CONSIDER FOR TALK

2021 SEA Symposium Abstract

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Complete genome sequence of Mycobacterium bacteriophage Stinson

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The complete genome sequence of *Siphoviridae* bacteriophage Stinson is 59.9kb long and comprises 78 putative protein coding genes. This subcluster K1 phage, infecting *Mycobacterium smegmatis* mc²155, has a temperate life cycle which is maintained by the immunity repressor protein, responsible for suppressing lysogenic gene expression.