Table S3: Common and unique isolates obtained from individuals.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | common bacteria and their % | | unique bacteria and their % | |
|  | *name of bacteria* | % | *name of bacteria* | % |
| Individual1 | *Staphylococcus warneri* | 15.4 | *Staphylococcus vitulinus* | 2.6 |
|  | *Staphylococcus epidermidis* | 7.7 |  |  |
|  | *Staphylococcus lentus* | 26 |  |  |
|  | *Staphylococcus arlettae* | 10.3 |  |  |
|  | *Staphylococcus aureus* | 5.1 |  |  |
|  | *Bacillus megaterium* | 2.6 |  |  |
|  | *Bacillus pumilus* | 2.6 |  |  |
|  | *Bacillus subtilis* | 2.6 |  |  |
|  | [*Bacillus licheniformis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_163119169) | 7.7 |  |  |
|  | *Pseudomonas stutzeri* | 5.1 |  |  |
|  | *Acinetobacter radioresistens* | 2.1 |  |  |
|  | *Bacillus sonorensis* | 5.1 |  |  |
|  | [*Corynebacterium striatum*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_237753946) | 5.1 |  |  |
| Individual2 | *Staphylococcus warneri* | 8.5 | *Citrobacter freundii* | 10.5 |
|  | *Staphylococcus epidermidis* | 21 | *Escherichia vulneris* | 5.3 |
|  | *Staphylococcus lentus* | 10.5 |  |  |
|  | *Staphylococcus arlettae* | 10.5 |  |  |
|  | *Bacillus pumilus* | 5.3 |  |  |
|  | *Bacillus subtilis* | 5.3 |  |  |
|  | *Bacillus amyloliquefaciens* | 5.3 |  |  |
|  | *Acinetobacter radioresistens* | 5.3 |  |  |
|  | *Exiguobacterium acetylicum* | 2 |  |  |
|  | *Staphylococcus aureus* | 10.5 |  |  |
| Individual3 | *Staphylococcus hominis* | 5.3 | [*Pseudonocardia*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_485037584) | 5.3 |
|  | *Staphylococcus epidermidis* | 10.4 | *Kocuriarhizophila* | 5.3 |
|  | *Staphylococcus haemolyticus* | 5.3 |  |  |
|  | *Staphylococcus aureus* | 5.3 |  |  |
|  | *Pseudomonas stutzeri* | 26.6 |  |  |
|  | *Acinetobacter radioresistens* | 15.7 |  |  |
|  | [*Corynebacterium striatum*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_237753946) | 10.2 |  |  |
|  | *Exiguobacterium acetylicum* | 5.3 |  |  |
|  | *Bacillus cereus* | 5.3 |  |  |
| Individual4 | [*Oceanobacillusiheyensis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_23097455) | 2 | *Geobacillus stearothermophilus* | 2 |
|  | *Corynebacterium striatum* | 4 | [*Bacillus thuringiensis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_49476684) | 20 |
|  | *Staphylococcus hominis* | 4 | [*Aerococcusurinae*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_326802614) | 4 |
|  | *Staphylococcus epidermidis* | 4 |  |  |
|  | *Staphylococcus haemolyticus* | 4 |  |  |
|  | *Bacillus subtilis* | 20 |  |  |
|  | *Bacillus amyloliquefaciens* | 20 |  |  |
|  | [*Bacillus licheniformis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_163119169) | 4 |  |  |
|  | *Acinetobacter radioresistens* | 4 |  |  |
|  | *Escherichia vulneris* | 4 |  |  |
|  | *Cronobactersakazakii* | 4 |  |  |
| Individual5 | *Bacillus subtilis* | 12.5 | *Staphylococcus sciuri* | 25 |
|  | [*Bacillus licheniformis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_163119169) | 25 | *Bacillus velezensis* | 12.5 |
|  | *Exiguobacteriumacetylicum* | 12.5 |  |  |
|  | *Bacillus sonorensis* | 12.5 |  |  |
| Individual6 | *Pseudomonas stutzeri* | 16.7 | *Planococcus* | 16.7 |
|  | *Exiguobacteriumacetylicum* | 16.7 | *Paenibacillus mucilaginosus* | 16.7 |
|  | *Bacillus cereus* | 16.5 |  |  |
|  | *Bacillus firmus* | 16.7 |  |  |
| Individual7 | *Staphylococcus pasteuri* | 7.1 | *Bacillus pumilus* | 7.3 |
|  | [*Bacillus licheniformis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_163119169) | 7.1 | *Exiguobacterium profundum* | 16.5 |
|  | *Pseudomonas stutzeri* | 35.7 | *Exiguobacterium mexicanum* | 12 |
|  | *Bacillus firmus* | 14.3 |  |  |
| Individual8 | *Staphylococcus aureus* | 7.7 | *Cronobacter sakazakii* | 7.7 |
|  | *Bacillus megaterium* | 7.7 | *Bacillus tequilensis* | 7.7 |
|  | *Bacillus subtilis* | 7.7 |  |  |
|  | *Bacillus amyloliquefaciens* | 15.4 |  |  |
|  | [*Bacillus licheniformis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_163119169) | 7.7 |  |  |
|  | *Exiguobacteriummexicanum* | 28.8 |  |  |
|  | *Bacillus cereus* | 9.6 |  |  |
| Individual9 | *Staphylococcus lentus* | 16.7 |  |  |
|  | *Staphylococcus pasteuri* | 16.7 |  |  |
|  | *Pseudomonas stutzeri* | 16.7 |  |  |
|  | *Acinetobacter radioresistens* | 16.7 |  |  |
|  | *Exiguobacteriumacetylicum* | 16.7 |  |  |
|  | *Bacillus cereus* | 16.5 |  |  |
| Individual10 | *Staphylococcus epidermidis* | 25 |  |  |
|  | *Staphylococcus haemolyticus* | 25 |  |  |
|  | *Staphylococcus aureus* | 20 |  |  |
|  | *Exiguobacterium acetylicum* | 15 |  |  |
|  | *Bacillus cereus* | 15 |  |  |
| Individual 11 | *Cronobacter sakazakii* | 20 | [*Staphylococcus saprophyticus*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_73661309) | 30 |
|  | *Pseudomonas stutzeri* | 15 |  |  |
|  | *Bacillus cereus* | 15 |  |  |
|  | [*Oceanobacillus iheyensis*](https://blast.ncbi.nlm.nih.gov/Blast.cgi#alnHdr_23097455) | 20 |  |  |