To SMART Review team:

Thank you for reviewing our submission of annotation for mycobacteriophage GinkgoMaracino of the A3 subcluster. We are fairly confident of our annotations, and request specific review of the following genome features:

1. Gene 40 and 41 have no overlap, both blast hit to dCMP deaminase and it is tempting to merge them into a single gene, but there is a stop codon in frame separating the two genes.
2. Gene 48, DNA Polymerase 1, appears to be a Klenow fragment
3. Genes 61,65 are not called by glimmer but are called by Genemark
4. Stop1-15, stoplysis, stoplysogeny are stoperator sites for gp79 immunity repressor activity.

Sincerely,

Jeff McLean and the RCC 2017 Phage Hunters

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