

University of Wisconsin Biotechnology Center

Gene Expression Center

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Client: Funding No. Direct Invoice Sub.No. M007087

Tissue RNA Extraction (miracidia (first larval stage) of Schistosoma mansoni)

Client Notes- 9 samples that were flash frozen in liquid nitrogen and stored in Trizol at -80C. Each sample includes 500-1200 miracidia (first larval stage) of *Schistosoma mansoni*. Each miracidium has 350-400 cells, so we may be dealing with a low-input situation. The genome size is about 400 Mb (<u>link to the database</u>). Our goal is to identify differentially expressed genes from different populations of miracidia (two populations, >3 biological replicates each)

7/23/2024 SMSB

Client will be sending samples in 1mL Trizol, on dry ice, in standard 1.5mL microfuge tube Suggested RNA extraction protocol provided

7/24/2024 SMSB

Approved service request for RNA Extraction & RNA Seq with sample being shipped next week

Date	Number of miracidia	Tissue Source	Label
4/18/24	600	Liver	Sma_NMRI_Mira_Liv-01
6/19/24	730	Liver	Sma_NMRI_Mira_Liv-02
7/11/24	350	Liver	Sma_NMRI_Mira_Liv-03
7/18/24	1200	Liver	Sma_NMRI_Mira_Liv-04
4/18/24	600	Intestine	Sma_NMRI_Mira_Int-01
6/6/24	152	Intestine	Sma_NMRI_Mira_Int-02
7/11/24	200	Intestine	Sma_NMRI_Mira_Int-03
7/18/24	450	Intestine	Sma_NMRI_Mira_Int-04

7/30/2024 LW

Samples (8) received by shipment on dry ice. All samples were frozen in ~1mL of trizol on arrival. Samples stored at 4C.

RNA Extraction- Organic homogenization w/ Qiagen RNeasy Kit w/ DNase treatment

8/13/2024 LW

Working in the biohood in room 2330A, samples transferred to 2mL Safelock tubes with 5mm stainless steel bead

RNA Extraction (8) – Batch 01

Nucleic Acid Extraction Room2330A

Samples homogenized by Tissuelyzer at 30Hz for 2min for 1 cycle, samples were allowed to rest at on ice for 5 minutes. Chloroform was added to samples, organic extraction performed.

Qiagen RNeasy Micro Kit (lot#172040952) used for cleanup.

On-column DNase treatment (Lot#178011710)

Sample eluted w/ 14uL nuclease-free water added to column

8/13/2024 LW

QC- each assayed on a Nanodrop One- A_{260}/A_{280} < 1.8 and phenol contamination Assayed samples on Agilent 4200Tapestation- RNA assayed on a High Sensitivity RNA screentape post denaturation

Profile comments added to EXCEL sheet

A visible measure of a core's impact is made through proper acknowledgement of usage in publications. To acknowledge our cores, please use the following statement:

The author(s) utilized the University of Wisconsin – Madison Biotechnology Center's Gene Expression Center Core Facility (Research Resource Identifier - RRID:SCR_017757) for RNA Extraction & QC services.