Lehigh University would like to submit mycobacteriophage Spongebob for annotation QC. We have checked all gaps and they are valid.

We have deleted gp13 (R 7808-79224), gp25 (R 21308-21622), gp57 (R 35648-362244), gp38 (R 28223- 28342) because they were reverse genes in a long stretch of forward genes and have little data to support the gene call.

We have also deleted feature 37 (tRNA 27821-27898) it falls within a reverse gene.

We have added gp53 (F 35612-36421) since it closes a large gap and has homology with many genes in other N cluster phages. We also added gp64 (F 40783-41091) because there is coding potential present and it has homology with other N cluster phages.

The following region we would like investigated more closely. We have a translational frameshift for gp14-15 (9081-9987). We have called a +1 translational frameshift at position 9558.