

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
2	32.361	Proteobacteria	Rhodobacteraceae	Rhodobacter	Rhodobacter_sphaeroides
4	23.356	Proteobacteria	Xanthobacteraceae	Rhodoplanes	iron-reducing_bacterium
6	11.864	Firmicutes	Thermotaleaceae	Geosporobacter	Geosporobacter_subterraneus
5	9.177	Firmicutes	SRB2	SRB2	iron-reducing_bacterium
13	3.974	Proteobacteria	Pseudomonadaceae	Pseudomonas	Pseudomonas_sp.
19	3.888	Firmicutes	Clostridiaceae	Clostridium sensu stricto 3	uncultured_bacterium
8	3.573	Bacteroidota	Dysgonomonadaceae	Dysgonomonas	uncultured_Dysgonomonas
17	2.058	Firmicutes	SRB2	SRB2	uncultured_bacterium
22	1.687	Firmicutes	Christensenellaceae	Christensenellaceae R-7 group	uncultured_Firmicutes
16	1.515	Proteobacteria	Comamonadaceae	Delftia	Delftia_acidovorans
31	1.058	Firmicutes	Desulfotomaculales	Desulfotomaculum	Desulfotomaculum_ruminis
20	0.829	Firmicutes	Oscillospirales	Hydrogenoanaerobacterium	uncultured_bacterium