Genome Annotation Submission Cover Sheet

Pre-SM*ART QC Phage Genome Annotation Checklist

			-	
Phage Name:		Name	,	
Your Name:		lame:	Deborah Tobiason	
Your Institution:		nstituti	on: Carthage College	
Your email:		mail:	dtobiason@carthage.edu	
		onal en orrespo	nails: ndence)	
ΡI	ease	e check	c each box indicating completion of each task.	
////////	2. 3. 4. 5. 6.	Have tRNA ends been adjusted with web-based Aragorn and/or tRNAscan SE? Has the frameshift in the tail assembly chaperone been annotated (where applicable?)		
	 For the items below, generate a genome profile, and review the following. For the YourPhageName_CompleteNotes.dnam5 file: 			
////////		a. b. c. d. e. For	Have any duplicate genes (or any with the same stop coordinate?) been removed? Does every gene have one and only one complete set of Notes Do the functions in the Notes match the official function list? Are all three lines of functional evidence described for EVERY gene? Do the notes contain the initial Glimmer/GeneMark data from the autoannotation? the YourPhageName .dnam5 file: Have any duplicate genes (or any with the same stop coordinate?) been removed? Is the Notes field empty for all the features with no known function (including hidden	
///		О. С.	marks? Do the function names in the Notes match the official function list?	

d. Is the function field EMPTY for all features?

- 9. Did you use PECAAN to annotate your phage?
 - a. If, so please describe how in the text field after question 10.
- 10. Describe any issues or specific genes that you were unable to satisfactorily resolve, and warrant further inspection in the Quality Control review.

Older annotation done in 2016 so missing starterator.