Complete Genome Sequences of Nine *Bacillus cereus* Group Phages

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We report the sequences of nine novel *Bacillus cereus* group bacteriophages: DIGNKC, Juglone, Nemo, Nigalana, NotTheCreek, Phrodo, SageFayege, Vinny, and Zuko. These bacteriophages are double-stranded DNA-containing *Myoviridae* isolated from soil samples using *B. thuringiensis* subsp. *kurstaki* as the host bacterium.

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The members of the 2013–2015 VCU Phage Hunters courses at VCU are listed at [https://wiki.vcu.edu/display/phagelab/VCU+Phage+Hunters](https://wiki.vcu.edu/display/phagelab/VCU+Phage+Hunters).

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**REFERENCES**


