

Supplementary Materials:

***Pectobacterium parmentieri* SCC 3193 Mutants with Altered Synthesis of Cell Surface Polysaccharides are Resistant to N4-Like Lytic Bacteriophage ϕ A38 (vB_Ppp_A38) but Express Decreased Virulence in Potato (*Solanum tuberosum* L.) Plants**

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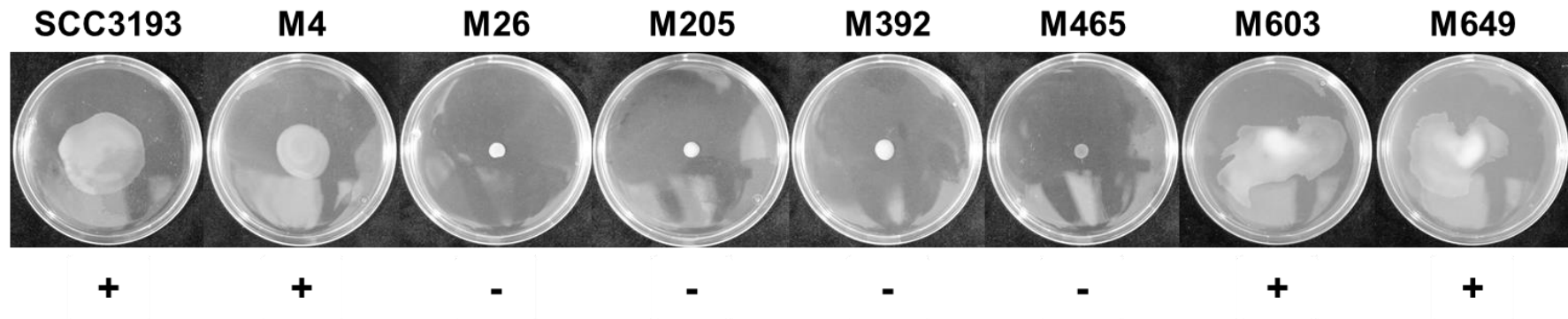
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Supplementary Material. Draft genome sequences of the seven Tn5 phage-resistant mutants (M4, M26, M205, M392, M465, M603 and M649) used to in detail localize the Tn5 transposon insertions in the genome of *P. parmentieri* strain SCC3193. Draft genome sequences are in FASTA format and compressed with Windows Zip.

a)



b)

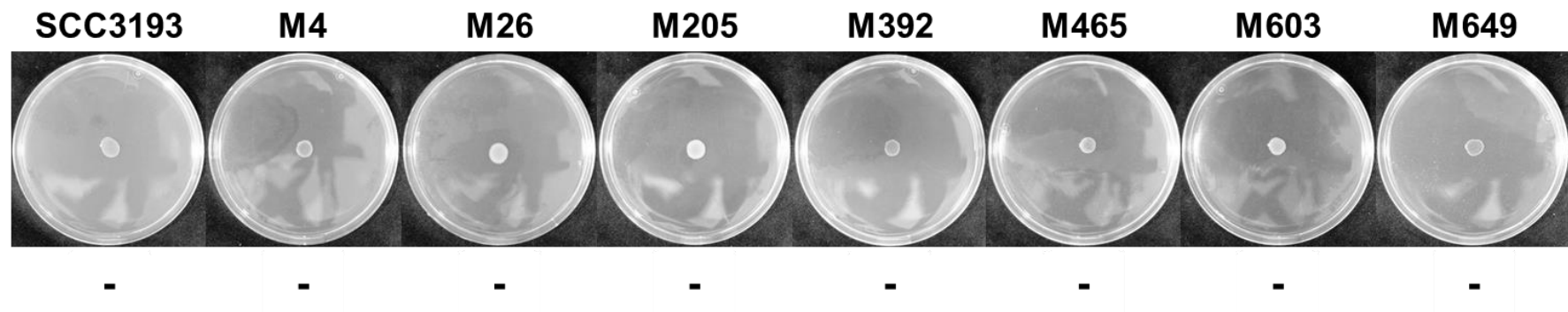
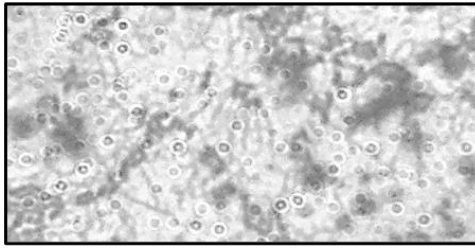
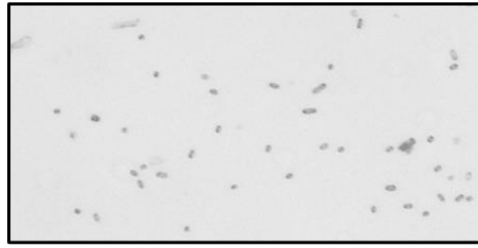


Figure S1. Assessment of swimming (a) and swarming (b) motility of SCC3193 WT strain and seven phage-resistant mutants (M4, M26, M205, M392, M465, M603 and M649). Representative photos are shown. (+) – indicates motility, (-) – indicates the lack of motility.

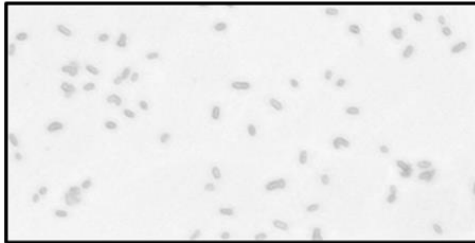
SCC3193



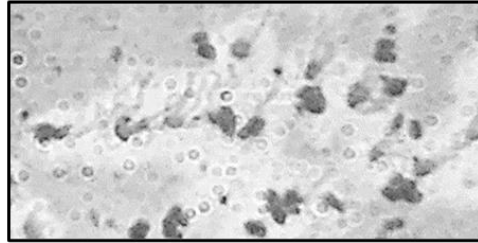
M4



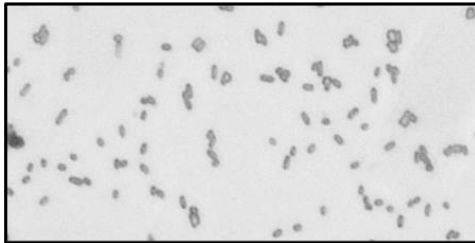
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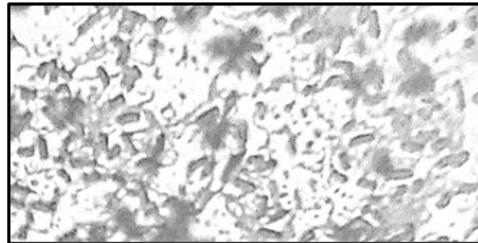
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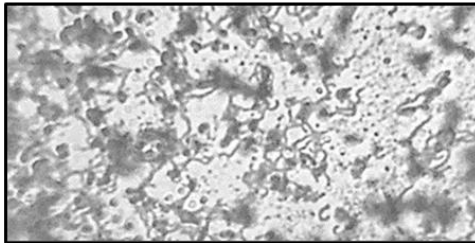
M392



M465



M603



M649

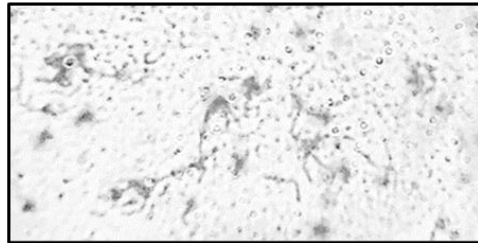


Figure S2. Visualization of the presence/absence of flagella of SCC3193 WT strain and seven phage-resistant mutants (M4, M26, M205, M392, M465, M603 and M649). Flagella staining was done using Ryu staining solution as described in [97]. Representative photos are shown.

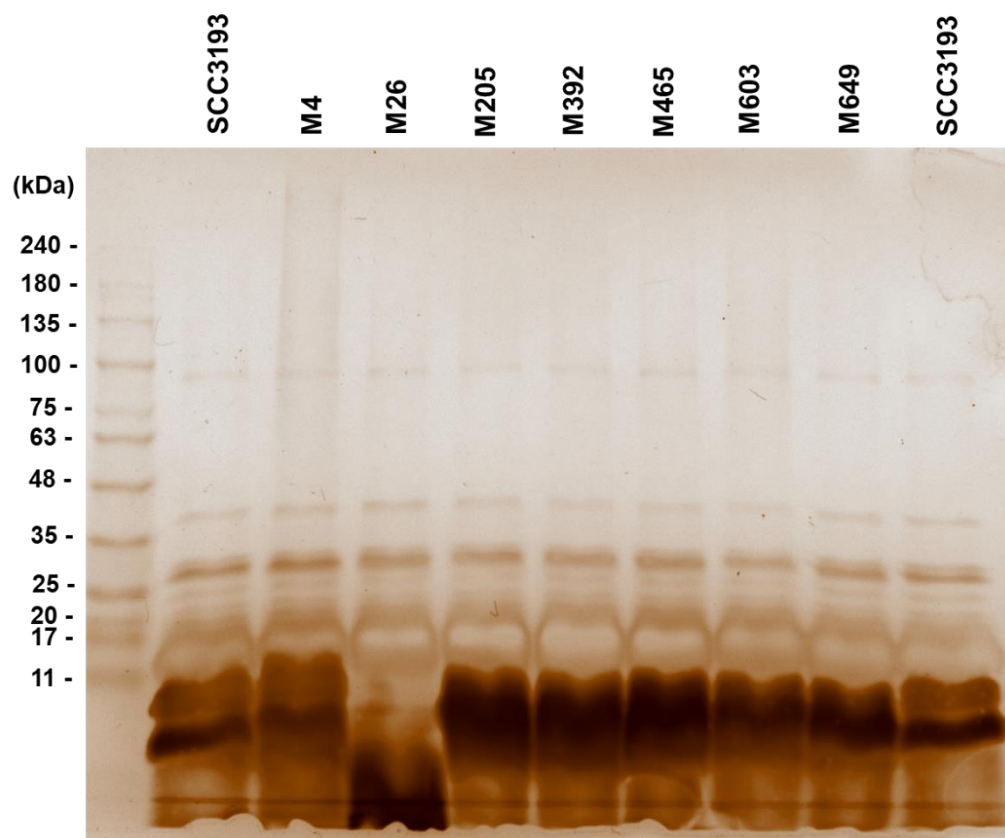


Figure S3. Characterization of lipopolysaccharide from wild type *P. parmentieri* strain SCC 3193 and seven phage-resistant mutants. (a) SDS-PAGE was performed using a gradient (4–20 %) polyacrylamide gel, and the LPS components were visualized by silver staining [77]. The size marker (11–245 kDa, Perfect Tricolor Protein Ladder, EURx, Poland) is shown in the first lane.