from Sterivex Filters

Prorked from DNA Extraction from Sterivex Filters

Christopher Neil Thornton¹, William Brazelton¹

¹University of Utah



dx.doi.org/10.17504/protocols.io.bykqpuvw

Brazelton Lab

William Brazelton

DISCLAIMER

DISCLAIMER - FOR INFORMATIONAL PURPOSES ONLY; USE AT YOUR OWN RISK

The protocol content here is for informational purposes only and does not constitute legal, medical, clinical, or safety advice, or otherwise; content added to protocols.io is not peer reviewed and may not have undergone a formal approval of any kind. Information presented in this protocol should not substitute for independent professional judgment, advice, diagnosis, or treatment. Any action you take or refrain from taking using or relying upon the information presented here is strictly at your own risk. You agree that neither the Company nor any of the authors, contributors, administrators, or anyone else associated with protocols.io, can be held responsible for your use of the information contained in or linked to this protocol or any of our Sites/Apps and Services.

ABSTRACT

Modified 2015 by the Brazelton Lab from protocols by Rika Anderson, Colleen Kellogg, Julie Huber, and Byron Crump. Incorporated some recommendations from Lever et al. (2015) Frontiers in Microbiology doi: 10.3389/fmicb.2015.00476.

DOI

dx.doi.org/10.17504/protocols.io.bykqpuvw

PROTOCOL CITATION

Christopher Neil Thornton, William Brazelton 2021. DNA Extraction from Sterivex Filters. **protocols.io** https://dx.doi.org/10.17504/protocols.io.bykqpuvw

FORK NOTE

FORK FROM

Forked from DNA Extraction from Sterivex Filters, Gigascience Database

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

CREATED

Sep 28, 2021

LAST MODIFIED

Sep 28, 2021

Citation: Christopher Neil Thornton, William Brazelton (09/28/2021). DNA Extraction from Sterivex Filters. https://dx.doi.org/10.17504/protocols.io.bykqpuvw

DISCLAIMER:

DISCLAIMER - FOR INFORMATIONAL PURPOSES ONLY; USE AT YOUR OWN RISK

The protocol content here is for informational purposes only and does not constitute legal, medical, clinical, or safety advice, or otherwise; content added to protocols.io is not peer reviewed and may not have undergone a formal approval of any kind. Information presented in this protocol should not substitute for independent professional judgment, advice, diagnosis, or treatment. Any action you take or refrain from taking using or relying upon the information presented here is strictly at your own risk. You agree that neither the Company nor any of the authors, contributors, administrators, or anyone else associated with protocols.io, can be held responsible for your use of the information contained in or linked to this protocol or any of our Sites/Apps and Services.

Prepare DEB

1 Prepare DNA Extraction Buffer (DEB):

0.1M Tris-HCl (pH 8) 4.5 mL of 1.0 M 0.1M Na-EDTA (pH 8) 9 mL of 0.5M

0.1M KH2PO4 (pH 8) 0.54 g 1.5M NaCl 13.5 mL of 5M

0.8M Guanidine HCl 3.44 g

0.5% Triton-X 100 0.225 mL (225 μL) of 100%

Add above ingredients to 50 mL tube.

Add milli-Q water to ~40 mL

Add NaOH to pH 10 (several drops at a time)

Add milli-Q water to 45 mL

Filter-sterilize to remove possible spores

Autoclave. Slightly loosen lid so that it is not air-tight. Recover from autoclave very soon after the autoclave cycle is completed.

Pour autoclaved solution into fresh 50 mL tube.

Aliquot into 1.5 mL tubes.

Hot Lysis

- Add 1.4 mL of DEB to each Sterivex with syringe and needle. Position the needle just below the mouth of the Sterivex so that it does not come back out the top. Do not fill to the top stop when solution covers white filter. Possible Stopping Point. Store at 20°C
 - 2.1 Possible Stopping Point. Store at 20°C
- 3 Place sterivex filter in 50mL tube with holes.
- 4 Incubate at 65°C for 30 mins. on Genemate spinning machine.
- 5 Vortex each sterivex again (inside the Falcon tube) for 30 seconds. Bead Beating:

protocols.io
2
09/28/2021

Citation: Christopher Neil Thornton, William Brazelton (09/28/2021). DNA Extraction from Sterivex Filters. https://dx.doi.org/10.17504/protocols.io.bykqpuvw

Bead Beating		
6	Using a syringe, withdraw fluid from each Sterivex and eject into bead tube (glass 0.1 mm for bacteria).	
7	Bead beat for 40 s.	
8	Centrifuge for 2 min at 5000 g.	
9	Add equal volume of phenol / chloroform / isoamyl alcohol (25:24:1, bought premixed with alkaline buffer) to each tube.	
Phenol/Chloroform extraction		
10	Transfer fluid avoiding beads into fresh Eppendorf tube. Add no more than 900 μL in each tube (or no more than 750 μL if using 1.5 mL tubes). Phenol / chloroform extraction:	
11	Gently shake a few times and then centrifuge at 14,000g for 1 minute.	
12	Remove supernatant to fresh tube.	
13	Add equal volume of chloroform / isoamyl alcohol (24:1) to each tube.	
14	Gently shake a few times and centrifuge.	
15	Remove supernatant to fresh tube, carefully avoiding the bottom organic layer. Ethanol precipitation:	
Ethanol precipitation		
16	Redistribute aqueous phase among 3 tubes so that each 2.0 mL tube has 550 μ L or less and each 1.6 mL tube has 450 μ L or less. For some samples, additional salt is not necessary, and you can skip the sodium acetate. In this case, you can add up to 600 μ L in a 2.0 mL tube.	
17	Add 0.1 volumes sodium acetate (3M, pH 5.2). (e.g. add 55 μ L to 550 μ L.)	

18	Add 2 Volumes 100% etnanol. (e.g. add 1210 μL to 605 μL.)
19	[optional for low biomass samples] Add 1.2 ul of glycogen (20 ug/ul).
20	Invert a few times to mix.
21	Incubate at 20°C for at least 1 hr. or overnight. Incubation on ice might work just as well and yield a cleaner pellet.
22	Centrifuge for 40 minutes at 16,000g. (Optional: used cooled centrifuge at 0°C)
23	Pour out supernatant. Do not completely invert tube; keep at a gentle angle to minimize the chance of the pellet falling out.
24	Add 500 μL of cold 70% ethanol to each tube.
25	Invert the tube to mix. Make sure the pellet is dislodged from the bottom so that it is properly washed.
26	Centrifuge at 16,000g for 10 minutes.
27	Remove liquid again with pipettor. Be careful to avoid pellet.
28	Place tubes with open lids in the Vacufuge. Spin for 7 minutes at 30°C on the VAL setting. If you can see ethanol in the tube, spin for another 25 minutes. If the pellets become powdery, they are too dry.
29	Resuspend in ${\sim}100~\mu\text{L}$ of low EDTA TE. Heat to 55°C
	Recipe for low EDTA TE: 10 mM TrisHCl 0.1 mM EDTA For 50 ml: 500 μ l 1 M TrisHCl (pH 8.0) autoclaved 10 μ l 0.5 M EDTA (pH 8.0) autoclaved \rightarrow to 50 ml with milliQ H2O \rightarrow filter sterilize with 0.22 μ m syringe filter TE is good for DNA storage, but EDTA inhibits PCR. So this low EDTA TE buffer is a good compromise for storing DNA for later PCR amplification. You can also just use EB (10 mM TrisHCl, pH 8 or 8.5).

for 10 or more minutes to dissolve pellet and store at 4°C. For longterm storage, place at 20 or 80°C, but avoid repeated freezing and thawing of the DNA. One strategy is to keep half at 4°C for the working sample and store the other half at 80°C as the archive sample.