

Major components

This table lists the most abundant types in the sample. For a bacterial culture or product, we interpret this as the list of types present at meaningful levels. Together these types make up at least 90% of the sample.

▼ Expand or collapse this section

Type	Percent of Community	Phylum	Family	Genus	Species
5	36.454	Proteobacteria	Rhodobacteraceae	Rhodobacter	Rhodobacter_sphaeroides
6	20.219	Bacteroidota	Dysgonomonadaceae	Dysgonomonas	uncultured_Dysgonomonas
13	7.720	Proteobacteria	Pseudomonadaceae	Pseudomonas	Pseudomonas_sp.
22	5.533	Firmicutes	Clostridiaceae	Clostridium sensu stricto 3	uncultured_bacterium
27	5.143	Proteobacteria	Comamonadaceae	Delftia	Delftia_acidovorans
25	4.830	Firmicutes	Oscillospirales	Hydrogenoanaerobacterium	uncultured_Firmicutes
28	4.726	Firmicutes	SRB2	SRB2	uncultured_bacterium
38	2.721	Firmicutes	SRB2	SRB2	uncultured_bacterium
8	2.526	Firmicutes	Peptostreptococcales-Tissierellales	Alkaliphilus	uncultured_bacterium
7	2.109	Firmicutes	Thermotaleaceae	Geosporobacter	Geosporobacter_subterraneus
19	1.823	Firmicutes	Christensenellaceae	Christensenellaceae R-7 group	uncultured_Firmicutes
16	0.976	Firmicutes	Thermotaleaceae	Geosporobacter	uncultured_Firmicutes
18	0.937	Firmicutes	Oscillospirales	Hydrogenoanaerobacterium	uncultured_bacterium