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# ONA extraction - Zooplankton - 50 tubes

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# OPEN ACCESS



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

working

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## Abstract

This protocol is derived from the QIAamp DNA micro kit (QIAGEN) protocol and was used to extract DNA from whole or parts of zooplanktonic freshwater crustaceans (Copepoda, Branchiopoda, ...) from New Caledonia.



# Materials

QIAamp (R) DNA Micro kit (50), QIAGEN

# Before start

Make sure to properly prepare your buffers (e.g. heating, adding ethanol)

1	Prepare your 50 tubes with one individual per well. Alternate genus in the wells to detect eventual contamination between wells.	1d
1.1	Collect one individual from a sample	1m
1.2	Note its genus and determine its sex with a binocular microscope	1m
1.3	For big individuals (more than 5 mm), dissect a few legs and put it in the well. Be careful not to damage the rest of the body and put it in a tagged Eppendorf tube.  For little individuals (less than 5 mm), put the whole body.	3m
	Note	
	If necessary, use alcool to get the biological material to fall at the bottom of the well	
1.4	When all tubes are filled, the biological material has to dry before going to lysis	12h
	Note	
	If necessary, use a micropipette to empty an excess of alcool in the well	
	Safety information	
	Make sure the plate is closed when you want to transport it elsewhere	
2	Prepare the lysis	15m
2.1	Add 4 180 µL ATL buffer in each well and equilibrate to room temperature	10m

10m



- 2.2 Add 🚨 20 µL K proteinase in each well, mix by pulse-vortexing for 🚷 00:00:15 and briefly centrifuge
- 15s
- 3 Put your tubes in a proofer at \$\mathbb{L}\$ 56 °C Overnight (6h or more) to lyse the tissues

4 Perform the DNA extraction manually

#### Note

When manually extracting DNA alone, do not do all 50 tubes at once as timing is short between each step. A maximum of twenty tubes can be done simultaneously.

- 4.1 Optional steps, if RNA carrier is needed: Quantity for twenty tubes (using 20 volumes + 2 margin to avoid errors = 22 volumes)
  - Mix Δ 22 μg solid RNA carrier | with Δ 22 μL AE buffer | and mix until dissolved.
  - Δ 200 μL per volume) and mix.
- 4.2 Add A 200 µL AL buffer (+RNAse if suited) in each tube and mix by pulse-vortexing for 00:00:15
- 15s
- 4.3 Add \( \Lambda \) 200 \( \mu L \) 96-100\( \% \) ethanol \( \min \) each tube, mix by pulse-vortexing for \( \hat{N} \) 00:00:15 \( \, \, \) 5m 15s incubate at room temperature for 60 00:05:00.
- 4.4 Briefly centrifuge to remove drops from the lid.
- 4.5 Prepare all the eluting columns by puting each in a 2mL collection tube.
- 4.6 Transfer the prepared DNA lysate into the eluting column with a pipette, centrifuge the column the column in a clean 2mL collection tube.
- 1m



#### **Safety information**

Make sure the column is empty of liquid. If not, centrifuge at higher speed until it is.

4.7 Add  $\perp$  500 µL AW1 buffer in the column, centrifuge  $\bigcirc$  8000 rpm, 00:01:00 , discard the collection tube containing the flow through and place the column in a clean 2mL collection tube.

1m

4.8 Add  $\perp$  500 µL AW2 buffer in the column, centrifuge  $\bigcirc$  8000 rpm, 00:01:00 , discard the collection tube containing the flow through and place the column in a clean 2mL collection tube.

1m

4.9 Centrifuge , 00:03:00 , Full speed to dry the membrane completely.

3m

- 4.10 Place the column in a clean, closable, eppendorf tube.
- 4.11 Add  $\perp$  50 µL AE buffer in the column, incubate at room temperature for (5) 00:01:00 and centrifuge 3, 00:01:00, Full speed. The DNA extract in the clean eppendorf tube, discard the column.

2m