

Resource Summary Report

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RNA extraction protocol (Trizol)

DOI:10.17504/protocols.io.ew7bfhn

Type: Protocol

Proper Citation

Hebert F.O., Grambauer S., Barber I., Landry C.R., Aubin-Horth N. 2016. RNA extraction protocol (Trizol). protocols.io dx.doi.org/10.17504/protocols.io.ew7bfhn

Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.ew7bfhn>

Authors: Hebert F.O., Grambauer S., Barber I., Landry C.R., Aubin-Horth N.

Group: GigaScience Press

Summary: This protocol describes how to extract total RNA from flatworms. It is from:Hebert, F, O; Grambauer, S; Barber, I; Landry, C, R; Aubin-Horth, N (2016): Reference transcriptome sequence resource for the study of the Cestode *Schistocephalus solidus*, a threespine stickleback parasite. GigaScience Database. <http://dx.doi.org/10.5524/100197>

Associated Publications: Hebert FO, Grambauer S, Barber I, Landry CR, Aubin-Horth N, Transcriptome sequences spanning key developmental states as a resource for the study of the cestode , a threespine stickleback parasite. GigaScience doi: [10.1186/s13742-016-0128-3](https://doi.org/10.1186/s13742-016-0128-3)

Affiliations: GigaScience, GigaScience, GigaScience, GigaScience, GigaScience

External URL: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4891850/>

Version: 1

Publication Date: 2016

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Record Creation Time: 20210329T220531+0000

Record Last Update: 20210329T220958+0000

Data and Source Information

Source: