

Resource Summary Report

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Identifying Viral Sequences Using VirSorter (Cyverse)

DOI:10.17504/protocols.io.eyjbfun

Type: Protocol

Proper Citation

Benjamin Bolduc 2016. Identifying Viral Sequences Using VirSorter (Cyverse). protocols.io dx.doi.org/10.17504/protocols.io.eyjbfun

Protocol Information

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

Authors: Benjamin Bolduc

Group: Sullivan Lab, iVirus

Summary: Identifying putative viral sequences from SPAdes-assembled data from the Ocean Sampling Day (2014) metagenomic datasets using VirSorter.

Associated Publications: Bolduc B, Youens-Clark K, Roux S, Hurwitz BL, Sullivan MB, iVirus: facilitating new insights in viral ecology with software and community data sets imbedded in a cyberinfrastructure. The ISME Journal 11(1). doi: [ismej201689](#)

Affiliations: The Ohio State University

External URL: <https://www.nature.com/articles/ismej201689.pdf?origin=ppub>

Version: 2

Publication Date: 2016

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