**Edugator**

I feel rather confident on my work with the bacteriophage known as Edugator. I made a few changes to start codons, as well as the deletion of certain genes.

* **Gene 44 and 45:** Was deleted due to the short ORF and no blast or NCBI results.
* **Gene 47:** Was deleted since it was in the other direction as the up and downstream genes. It was also had a very short ORF length of only 207bp.
* **Gene 52:** Had no coding potential at all and was also rather short at 225bp.
* **Gene 38:** was in the other direction of the up and downstream genes and only had a ORF of 132bp. It was also different from every other genome in phamerator when compared to bacteriophages in the same sub-cluster.
* **Gene 40:** Was a gene that did not have the best SD score, and was only 147bp long, so it was deleted.
* **Gene 84:** Did not have very good coding potential I would like someone to look at both genes 84 and 85.
* **Gene 85:** Was rather odd it only had one possible start codon, and was only 153bp in length. I left this as a gene though since it did have good coding potential.
* **Gene Numbers 97 and 98:** Were both deleted since they had no coding potential and where rather short at 150bp for gene 97 and 306bp for gene number 98.