Table S2: GenBank accession numbers of sequences obtained in this study in alphabetical order according to bacterial species

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| **Bacterial species** | **Accession number** |
| *Achromobacter xylosoxidans* | MT664080 |
| *Acinetobacter haemolyticus* | MT664081 |
| *Bacillus* sp. | MT664082 |
| *Bacillus* sp. | MT664083 |
| *Brachybacterium* sp. | MT664084 |
| *Chryseobacterium culicis* | MT664085 |
| *Cloacibacterium normanense* | MT664086 |
| *Corynebacterium xerosis* | MT664087 |
| *Deinococcus aquaticus* | MT664088 |
| *Deinococcus indicus* | MT664089 |
| *Desemzia* sp. | MT664090 |
| *Devriesea agamarum* | MT664091 |
| *Devriesea agamarum* | MT664092 |
| *Devriesea agamarum* | MT664093 |
| *Exiguobacterium acetylicum* | MT664094 |
| *Exiguobacterium* sp. | MT664095 |
| *Micrococcus* sp. | MT664096 |
| *Ochrobactrum* sp. | MT664097 |
| *Pseudomonas* sp. | MT664098 |
| *Serratia marcescens* | MT664099 |
| *Sporosarcina thermotolerans* | MT664100 |
| *Staphylococcus kloosii* | MT664101 |
| *Stenotrophomonas rhizophila* | MT664102 |
| *Uruburuella testudinis* | MT664103 |
| *Uruburuella testudinis* | MT664104 |