|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **GENUS** | **C** | **CD** | **p** | **FDR** |
| **Roseburia** | 4.366 | 1.127 | 0.012 | 1 |
| **Sutterella** | 0.175 | 1.846 | 0.014 | 1 |
| **Terrahaemophilus** | 0.016 | 0.108 | 0.015 | 1 |
| **Oribacterium** | 0.000 | 0.007 | 0.027 | 1 |
| **Aggregatibacter** | 0.001 | 0.011 | 0.036 | 1 |
| **Eubacterium** | 7.756 | 3.297 | 0.037 | 1 |
| **Malikia** | 0.000 | 0.004 | 0.040 | 1 |
| **Subdoligranulum** | 1.107 | 0.304 | 0.045 | 1 |

Supplementary table 1 Bacterial genera with significant (p=uncorrected, unpaired two tailed T test) difference in abundance between controls (C) and treatment naïve pediatric Crohn disease (CD) patients. The numbers represent percent values of abundance. FDR: false dicovery rate