**Mycobacteriophage HamSlice Annotation Cover Sheet**

Gp38 (inclusion/exclusion): very short ORF at 87 bp; matches 1:1 with gp39 of Peaches and Flux. There are no other start sites that make the ORF longer, and GeneMark-Smeg output contains coding potential that encompasses ORF.

Gp54 (inclusion/exclusion): gene was manually added. Not called by either Glimmer or GeneMark, manually added to minimize gap and because GeneMark output revealed coding potential; preserves gene order between Peaches and Hamslice; no significant Blast matches found

Gp73 (start choice): Disagrees with both Glimmer and GeneMark predictions (45952); manually changed to minimize gap and TTG start

Note:other similar phages (Peaches) call a gene before 803 bp (HamSlice current gp1). We left this unannotated due to lack of coding potential shown in GeneMark-Smeg output and no calls from GeneMark or Glimmer.