CONSIDER FOR TALK

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Complete genome Sequence of Mycobacterium smegmatis Phage Rummer, a Subcluster A3 Actinophage

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Bacteriophage Rummer is a siphovirus morphology actinophage isolated from *Mycobacterium smegmatis*. Rummer has a 50,908 base pair (bp) genome encoding 89 predicted protein-coding genes and three (3) tRNAs. Based on gene content similarity to sequenced actinobacteriophages, Rummer is assigned to phage subcluster A3.