|  |  |  |
| --- | --- | --- |
| **Table S1.** |  | Physicochemical parameters of lake water. |
| **pH** | **NH3-N** | **TP** | **COD** | **BOD5** | **TOC** | **nitrates** | **nitrite** | **sulfates** | **K** | **Fe** | **Na** | **Cd** | **Pb** | **Cr** | **As** | **Ni** | **Ca** | **Mg** | **TDS** |
|  | mg/L |
| 8.1 | 0.142 | 0.02 | 11 | 2.3 | 7.7 | 0.805 | - | 20.9 | 3.15 | 0.0240 | 18.6 | - | - | - | 0.00157 | 0.00160 | 37.0 | 8.57 | 71 |
|  |  | -, means under limit of detection |
|  |  |  |
| **Table S2.** |  | Molecular identification and functional profile of culturable endobacterial isolates using 16S rRNA as query sequences. |
| **Origin** | **Isolate name** | **Closet relative (NCBI)** | **Closest type strain (accession number)** | **%Similarity (EzTaxon)** | **Phylum** | **Family** |
| leaf | EGA240353 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240354 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.86 | Firmicutes | Bacillaceae |
| EGA240355 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.86 | Firmicutes | Bacillaceae |
| EGA240356 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.51 | Firmicutes | Bacillaceae |
| EGA240357 | *Cytophaga* sp. VM1T | *Cytophaga massiliensis* (EF394924) | 99.21 | Bacteroidetes | Cytophagaceae |
| EGA240358 | *Roseateles depolymerans* | *Roseateles depolymerans* (CP013729) | 99.72 | Proteobacteria | norank |
| EGA240359 | *Herbaspirillum huttiense* | *Herbaspirillum aquaticum* (NJGV01000018) | 99.93 | Proteobacteria | Oxalobacteraceae |
| EGA240360 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240361 | *Acinetobacter soli* | *Acinetobacter soli* (APPU01000012) | 99.30 | Proteobacteria | Moraxellaceae |
| EGA240363 | *Acinetobacter* sp. | *Acinetobacter seifertii* (KB851199) | 99.15 | Proteobacteria | Moraxellaceae |
| EGA240364 | *Acinetobacter seifertii* | *Acinetobacter seifertii* (KB851199) | 99.21 | Proteobacteria | Moraxellaceae |
| EGA240365 | *Agrobacterium vitis* |  CP000633\_s (CP000633) | 98.90 | Proteobacteria | Rhizobiaceae |
| EGA240366 | *Sphingomonas azotifigens* | *Sphingomonas azotifigens* (BCTR01000108) | 99.49 | Proteobacteria | Sphingomonadaceae |
| EGA240367 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.44 | Firmicutes | Bacillaceae |
| EGA240368 | *Sphingomonas trueperi* | *Sphingomonas trueperi* (X97776) | 99.56 | Proteobacteria | Sphingomonadaceae |
| EGA240369 | *Microbacterium* sp HBUM179310 | *Microbacterium hydrothermale* (HM222660) | 98.71 | Actinobacteria | Microbacteriaceae |
| stem | EGA240370 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240371 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.93 | Firmicutes | Bacillaceae |
| EGA240372 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.51 | Firmicutes | Bacillaceae |
| EGA240373 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240374 | *Zavarzinia compransoris* | *Zavarzinia compransoris* (JX986958) | 94.65 | Proteobacteria | Acetobacteraceae |
| EGA240375 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.93 | Firmicutes | Bacillaceae |
| EGA240376 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.72 | Firmicutes | Bacillaceae |
| EGA240377 | *Stenotrophomonas* sp. | CP026001\_s (CP026001) | 99.44 | Proteobacteria | Xanthomonadaceae |
| root | EGA240378 | *Phycicoccus jejuensis* | *Phycicoccus jejuensis* (JOEE01000020) | 99.72 | Actinobacteria | Lntrasporangiaceae |
| EGA240379 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.79 | Firmicutes | Bacillaceae |
| EGA240380 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240381 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.65 | Firmicutes | Bacillaceae |
| EGA240382 | *Rhizobium sp.* strain MERSZ-6 | *Rhizobium wuzhouense* (MG857114) | 99.78 | Proteobacteria | Rhizobiaceae |
| EGA240383 | *Bacillus* sp. S2(2014) | *Bacillus tequilensis* (AYTO01000043) | 99.72 | Firmicutes | Bacillaceae |
| EGA240384 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.58 | Firmicutes | Bacillaceae |
| EGA240386 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.70 | Firmicutes | Bacillaceae |
| EGA240387 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.86 | Firmicutes | Bacillaceae |
| EGA240388 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.51 | Firmicutes | Bacillaceae |
| EGA240389 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.51 | Firmicutes | Bacillaceae |
| EGA240390 | *Bacillus amyloliquefaciens* | *Bacillus tequilensis* (AYTO01000043) | 99.93 | Firmicutes | Bacillaceae |
| EGA240391 | *Microbacterium* sp. HBUM179310 | *Microbacterium hydrothermale* (HM222660) | 98.63 | Actinobacteria | Microbacteriaceae |
| EGA240393 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.72 | Firmicutes | Bacillaceae |
| EGA240394 | *Rhizobium* sp. HBU08115 | *Rhizobium oryzicola* ([JX446583](https://www.ezbiocloud.net/16SrRNA?ac=JX446583)) | 99.45 | Proteobacteria | Rhizobiaceae |
| EGA240395 | *Mycobacterium* sp. M26 | LN929908\_s (LN929908) | 99.36 | Actinobacteria | Mycobacteriaceae |
| EGA240396 | *Rhizobium* sp. | *Rhizobium aquaticum* (KM083136) | 99.61 | Proteobacteria | Rhizobiaceae |
| EGA240397 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.44 | Firmicutes | Bacillaceae |
| EGA240398 | *Bacillus velezensis* | *Bacillus siamensis* ([AJVF01000043](https://www.ezbiocloud.net/16SrRNA?ac=AJVF01000043)) | 99.86 | Firmicutes | Bacillaceae |
| EGA240399 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.93 | Firmicutes | Bacillaceae |
| EGA240400 | *Priestia megaterium* | *Priestia aryabhattai* (EF114313) | 99.86 | Firmicutes | Bacillaceae |
| EGA240401 | *Chryseobacterium* sp. zd2 | *Chryseobacterium massiliae* (AF531766) | 99.40 | Bacteroidetes | Weeksellaceae |
| EGA240402 | Uncultured *Xanthobacter* sp. | *Aquabacter spiritensis* (FR733686) | 98.25 | Proteobacteria | Xanthobacteraceae |
| EGA240403 | *Aeromonas veronii* | *Aeromonas veronii* (CDDK01000015) | 99.02 | Proteobacteria | Aeromonadaceae |
| EGA240404 | *Pseudomonas oryzihabitans* | *Pseudomonas oryzihabitans* (BBIT01000012) | 99.02 | Proteobacteria | Pseudomonadaceae |
| EGA240405 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.37 | Firmicutes | Bacillaceae |
| EGA240406 | *Fictibacillus* sp. | Fictibacillus halophilus (KP265300) | 99.93 | Firmicutes | Bacillaceae |
| EGA240407 | *Bacillus* sp. (in: Bacteria) | *Bacillus tequilensis* (AYTO01000043) | 99.65 | Firmicutes | Bacillaceae |
| EGA240408 | *Roseomonas* sp. | *Roseomonas lacus* (AJ786000) | 96.27 | Proteobacteria | Acetobacteraceae |
| EGA240409 | *Bacillus subtilis* | *Bacillus tequilensis* (AYTO01000043) | 99.51 | Firmicutes | Bacillaceae |
| EGA240410 | *Bacillus* sp MN13 | *Bacillus tequilensis* (AYTO01000043) | 99.37 | Firmicutes | Bacillaceae |

|  |  |
| --- | --- |
| **Table S3** | Microbiological validation of OTUs of endophytic bacteriome in *E. crassipes*.  |

|  |  |  |
| --- | --- | --- |
| **Genus** | **Bacteriome based on 16S rRNA gene sequencing** | **Microbiological tools**  |
| Root | Stem | Leaf | Total OTUs | Frequency (%) | Root | Stem | Leaf | Total isolates (n) | Frequency (%) |
| *Bacillus* | 3598 | 82 | 568 | 4248 | 100 | 18 | 6 | 6 | 30 | 100 |
| *Cytophaga* | – | – | – | – | – | – | – | 1 | 1 | 33.3 |
| *Roseateles* | – | – | – | – | – | – | – | 1 | 1 | 33.3 |
| *Herbaspirillum* | 58 | 5932 | 8872 | 14862 | 100 | – | – | 1 | 1 | 33.3 |
| *Acinetobacter* | 0 | 1 | 5 | 6 | 66.7 | – | – | 3 | 3 | 33.3 |
| *Agrobacterium* | – | – | – | – | – | – | – | 1 | 1 | 33.3 |
| *Sphingomonas* | 874 | 4716 | 15406 | 20996 | 100 | – | – | 2 | 2 | 33.3 |
| *Microbacterium* | 141 | 98 | 863 | 1102 | 100 | 1 | – | 1 | 2 | 66.7 |
| *Zavarzinia* | – | – | – | – | – | – | 1 | – | 1 | 33.3 |
| *Stenotrophomonas* | 86 | 112 | 284 | 482 | 100 | – | 1 | – | 1 | 33.3 |
| *Phycicoccus* | 10 | 15 | 135 | 160 | 100 | 1 | – | – | 1 | 33.3 |
| *Allorhizobium-Neorhizobium-Pararhizobium-Rhizobium* | 558 | 597 | 15891 | 17046 | 100 | 3 | – | – | 3 | 33.3 |
| *Mycobacterium* | 937 | 84 | 246 | 1267 | 100 | 1 | – | – | 1 | 33.3 |
| *Priestia* | – | – | – | – | – | 1 | – | – | 1 | 33.3 |
| *Chryseobacterium* | 7 | 195 | 1007 | 1209 | 100 | 1 | – | – | 1 | 33.3 |
| *Aquabacter* | – | – | – | – | – | 1 | – | – | 1 | 33.3 |
| *Aeromonas* | 123 | 32 | 90 | 245 | 100 | 1 | – | – | 1 | 33.3 |
| *Pseudomonas* | 141 | 336 | 1981 | 2458 | 100 | 1 | – | – | 1 | 33.3 |
| *Fictibacillus* | 45 | 0 | 1 | 46 | 66.7 | 1 | – | – | 1 | 33.3 |
| *Roseomonas* | 1357 | 50 | 119 | 1526 | 100 | 1 | – | – | 1 | 33.3 |

|  |  |
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| **Table S4** | *In vitro* functional profile of bacterial isolates from *E. crassipes* on plant growth promotions traits. |

|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Isolate name** | **Genus** | **ACC deaminase** | **Organic phosphorus solubilization** | **Inorganic phosphorus solubilization** | **Siderophore** | **IAA** | **Nitrogen fixation** | **Cellulase** | **Chitinase** | **Protease** |
| EGA240353 | *Bacillus*  | + |  |  |  |  |  | + | + | + |
| EGA240354 | *Bacillus* | + |  |  |  | + |  | + |  |  |
| EGA240355 | *Bacillus* | + |  | + |  | + |  | + | + | + |
| EGA240356 | *Bacillus*  | – |  | + |  | + | + |  |  | + |
| EGA240357 | *Cytophaga*  | – | + | + | + | + |  |  | + |  |
| EGA240358 | *Roseateles*  | + |  | + |  | + | + |  |  |  |
| EGA240359 | *Herbaspirillum*  | – |  |  |  |  | + |  |  |  |
| EGA240360 | *Bacillus* | – |  | + |  |  |  | + |  | + |
| EGA240361 | *Acinetobacter*  | – | + |  | + | + |  |  |  |  |
| EGA240363 | *Acinetobacter*  | – | + | + | + | + |  |  |  |  |
| EGA240364 | *Acinetobacter* | + | + | + | + | + |  |  |  |  |
| EGA240365 | *Agrobacterium*  | – |  | + |  |  |  |  |  |  |
| EGA240366 | *Sphingomonas*  | – | + | + |  |  |  |  |  |  |
| EGA240367 | *Bacillus* | + |  |  |  | + |  |  |  |  |
| EGA240368 | *Sphingomonas*  | + | + | + |  |  |  |  |  |  |
| EGA240369 | *Microbacterium* | – | + | + |  |  |  |  |  |  |
| EGA240370 | *Bacillus* | + |  | + |  | + |  | + |  | + |
| EGA240371 | *Bacillus* | + |  | + |  |  |  | + | + |  |
| EGA240372 | *Bacillus*  | + |  |  |  | + |  | + |  | + |
| EGA240373 | *Bacillus*  | – |  |  |  | + |  |  |  |  |
| EGA240374 | *Zavarzinia*  | + | + | + |  |  |  | + |  |  |
| EGA240375 | *Bacillus* | – |  |  |  | + |  |  | + | + |
| EGA240376 | *Bacillus* | – |  | + |  | + |  | + |  |  |
| EGA240377 | *Stenotrophomonas*  | – |  |  | + | + |  |  |  | + |
| EGA240378 | *Phycicoccus*  | – |  | + | + | + |  |  |  |  |
| EGA240379 | *Bacillus*  |  |  | + |  | + |  | + | + | + |
| EGA240380 | *Bacillus*  |  |  | + | + | + |  |  | + | + |
| EGA240381 | *Bacillus*  | + |  |  |  |  |  | + |  |  |
| EGA240382 | *Rhizobium*  | + |  | + |  |  | + |  |  |  |
| EGA240383 | *Bacillus*  |  |  |  |  | + |  | + |  | + |
| EGA240384 | *Bacillus*  | + |  |  |  | + | + |  | + | + |
| EGA240386 | *Bacillus*  |  |  | + |  | + |  | + |  | + |
| EGA240387 | *Bacillus*  |  |  |  |  | + |  | + | + |  |
| EGA240388 | *Bacillus*  | + |  | + | + |  |  | + |  | + |
| EGA240389 | *Bacillus*  |  |  |  |  | + |  |  | + |  |
| EGA240390 | *Bacillus*  | + |  | + |  | + |  |  | + |  |
| EGA240391 | *Microbacterium*  | + | + | + | + |  |  |  |  |  |
| EGA240393 | *Bacillus*  |  |  |  |  |  |  | + | + | + |
| EGA240394 | *Rhizobium*  | + |  | + |  |  | + |  |  |  |
| EGA240395 | *Mycobacterium*  |  | + |  | + |  |  |  |  |  |
| EGA240396 | *Rhizobium*  | + |  | + |  | + | + |  |  |  |
| EGA240397 | *Bacillus* |  |  |  |  |  |  | + | + | + |
| EGA240398 | *Bacillus*  | + |  | + |  | + | + | + | + | + |
| EGA240399 | *Bacillus*  |  |  |  | + | + |  | + | + |  |
| EGA240400 | *Priestia* | + | + | + |  |  |  |  |  |  |
| EGA240401 | *Chryseobacterium*  | + | + |  |  |  |  |  |  |  |
| EGA240402 | *Aquabacte* | + | + | + |  |  | + |  | + |  |
| EGA240403 | *Aeromonas*  | + | + |  |  |  |  |  |  |  |
| EGA240404 | *Pseudomonas*  | + | + | + | + | + | + | + |  | + |
| EGA240405 | *Bacillus*  |  |  | + |  |  |  | + |  | + |
| EGA240406 | *Fictibacillus*  | + |  |  |  | + | + |  | + |  |
| EGA240407 | *Bacillus*  |  |  | + |  |  |  | + | + |  |
| EGA240408 | *Roseomonas*  | + | + | + | + | + |  |  |  |  |
| EGA240409 | *Bacillus* |  |  |  |  | + |  |  | + | + |
| EGA240410 | *Bacillus*  |  |  |  |  | + |  | + | + | + |