**Supplementary Table 4:** **Appearance of identified bacterial strains in NDD patient subgroups (different diagnoses) and Control group**.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Bacterial species obtained with** **universal primers** | MSDD (%)n=12 | PDD-NOS (%)n=10 | ERLD (%)n=9 | CHA (%)n=5 | CTRL (%)n=28 |
| *E. ramosum* strain JCM 1298 | 41.7 | 10,0 | 55.6 | 40,0 | 25,0 |
| *I. bartlettii* strain WAL 16138 | 100,0\* | 70,0 | 100,0 | 100,0 | 64.3 |
| *R. ilealis* strain CRIB (2) | 91.7\*\* | 80,0\* | 88.9\* | 60,0 | 32.1 |
| *D. guttoideum* strain DSM 402 | 16.7 | 30,0 | 55.6\*\* | 40,0 | 3.6 |
| **Bacterial species obtained with** **LB-specific primers** |  |  |  |  |  |
| *E. faecalis* strain NBRC 100480 | 58.3 | 60,0 | 55.6 | 60,0 | 89.3 |
| *E. rectale* strain ATCC 33656 | 33.3 | 20,0 | 22.2 | / | 46.4 |
| *S. pasteurianus* | /\* | 20,0 | 33.3 | / | 42.9 |
| *B. pullicaecorum* strain 25-3 | /\* | 20,0 | 33.3 | / | 42.9 |
| *D. invisus* strain JCM 17566 | /\* | 20,0 | 33.3 | / | 42.9 |
| *R. champanellensis* strain 18P13 | 16.7 | 20,0 | 11.1 | / | 53.6 |
| *E. gallinarum* strain LMG 13129 | 16.7 | 20,0 | 11.1 | / | 53.6 |
| *L. sakei* subsp. *carnosus* strain CCUG 31331 | 41.7 | 30,0 | 22.2\* | 20,0 | 67.9 |
| *F. prausnitzii* strain ATCC 27768  | 41.7 | 30,0 | 22.2\* | 20,0 | 67.9 |
| *L. rhamnosus* strain NBRC 3425 | /\*\* | 20,0 | 11.1 | 20,0 | 42.9 |
| *L. sakei* (2) | 50,0 | 60,0 | 11.1\*\* | 80,0 | 75,0 |
| *D. invisus* strain JCM 17566 (2) | 41.7\* | 40,0\* | 33.3\* | 40,0 | 82.1 |
| **Bacterial species obtained with** **BB-specific primers** |  |  |  |  |  |
| *B. pseudocatenulatum* strain B1279 | 16.7 | 10,0 | 22.2 | / | 35.7 |
| *B. adolescentis* strain ATCC 15703 | / | / | / | / | 28.6 |
| *B. animalis* subsp. *lactis* strain YIT 4121 | 16.7 | 40,0\*\* | 11.1 | / | / |
| *B. faecale*  | 91.7 | 70,0\* | 66.7\* | 80,0 | 100 |
| *B. longum* subsp. *infantis* strain ATCC 15697 | 50,0 | 60,0 | 55.6 | 40,0 | 71.4 |
| *B. stercoris* strain Eg1 16S ribosomal | 66.7\*\* | 60,0\*\* | 66.7\* | 80,0 | 100,0 |

Comparisons between patient subgroups (different diagnoses) and Control group for each bacterial strain were performed with Pearson’s Chi-square test. n – number of participants per group. Results were considered statistically significant at P<0.05. \*P<0.05; \*\*P<0.01