





Oct 11, 2022

# Invertebrate bulk sample metabarcoding protocol collection

## Dominik Buchner<sup>1</sup>

<sup>1</sup>University of Duisburg-Essen, Aquatic Ecosystem Research



dx.doi.org/10.17504/protocols.io.j8nlkw4n6l5r/v1



#### **ABSTRACT**

This is a collection of protocols currently used in the LeeseLab to perform invertebrate bulk sample metabarcoding. Feel free to contact us if any questions arise.

DOI

dx.doi.org/10.17504/protocols.io.j8nlkw4n6l5r/v1

#### **COLLECTION CITATION**

Dominik Buchner 2022. Invertebrate bulk sample metabarcoding protocol collection . **protocols.io** 

https://dx.doi.org/10.17504/protocols.io.j8nlkw4n6l5r/v1

**KEYWORDS** 

metabarcoding, malaise trap, lysis, DNA extraction, PCR cleanup, PCR normalization

**LICENSE** 

This is an open access collection 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** 

Oct 11, 2022

LAST MODIFIED

Oct 11, 2022

**COLLECTION INTEGER ID** 

71149

ABSTRACT



1

This is a collection of protocols currently used in the LeeseLab to perform invertebrate bulk sample metabarcoding. Feel free to contact us if any questions arise.

### **FILES**

