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## Processing a Viral Metagenome Using iVirus V.3

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1 Works for me

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Sullivan Lab iVirus

Benjamin Bolduc

**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

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## Downloading Viral Metagenome Data Version 1

by Benjamin Bolduc



	<u></u>	Assembling Viral Metagenomic Data with SPAdes (Cyverse)  Version 2  by Benjamin Bolduc
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		Clustering Viral Genomes in iVirus  Version 1 by Benjamin Bolduc
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