**Genome Announcements**

**General Questions**

1. What phage(s) are in this Genome Announcement (GA)?
2. Where was the phage(s) found?
3. What is its host?
4. Was it isolated from direct plating or after enrichment?
5. What types of plaques does it make?
6. What is the virion morphology? Are head diameter and tail length reportable?
7. What type of sequencing technique(s) was used?
8. With what software was the genome assembled?
9. What is the sequence coverage?
10. How long is the genome?
11. What type of ends does this genome(s) have?
12. What is the GC content of your phage(s)? How does that compare to the host?
13. How many protein-coding genes does your phage have?
14. How many tRNA genes are there? Where are they located? Is there a tmRNA?
15. If there are more than one phage in your GA, what is the range of GC content, gene numbers, sequence lengths, etc. across your genomes?
16. Are there any publications about similar phages?