



The Missing Tailed Phages: Prediction of Small Capsid Candidates

Antoni Luque^{1,2,3,*}, Sean Benler⁴, Diana Lee^{1,2}, Colin Brown^{1,5}, and Simon White⁶

¹ Viral Information Institute, San Diego State University, San Diego, CA, USA.

² Computational Science Research Center, San Diego State University, San Diego, USA.

³ Department of Mathematics and Statistics, San Diego State University, San Diego, USA.

⁴ National Center for Biotechnology Information (NCBI), Bethesda, MD, USA.

⁵ Department of Physics, San Diego State University, San Diego, USA.

⁶ Department of Molecular and Cell Biology, University of Connecticut, Storrs, CT, USA.

* Correspondence: aluque@sdsu.edu.

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Supplementary Figures and Tables

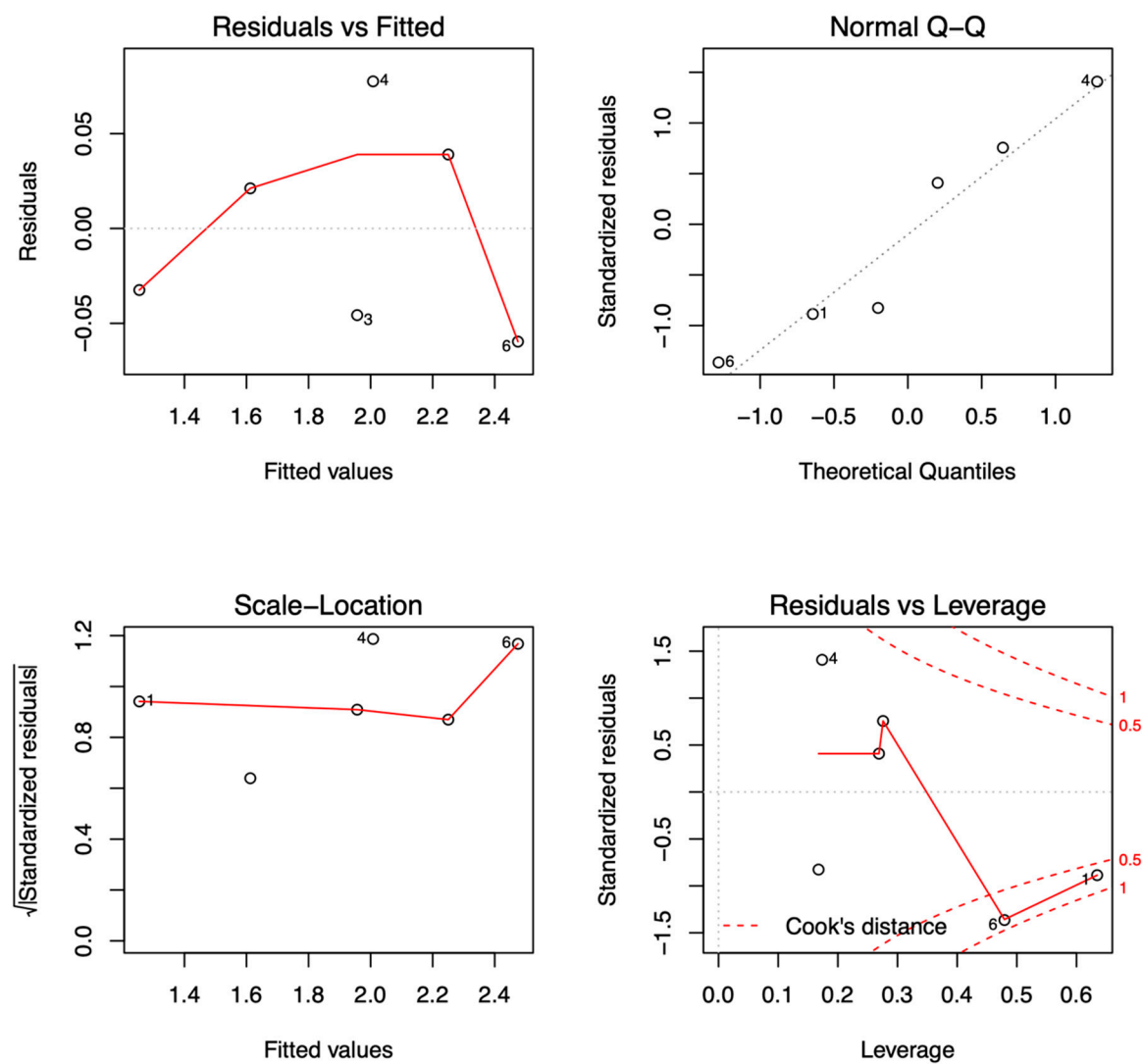


Figure S1. Residual diagnostics for the genome length model.

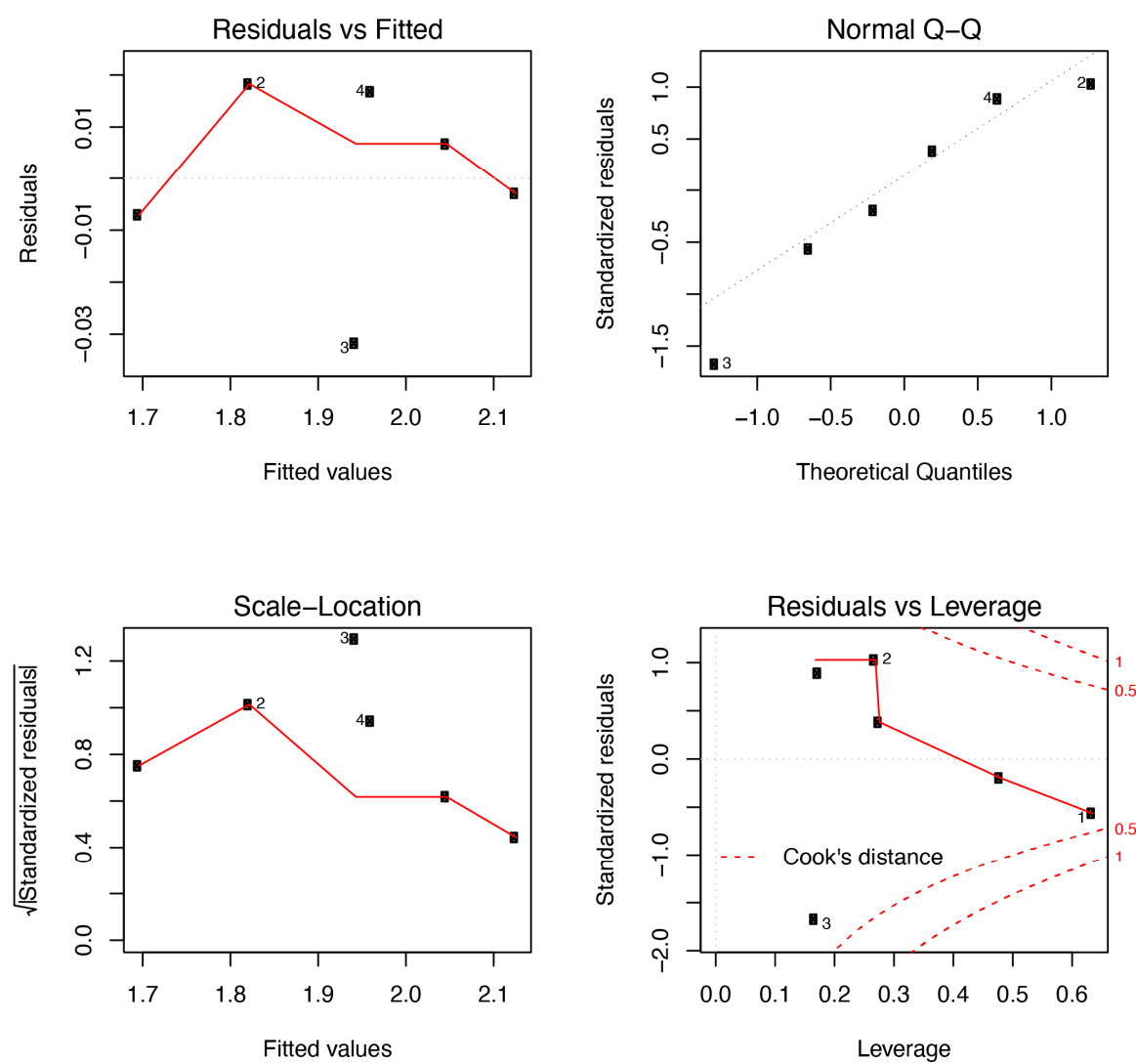


Figure S2. Residual diagnostics for the capsid diameter model.

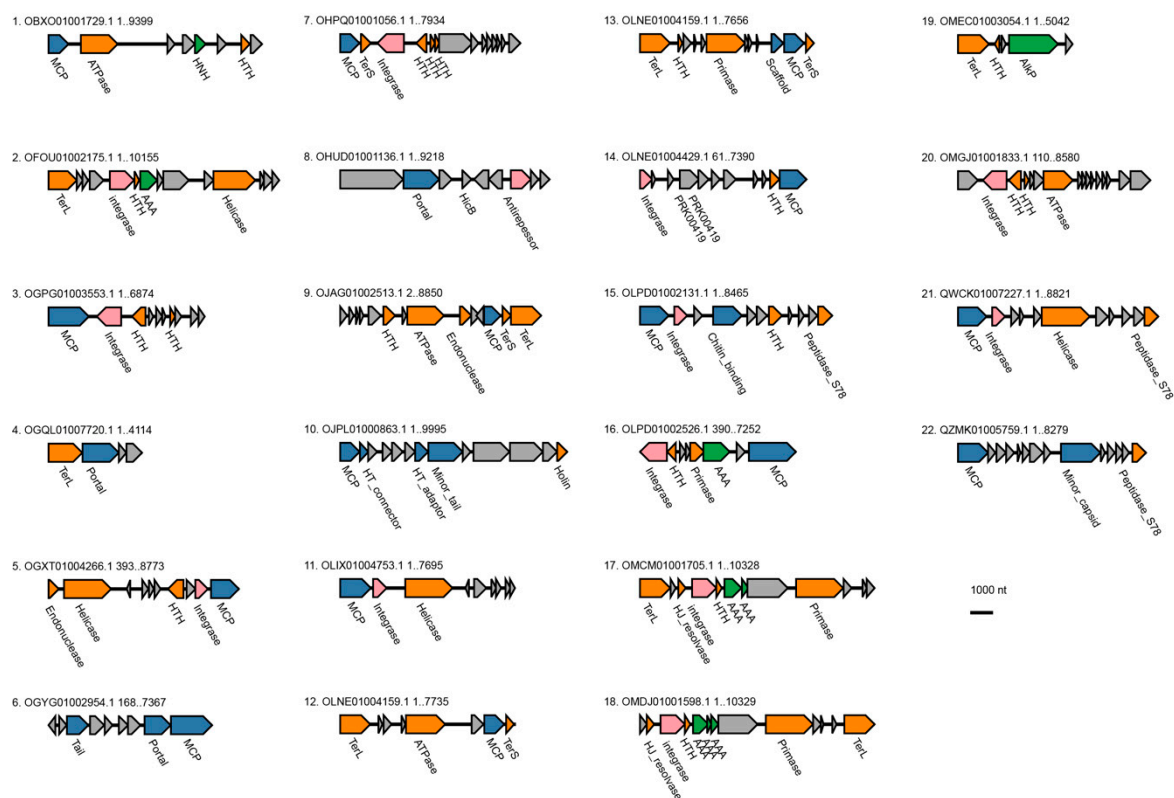


Figure S3. Annotated metagenome-assembled tailed phage genomes for predicted $T \leq 3$ capsid architectures.

Property	rho	p-value
Interior volume	0.924	$4.09 \cdot 10^{-6}$ ***
Interior surface	0.923	$4.08 \cdot 10^{-6}$ ***
Exterior surface	0.989	$2.84 \cdot 10^{-6}$ ***
Genome size	0.817	$3.37 \cdot 10^{-6}$ ***
Capsids thickness	0.663	0.0010 **
Interior sphericity	−0.566	0.0069 **
Exterior sphericity	−0.684	0.0006 ***
Genome density	−0.162	0.4695
MCP interior area [§]	0.484	0.0242 *
MCP exterior area [§]	0.380	0.0819

Table S1. Correlation analysis with capsid size. Spearman's correlation coefficient (rho) for structural properties as a function of the external capsid diameter. The stars in the p-value are associated to standard cut-offs for statistical significance: (*) $p < 0.05$, (**) $p < 0.01$, and (***) $p < 0.001$.