API480: features towards therapy in honeybee hives





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Introduction

American foulbrood disease (AFB) is a devastating bacterial disease affecting honeybees. It is caused by *Paenibacillus larvae*, a worldwide-distributed spore forming Gram-positive bacterium which spread easily across apiaries producing highly resistant spores. When AFB symptoms are found the burning of contaminated hives is mandatory causing serious economic losses [1]. Bacteriophages (phages) are being considered valuable solutions to the control of this infection [2-5]. So far, 48 *Siphoviridae P. larvae* phages sequences are known and most encode integration genes suggesting a temperate lifestyle. All of these 48 phages seem to have a common evolutionary ancestor showing an overall common structure. Their genomes were grouped into four clusters (with Fern, Harrison, Vegas and Halcyone as representative phages) and one singleton (phage Lily) [6].

Goal of the study

The main goal of this study was the characterization of the first podovirus infecting *P. larvae* (API480), envisaging AFB control.

Methods Phage viability in simulated field Genome sequencing, annotation and **Phage isolation** Phage characterization conditions compative analysis 50% (w/v) sucrose Larval homogenised TEM Illumina MiSeq Lytic spetra, EOP Royal jelly (RJ) Adsorption, OSGC (MOI=0.1) pH solutions (universal buffer) ranging Specificity test 3.0 to 5.0 Statistical significance (p<0.05) **☆**SplitsTree4 놀 python™ Culture condictions: MYPGP medium 37°C, 5% CO₂, O/N Cut-off of > 40% homology **PHAMERATOR** 10⁷ PFU.mL individual cluster

Results

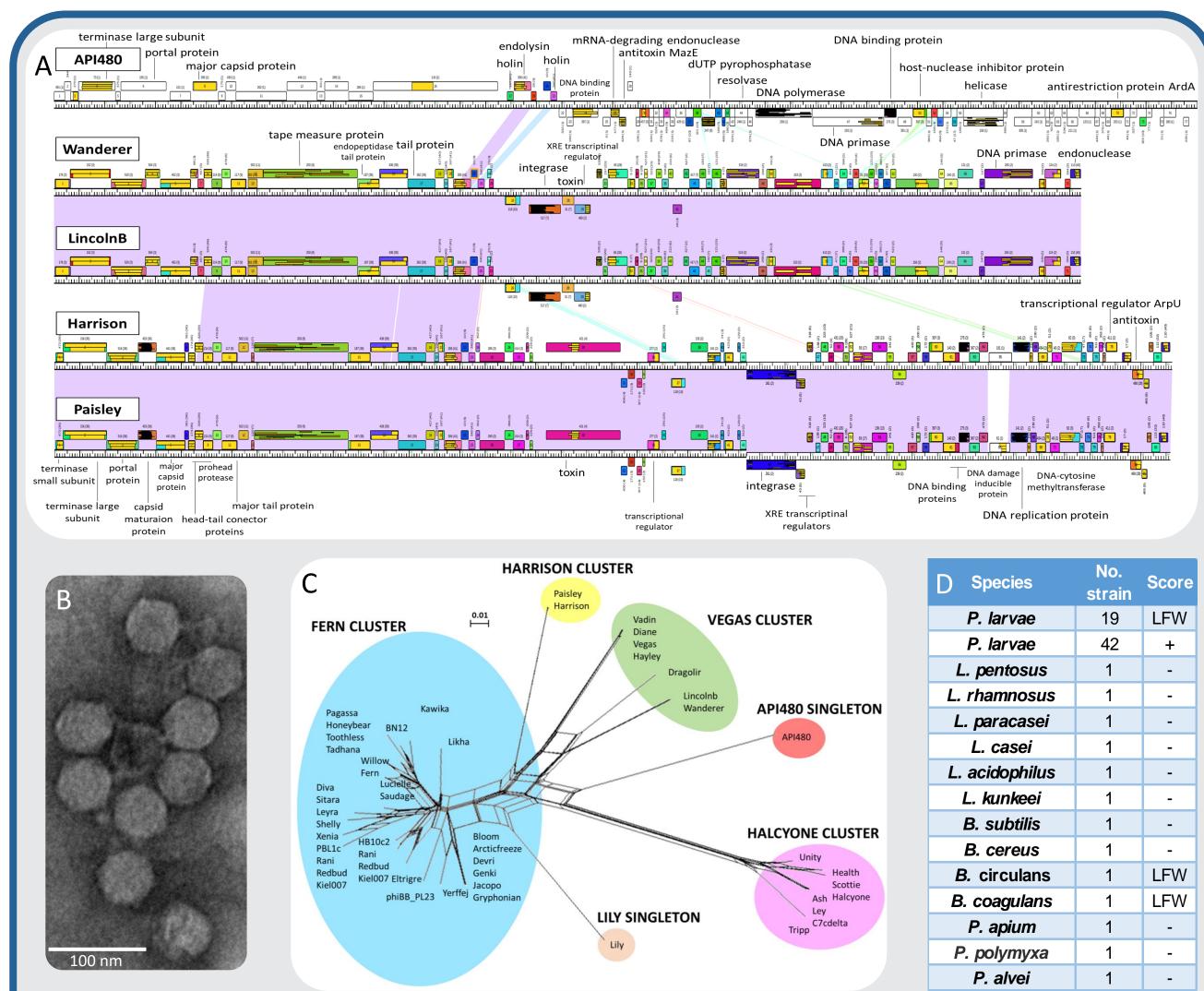


Figure 1. (A) Pairwise genome maps of API480 with the closest relatives phages. **(B)** TEM micrograph showing the virion particle morphology. **(C)** Relationship of shared gene content from a total 49 *P. larvae* phages. **(D)** API480 lytic spectra and EOP (susceptible (+), non-susceptible (-) and lysis from without (LFW)).

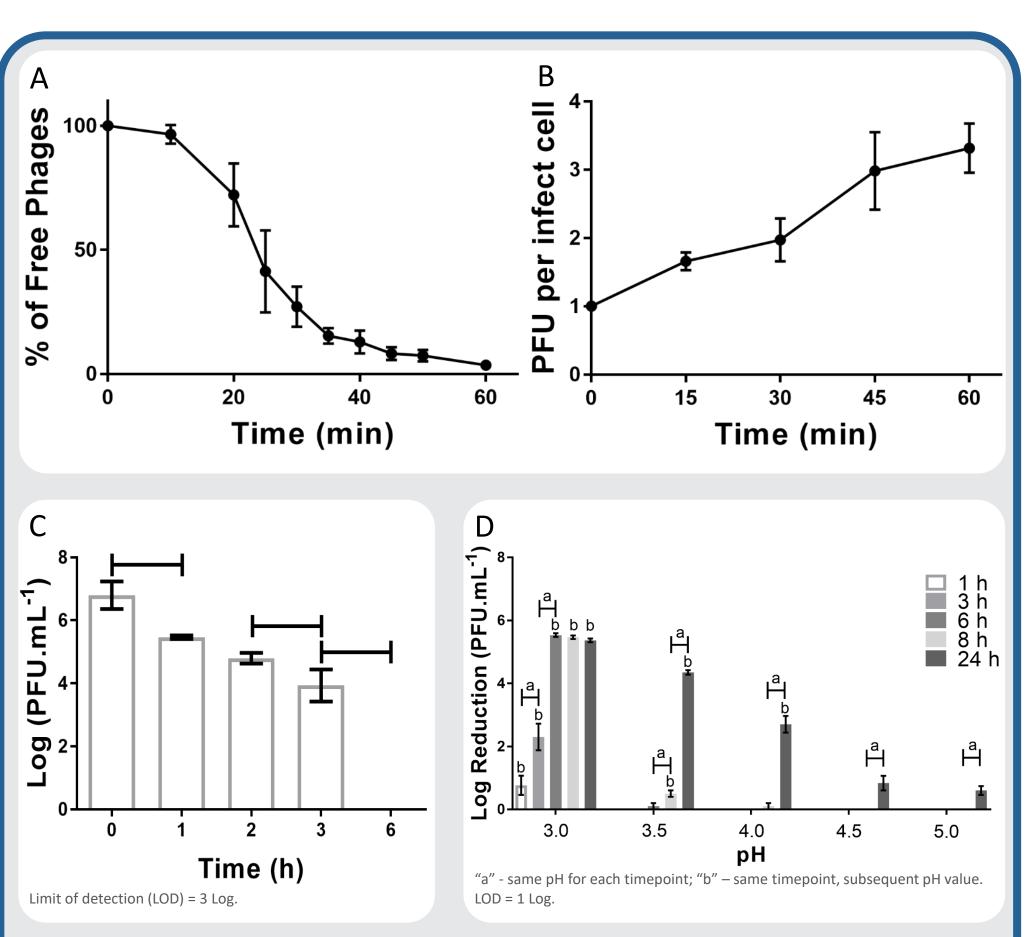


Figure 2. (A) Phage adsorption. (B) OSGC. (C) Effect of RJ on API480 concentration. (D) Effect of acidic pH in API480 concentration.

No negative effects caused in API480 phage viability by sucrose and larval homogenate after 24 hours.

Main conclusions

- API480 is a distinct phage (4% genome coverage and 14% shared proteins), suggesting the creation of a new species within the *Podoviridae* family.
- Besides no lysogenic module being identified, API480 is a temperate phage.
- API480 has a broad lytic spectrum and is specific to *P. larvae*. The larval commensal (*L. kunkeei* and *P. apium*) are not affected by this phage.
- This phage is very stable when exposed to high sucrose concentration and to larval homogenized content.
- Overall, results suggest that this phage holds potential to be used in the biocontrol of American Foulbrood disease.

Reference

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