

Table S3 Percentage of ORFs homologous between bacteriophages of the PAK_P1-like and KPP10-like clades

| Name (<i>number of predicted ORFs</i>) | JG004 | PAK_P1 | PAK_P2 | PAK_P4 | PaP1 | C2_10-Ab1 | PAK_P5 | CHA_P1 | KPP10 | LSL4 | PAK_P3 |
|--|------------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| JG004 (161) | 100^a | 89 | 89 | 88 | 83 | 84 | 23 | 24 | 22 | 23 | 24 |
| PAK_P1 (181) | 79 | 100 | 90 | 90 | 83 | 78 | 24 | 24 | 23 | 24 | 24 |
| PAK_P2 (175) | 82 | 93 | 100 | 95 | 82 | 77 | 25 | 25 | 23 | 24 | 25 |
| PAK_P4 (174) | 82 | 94 | 95 | 100 | 83 | 78 | 25 | 25 | 24 | 25 | 25 |
| PaP1 (157) | 85 | 96 | 91 | 92 | 100 | 84 | 24 | 24 | 24 | 25 | 25 |
| C2_10-Ab1 (157) | 87 | 90 | 86 | 86 | 84 | 100 | 24 | 24 | 24 | 25 | 25 |
| PAK_P5 (164) | 23 | 27 | 26 | 27 | 23 | 23 | 100 | 96 | 83 | 94 | 93 |
| CHA_P1 (164) | 23 | 26 | 26 | 26 | 23 | 23 | 96 | 100 | 84 | 93 | 94 |
| KPP10 (146) | 25 | 28 | 28 | 29 | 26 | 25 | 93 | 95 | 100 | 91 | 95 |
| LSL4 (165) | 22 | 27 | 25 | 26 | 24 | 24 | 93 | 92 | 81 | 100 | 91 |
| PAK_P3 (165) | 23 | 27 | 26 | 27 | 24 | 24 | 93 | 93 | 84 | 91 | 100 |

^a: in bold, percentages corresponding to the two clades