

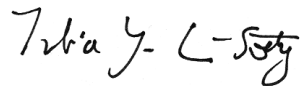
May 19, 2015

Dear Colleagues,

Thank you for your review of **Arthrobacter phage Brent**. I did not format the second “minimalistic” Final dnam5 file since the collection of Arthrobacter phages is not quite as vast as those for Mycobacteriophages and the functions of many gene features are still unknown.

Most of our genes BLASTed to phage vB_ArtM-ArV1 from Lithuania, but the genome of Brent is most similar to Jawnski based on local BLAST within PhagesDB. One gene feature we are not sure about is number 64. GeneMark did not predict substantial coding potential; Glimmer call was rather weak. However, it did appear on Phamerator and was “corroborated” by Jawnski. So, we decided to keep it.

Best,



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