# **Supplementary Table 6: Classification of B91 strains with the use of CLARK.** B91 strain genomes were compared against genomes for the *Campylobacter* species - *coli, concisus, corcagienisis, cuniculorum, curvus, fetus, gracilis, hepaticus, hominis, iguaniorum, insulaenigrae, jejuni, lari, mucosalis, peloridis, rectus, sputorum, subantarcticus, showae, upsaliensis, ureolyticus, volucris*. More information on the genomes in the methods. Classification of NA indicates that CLARK was unable to classify the scaffold.

|  |  |  |  |
| --- | --- | --- | --- |
| **Isolate** | **Scaffold** | **length (bp)** | **Clark classification** |
| B91\_SC | scaffold1 | 2160580 | *C. showae* |
| B91\_SCBr | scaffold1 | 1586030 | *C. showae* |
| scaffold2 | 359805 | *C. showae* |
| scaffold3 | 1611 | NA |
| scaffold4 | 191055 | *C. showae* |
| scaffold5 | 19196 | *C. showae* |
| B91\_TPP | scaffold1 | 2131013 | *C. showae* |
| scaffold2 | 14986 | *C. showae* |
| scaffold3 | 916 | NA |
| scaffold4 | 847 | NA |
| scaffold5 | 751 | *C. concisus* |
| scaffold6 | 605 | NA |

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