

# The Gnatwork Intended use of resource / data

Open access resources and data provided by The Gnatwork should be used for the intended purpose only, as specified below.

#### Title of resource

DNA Extraction Protocol for Simulium sp. and Onchocerca volvulus

# **Authored by**

When using this protocol, the following should be referenced:

Samuel Armoo<sup>1</sup>, Romain Derelle<sup>2</sup>, Daniel Boakye<sup>3</sup>, Mike Y. Osei-Atweneboana<sup>1</sup>, Charles Brockhouse<sup>4</sup>, John Colbourne<sup>2</sup>. Adapted from Brockhouse *et al* (1993).

<sup>1</sup>Council for Scientific and Industrial Research – Water Research Institute, Accra, Ghana.

#### DOI

N/A

# **Description**

This protocol is for extraction of genomic DNA from *Simulium* sp and *Onchocerca volvulus* tissues. This is a CTAB-based protocol, which is adapted from Brockhouse *et al* (1993).

#### Intended use

Scientific research use and training purposes.

#### **Restrictions on use**

Content is not to be redistributed in the public domain (e.g. presentation, lecture, online or in publications).

#### **Resource history**

<sup>&</sup>lt;sup>2</sup>University of Birmingham, Birmingham, UK

<sup>&</sup>lt;sup>3</sup>Noguchi Memorial Institute for Medical Research, Accra, Ghana

<sup>&</sup>lt;sup>4</sup>Creighton University, Omaha, NE, USA

#### GN 24 DNA Extraction Protocol for Simulium sp. and Onchocerca volvulus

When using this protocol, the following should be referenced: Samuel Armoo, Romain Derelle, Daniel Boakye, Mike Y. Osei-Atweneboana, Charles Brockhouse, John Colbourne. Adapted from Brockhouse *et al* (1993)

#### 1.0 Introduction

Due to the high pigmentation of blackflies, and the PCR inhibition properties of the pigments, the CTAB-based protocol used by Brockhouse and co (Brockhouse et al., 1993) remains a viable choice for DNA extraction given the consistency of high quality DNA produced. Therefore, this protocol was used for DNA extraction in this project with slight modifications. More details below.

#### 2.0 Preparing Buffers and Solutions

Buffers and solutions are prepared in advance and stored for downstream applications

Details of buffer and solution preparation steps below.

## 2.1 CTAB homogenization buffer

How to prepare a 200 mL solution of CTAB homogenization buffer (100nM Tris-HCL pH 8; 1.4M NaCl; 0.2%  $\beta$ -mercaptoethanol; 0.02 Na<sub>2</sub>EDTA; 2% CTAB)

Add the reagents below to160 mL of distilled water. Use a fume hood.

Reagent	Mass / Volume
NaCl	16.36 g
β-mercaptoethanol	400 uL
Tris-HCI*	20 mL
Na <sub>2</sub> EDTA (0.5M)*	8 mL
СТАВ	4 g

<sup>\*</sup>See appendix for preparation

- Top up to 200 mL with distilled water.
- Filter solution through a Stericup<sup>®</sup> filter unit or any portable filtration.
- Store at 4 °C

# 2.2 chloroform: isoamyl alcohol (24:1)

• Mix the following in a fume hood.

Reagent	Volume
Chloroform	192 mL
Isoamyl alcohol	8 mL

Store at room temperature in a fume hood.

#### 2.3 Sample Preparation

DNA was extracted from alcohol-dehydrated blackfly samples, and air-dried individual *O. volvulus* larvae on glass slides. Blackfly heads, and whole larval worms were used for DNA extractions. To re-hydrate the

preserved samples, blackfly heads and worm larvae were transferred into TE buffer and stored overnight at 4 °C.

### 2.4 Sample Homogenization and DNA Extraction

- 1. Transfer rehydrated blackfly or worm tissue into 500 uL of pre-warmed (60 °C) CTAB homogenization buffer in a 1.5 mL tube.
- 2. The animal tissue was crushed in the CTAB homogenization buffer using BioEcho tissue grinding pestle.
- 3. Add 1 mg/mL of proteinase K and mix.
- 4. Incubate at 60 °C overnight in a mixing incubator at low speed.
- 5. Add 500 uL of chloroform/isoamylalcohol (24:1) solution (in fume hood), and gently mix for 3 mins by inverting the tube.
- 6. Centrifuge at 14,000 xg for 12 min.
- 7. After centrifugation the mixture separates into a colorless aqueous upper layer, an interphase, and a lower chloroform phase. Transfer the upper aqueous layer into a new 1.5 mL tube, and discard the old one.
- 8. Add 1 uL DNase-free RNase to the recovered mixture and incubate at 37 °C for. 30 min.
- 9. Add 350 uL of isopropanol (or 2/3 of the recovered volume), and gently mix by inverting the tube.
- 10. Incubate mixture overnight at room temperature.
- 11. Centrifuge the mixture at 14,000 xg for 15 mins to pellet the DNA.
- 12. Carefully remove and discard the supernatant, then add 100 uL of cold freshly prepared 70% ethanol.
- 13. Centrifuge the mixture at 14,000 xg for 15 mins to pellet the DNA.
- 14. Carefully remove and discard the supernatant, then add 100 uL of cold freshly prepared 70% ethanol.
- 15. Centrifuge the mixture at 14,000 xg for 15 mins to pellet the DNA.
- 16. Carefully remove and discard the supernatant, then dry the pellet by leaving tube open at room temperature for 5 minutes.
- 17. Resuspend the pellet in 100 uL of low TE buffer (pH 8), and store at 20 °C

#### Reference

**Brockhouse, C.L., Vajime, C.G., Marin, R., and Tanguay, R.M.** (1993). Molecular identification of onchocerciasis vector sibling species in black flies (Diptera: Simuliidae). Biochem Biophys Res Commun *194*, 628-634.

#### **Appendix**

How to prepare a100 mL of 1M Tris-HCL (pH 8.0) solution

Reagent	Mass / Volume
Tris salt	12.11 g
Distilled water	80 mL

Add 12.11 g of Tris base to 80 mL of distilled water. Adjust to pH 8 with HCl (usually 4.2 mL of HCl will be required).

# How to prepare a 100 mL of 0.5M EDTA (pH 8.0) solution

Reagent	Mass / Volume
EDTA disodium salt	18.61 g
Distilled water	80 mL

Add 18.61 g of EDTA disodium salt to 80 mL of distilled water. Adjust pH to 8 with a few pellets of NaOH (usually 2g NaOH pellets will be required). Top up with distilled water to 100 mL.

# How to prepare a100 mL solution of TE buffer

Reagent	Volume
0.5M EDTA (pH 8)	0.2 mL
Tris-HCL (pH 8)	1 mL

- Top-up the above mixture with distilled water to 100 mL.
- Sterilize by autoclaving
- Store at 4 °C