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# Processing a Viral Metagenome Using iVirus V.2

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1 Works for me

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Sullivan Lab iVirus

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#### **ABSTRACT**

A collection of protocols designed to guide the user in processing a viral metagenome from raw sequence data to assembly, and subsequent analysis. The user uses *actual* reads from <u>Ocean Sampling Day (2014)</u> and processes them entirely within Cyverse, a NSF-supported cyberinfrastructure.

**EXTERNAL LINK** 

https://www.nature.com/articles/ismej201689.pdf?origin=ppub

THIS COLLECTION ACCOMPANIES THE FOLLOWING PUBLICATION

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: <a href="mailto:ismej201689">ismej201689</a>

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