

Additional file 3 – Sequence analysis of prophage 03 of *P. fluorescens* Pf-5

| CDS | Gene name | Position | Protein length (residues) | Protein mol weight (kDa) | Putative ribosome binding site | Closest protein of phage or plasmid origin (accession no.), blastp E-value ^a | Other similar proteins (accession no.), blastP E-value ^a | Predicted function (functional domains) |
|----------------|-------------|----------------------|---------------------------|--------------------------|--------------------------------|--|---|---|
| PFL_1976 | <i>int2</i> | 2207060..2208202 | 380 | 43.927 | GTAG | Shufflon-specific recombinase from plasmid ColIb-P9 (AAZ05392), 4e-21 | Phage integrase family protein PputGB1_1711 from <i>Pseudomonas putida</i> GB-1 (YP_001667950), 0.0 | Putative phage integrase |
| 1 ^b | <i>orf1</i> | 2208171..2208452 (-) | 93 | 10.325 | GGAG | Transcriptional regulator PrtN from <i>Pseudomonas aeruginosa</i> pyocin (Q06552); 5e ⁻⁰⁵ | Conserved hypothetical protein PputGB1_1711 from <i>Pseudomonas putida</i> GB-1 (ABY97615), 3e-36 | Putative phage-related positive regulator |
| PFL_1977 | PFL_1977 | 2208449..2208922 (-) | 157 | 16.913 | GGAG | None detected | None detected | Hypothetical protein |
| 2 ^b | <i>orf2</i> | 2209446..2211179 (-) | 176 | 19.300 | AGGAG | Hypothetical protein gp37 from <i>Burkholderia</i> phage BcepC6B (AAT38396); 1e-15 | Hypothetical protein PA2G_00782 from <i>Pseudomonas aeruginosa</i> 2192 (EAZ57582); 2e ⁻¹³ | Conserved hypothetical phage protein |
| PFL_1978 | PFL_1978 | 2209446..2211179 (-) | 577 | 63.298 | GGAG | Hypothetical protein gp41 from <i>Pseudomonas aeruginosa</i> phage CTX (NP_490641); 0.0 | Hypothetical protein Orf27 from <i>Pseudomonas putida</i> plasmid pWVO (NP_542819); 0.0 | Conserved hypothetical protein (COG0270) |
| PFL_1979 | PFL_1979 | 2211169..2211507 (-) | 112 | 12.047 | AGGAG | None detected | Hypothetical protein PP3030 from <i>Pseudomonas putida</i> KT2440 (NP_745174); 4e ⁻¹⁰ | Conserved hypothetical protein |
| PFL_1980 | PFL_1980 | 2211507..2211728 (-) | 73 | 8.368 | AGGAG | None detected | None detected | Hypothetical protein |

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|----------------|-------------|----------------------|-----|--------|---------------|--|--|--|
| PFL_1981 | PFL_1981 | 2211725..2212144 (-) | 139 | 15.210 | GGAG | None detected | Hypothetical protein PSPTO3426 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793209), 6e ⁻²⁴ | Conserved hypothetical protein |
| PFL_1982 | <i>prtN</i> | 2212189..2212515 (-) | 108 | 12.260 | GCAAG | Transcriptional regulator PrtN from <i>Pseudomonas aeruginosa</i> pyocin (Q06552); 5e ⁻⁰⁵ | Putative transcriptional regulator PSPTO3425 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793208), 3e ⁻⁴⁰ | Putative phage-related positive regulator |
| PFL_1983 | PFL_1983 | 2212674..2212958 (-) | 94 | 10.986 | GGAG | None detected | Putative DNA-binding Roi-like protein PSPTO3423 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793206); 1e ⁻³³ | Conserved hypothetical phage protein |
| PFL_1984 | PFL_1984 | 2212968..2213552 (-) | 194 | 21.429 | GGAG | None detected | Hypothetical protein PSPTO3422 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793205), 6e ⁻⁴⁹ | Conserved hypothetical protein |
| PFL_1985 | PFL_1985 | 2213627..2213758 | 43 | 4.439 | TAAG | None detected | None detected | Hypothetical protein |
| PFL_1986 | PFL_1986 | 2213809..2214522 (-) | 237 | 26.143 | None selected | Repressor C2 from <i>Salmonella typhimurium</i> bacteriophage ST64T (NP_720299); 5e ⁻²¹ | Repressor protein C1/C2 family PP3033 from <i>Pseudomonas putida</i> KT2440 (AAN68641); 2e ⁻⁶⁴ | Putative LexA-like transcriptional repressor of C1/C2 family (COG1974) |
| 3 ^b | <i>orf3</i> | 2214599..2214871 | 90 | 9.740 | GAGG | None detected | Putative regulatory protein PP3034 from | Conserved hypothetical |

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| PFL_1987 | PFL_1987 | 2215205..2215369 (-) | 54 | 5.197 | GGGA | None detected | <i>Pseudomonas putida</i> KT2440 (NP_745178); 4e ⁻²⁹ | protein |
| PFL_1988 ^c | PFL_1988 | 2215220..2215738 | 172 | 18.872 | AGGA | Hypothetical protein gp14 from <i>Burkholderia</i> phage Bcep176 (ABA60015), 5e ⁻⁰⁷ | Hypothetical protein PSPTO3420 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793203), 2e ⁻⁸⁵ | Hypothetical protein Conserved hypothetical phage protein |
| PFL_1989 | PFL_1989 | 2215930..2218167 | 746 | 83.604 | GGAG | Hypothetical protein ORF002 from <i>Staphylococcus</i> phage 47 (AAV91193), 9e ⁻⁵⁴ | DNA primase domain protein PSPTO3418 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793201); 0.0 | Putative Ssb-like protein (COG5545) |
| PFL_1990 | PFL_1990 | 2218181..2218525 | 114 | 12.595 | GCGAA | None detected | Conserved hypothetical protein PSPTO3417 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (NP_793200), 9e ⁻⁴⁵ | Conserved hypothetical protein |
| PFL_1991 | hol | 2219046..2219390 | 114 | 11.988 | AGGGG | Holin from pyocin locus of <i>Pseudomonas aeruginosa</i> (BAA83151); 5e ⁻²⁸ | Hypothetical protein Pfl01_1137 from <i>Pseudomonas fluorescens</i> Pf0-1 (ABA72880); 2e ⁻⁴⁷ | Holin |
| PFL_1992 | PFL_1992 | 2219596..2220168 | 190 | 20.878 | GGAC | None detected | Conserved hypothetical protein PSPTO3413 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56891), 2e ⁻⁷⁹ | Conserved hypothetical protein |

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|----------|----------|------------------|-----|--------|-------|--|---|---------------------------------------|
| PFL_1993 | PFL_1993 | 2220134..2222185 | 683 | 76.840 | CAGAG | Phage terminase gp15 from <i>Wolbachia</i> phage WO (BAA89640), 6e ⁻¹⁰⁹ | Putative terminase GpA from <i>Pseudomonas putida</i> GB-1 (ABY97103), 0.0 | Phage terminase (COG5525) |
| PFL_1994 | PFL_1994 | 2222187..2222393 | 68 | 7.601 | AAGAG | None detected | Conserved hypothetical protein PSPTO3411 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AA056889), 2e ⁻²³ | Conserved hypothetical protein |
| PFL_1995 | PFL_1995 | 2222393..2223874 | 493 | 55.288 | CAAGG | Phage portal protein from <i>Wolbachia</i> sp. bacteriophage WOcauB1(BAD16786), 1e ⁻¹⁰⁵ | Phage portal protein, lambda family PP_3044 from <i>Pseudomonas putida</i> KT2440 (AE016495), 0.0 | Portal protein (COG5511) |
| PFL_1996 | PFL_1996 | 2223871..2225013 | 380 | 40.265 | AGGAG | Putative protease Z3097 from <i>E.coli</i> O157:H7 prophage CP-933U (AAG57011), 8e ⁻³¹ | Clp protease PSPTO3409 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56887), 7e ⁻¹⁴² | Putative ClpP-like protease (COG0740) |
| PFL_1997 | PFL_1997 | 2225010..2225354 | 114 | 11.885 | AGGAA | None detected | Conserved hypothetical protein PputGB1_1199 from <i>Pseudomonas putida</i> GB-1 (ABY97107), 2e ⁻⁴³ | Conserved hypothetical protein |
| PFL_1998 | PFL_1998 | 2225418..2226413 | 331 | 36.382 | GGAG | Phage protein gp20 from <i>Wolbachia</i> sp. bacteriophage WO (BAA89645), 6e ⁻⁵¹ | Conserved hypothetical protein PSPTO3407 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56885), 8e ⁻¹⁵⁶ | Conserved hypothetical phage protein |

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|-----------------------|----------|------------------|-----|--------|------|--|--|--|
| PFL_1999 | PFL_1999 | 2226416..2226730 | 104 | 11.328 | GGAG | None detected | Conserved hypothetical protein PSPTO3406 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56884), 4e ⁻³⁹ | Conserved hypothetical protein |
| PFL_2000 | PFL_2000 | 2226727..2227380 | 217 | 24.691 | GGAG | Putative phage-related protein from phage VP882 of <i>Vibrio parahaemolyticus</i> 882 (AAS38501), 1e ⁻⁶⁶ | Conserved hypothetical protein PSPTO3405 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56883), 8e ⁻⁷⁵ | Hypothetical phage protein |
| PFL_2001 | PFL_2001 | 2227373..2227885 | 170 | 18.805 | GGTG | Hypothetical protein PRF10 from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83152), 7e ⁻²⁸ | Conserved hypothetical protein PSPTO3404 <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56882), 5e ⁻⁶⁶ | Hypothetical phage protein |
| PFL_2002 ^d | PFL_2002 | 2227882..2228472 | 196 | 20.198 | GGAG | Putative baseplate assembly protein V from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83153), 4e ⁻⁵³ | Putative tail spike protein gpV from <i>Pseudomonas entomophila</i> L48 (AAO56881), 4e ⁻⁶⁷ | Baseplate assembly protein V (COG4540) |
| PFL_2003 | PFL_2003 | 2228537..2228737 | 66 | 7.144 | GGAG | None detected | Conserved hypothetical protein PSPTO3402 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56880), 3e ⁻⁰⁹ | Conserved hypothetical protein |
| PFL_2004 | PFL_2004 | 2228742..2229068 | 108 | 12.072 | GGAG | Putative baseplate assembly protein W from R pyocin region of <i>P.aeruginosa</i> PAO1 | Putative baseplate assembly protein gpW from <i>Pseudomonas</i> | Baseplate assembly protein W (COG3628) |

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|----------|----------|------------------|-----|--------|-------|---|---|-----------------------------------|
| PFL_2005 | PFL_2005 | 2229065..2229946 | 293 | 31.192 | GGAG | (BAA83155), 1e ⁻³⁵ Baseplate J-like protein (baseplate or base of tail fiber) phage phi CTX of <i>P.aeruginosa</i> PAO1 (NP_490617), 3e ⁻⁷¹ | <i>entomophila</i> L48 (PSEEN4162), 6e ⁻⁴³ Putative baseplate J-like protein gpJ from <i>Pseudomonas</i> <i>entomophila</i> L48 (PSEEN4161), 9e ⁻¹¹⁰ | Baseplate J protein (COG3948) |
| PFL_2006 | PFL_2006 | 2229948..2230553 | 201 | 22.268 | GGAG | Baseplate assembly protein gpI from Enterobacteria phage P2 (AAD03285), 2e ⁻⁴⁸ | Tail protein I PSPTO3399 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56877), 4e ⁻⁷⁷ | Tail protein I (COG4385) |
| PFL_2007 | PFL_2007 | 2230550..2231776 | 408 | 42.749 | GGAC | Phage-related tail fibre protein HR2 from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83158), 5e ⁻⁴¹ | Putative tail fiber protein H PSPTO3398 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56876), 3e ⁻⁷² | Tail fiber protein H (COG5301) |
| PFL_2008 | PFL_2008 | 2231776..2232558 | 260 | 25.745 | GGTG | None detected | None detected | Hypothetical protein |
| PFL_2009 | PFL_2009 | 2232704..2233870 | 388 | 42.051 | GGAG | Phage tail sheath protein from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83160), 1e ⁻¹⁵⁶ | Putative baseplate J-like protein gpFI from <i>Pseudomonas</i> <i>entomophila</i> L48 (PSEEN4155), 0.0 | Tail sheath protein (COG3497) |
| PFL_2010 | PFL_2010 | 2233889..2234392 | 169 | 18.403 | GGAG | Contractile tube protein of <i>P.aeruginosa</i> phage PS17 (JC5192), 2e ⁻⁵⁷ | Major tail tube protein PSPTO3395 from <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 (AAO56873), 8e ⁻⁸⁴ | Major tail tube protein (COG3498) |
| PFL_2011 | PFL_2011 | 2234404..2234709 | 101 | 10.981 | AGGAA | Conserved hypothetical protein | Conserved hypothetical protein | Conserved hypothetical |

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|----------|----------|----------------------|-----|--------|------|---|--|------------------------------------|
| PFL_2012 | PFL_2012 | 2234717..2234821 | 33 | 3.964 | GGCG | gp23 from bacteriophage WOcauB1 of <i>Wolbachia</i> sp. wCauB (BAD16807), 3e ^{-05\} | PputW619_4038 from <i>Pseudomonas putida</i> W619 (ACA74518), 8e ⁻³² | protein |
| PFL_2013 | PFL_2013 | 2234851..2237418 | 855 | 88.148 | GGTG | Tail tape measurure protein PRF20 from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83164), 4e ⁻⁴⁷ | Tail tape measurure protein PSPTO3393 from <i>P. syringae</i> pv. <i>tomato</i> DC3000 (AAO56871), 0.0 | Hypothetical protein |
| PFL_2014 | PFL_2014 | 2237427..2238272 | 281 | 29.763 | GGAG | Protein PRF21 from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83165), 3e ⁻⁶⁵ | Hypothetical protein PSPTO3392 from <i>P. syringae</i> pv. <i>tomato</i> DC3000 (AAO56870), 5e ⁻¹¹⁵ | Tape measure protein (COG3494) |
| PFL_2015 | PFL_2015 | 2238247..2238453 | 68 | 7.322 | GGAG | Phage tail protein PRF22 (GpX) from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83166), 6e ⁻²¹ | Tail protein X PSPTO3391 from <i>P. syringae</i> pv. <i>tomato</i> DC3000 (AAO56869), 3e ⁻²⁴ | Tail protein X (COG5004) |
| PFL_2016 | PFL_2016 | 2238450..2238920 (-) | 156 | 16.732 | GGAG | None detected | None detected | Hypothetical protein |
| PFL_2017 | PFL_2017 | 2239001..2240044 | 347 | 38.005 | GGAG | Phage late control gene D protein PRF23(GpD) from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83167), 2e ⁻⁹⁶ | Tail protein D PSPTO3390 from <i>P. syringae</i> pv. <i>tomato</i> DC3000 (AAO56868), 2e ⁻¹⁶⁰ | Late control protein gpD (COG3500) |
| PFL_2018 | lys | 2240071..2240622 | 183 | 19.846 | GGAG | Lytic enzyme PRF24 from R pyocin region of <i>P.aeruginosa</i> PAO1 (BAA83168), 1e ⁻³⁵ | Putative lysozyme PP_3066 from <i>Pseudomonas putida</i> KT2440 (AE016496), 1e ⁻⁸² | Endolysin (COG3179) |
| PFL_2019 | PFL_2019 | 2240619..2241140 | 173 | 18.154 | GGTG | Conserved phage protein gp78 from | Hypothetical protein PSPPH_0666 from | Conserved phage protein |

Burkholderia cepacia *P. syringae* pv.
phage Bcep22 *phaseolicola* 1448a
(AAQ55012), 4e⁻⁰⁵ (AAZ35483), 2e⁻⁴²

^a Only expectation values of 1e⁻⁰⁵ and below were considered as significant matches during BLAST database searches; ^b open reading frame(s) not present in the original genome annotation; ^d open reading frame has an alternative start site.