

May 13, 2015

Greetings,

I am respectfully submitting the annotated genome of the bacteriophage SamScheppers. We feel confident in the majority of our annotation, but appreciate a second look at the following regions:

Number 9 - Start 5096 is questionable as it has a large overlap (20 bp) with the previous gene.

Number 19 - Start: 11957 - Is a translational frameshift.

Number 73- Start: 45586 - We are not completely certain on the start. There is another possible start that would close the gap, but its SD score is not as good, neither is the BLAST.

The function at several loci do not line up with the functions found on the Hatfull approved map, this may be because the genomes being compared to not line up well.

Thank you!

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