**S23 Table.** Statistical analysis of nematode survival after exposure to cry-toxin-expressing *B. thuringiensis*1

|  |  |  |  |
| --- | --- | --- | --- |
| **Reference** | **Comparison** | ***Z*** | ***P*** |
| BT-679\_Cry- | BT-679\_Cry+ | 3.44 | **0.0023** |
|  | BT-679\_Cry-\_+14 | 3.05 | **0.0092** |
|  | BT-679\_Cry-\_+21 | 1.62 | 0.4190 |
|  | BT-679\_Cry-\_0 | 0.04 | >0.99 |
| BT-679\_Cry+ | BT-679\_Cry-\_+14 | 0.35 | >0.99 |
|  | BT-679\_Cry-\_+21 | 1.77 | 0.3045 |
|  | BT-679\_Cry-\_0 | 3.35 | **0.0032** |

1 Non-parametric comparisons with control using Dunn method for joint ranking, as implemented in JMP 9.0.2 (SAS Institute Inc.). Significant probabilities are given in bold. Strain abbreviations: BT-679\_Cry-, BT-679 without plasmid which contains the toxin genes *cry14Aa1* and *cry21Aa2*; BT-679\_Cry+, BT-679 wildtype with the toxin-containing plasmids; BT-679\_Cry-\_+14, BT-679\_Cry- with a *cry14Aa1*-expressing vector; BT-679\_Cry-\_+21, BT-679\_Cry- with a *cry21Aa2*-expressing vector; BT-679\_Cry-\_+0, BT-679\_Cry- with a *rfp*-expressing vector as a negative control. The data is shown in S6 Data.