

Resource Summary Report

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Script R15: CoNet Analysis - Formatting Relative Abundance Files

DOI:10.17504/protocols.io.ejpbcmn

Type: Protocol

Proper Citation

HANNIGAN GD, GRICE EA, ET AL. 2016. Script R15: CoNet Analysis - Formatting Relative Abundance Files. protocols.io dx.doi.org/10.17504/protocols.io.ejpbcmn

Protocol Information

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

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Group: VERVE Net, Club Grice

Summary: This protocol outlines the analysis used to generate input files for CoNet for the phage-bacteria network. Based on methods from the following publication:Hannigan, Geoffrey D., et al. "The Human Skin Double-Stranded DNA Virome: Topographical and Temporal Diversity, Genetic Enrichment, and Dynamic Associations with the Host Microbiome." mBio 6.5 (2015): e01578-15.

Associated Publications: Kindler L, Stoliartchouk A, Teytelman L, Hurwitz BL, Method-centered digital communities on protocols.io for fast-paced scientific innovation. F1000Research doi: [10.12688/f1000research.9453.2](https://doi.org/10.12688/f1000research.9453.2)

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