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Note: Where there are blanks in the table below, the taxonomy could not be resolved in finer detail.
Note: The percentages listed represent the relative abundance of each taxon. This summary is based off of normalised data. Because of limitations in the way the samples are processed, we cannot reliably obtain species level resolution. As such, the data shown are collapsed at the genus level.
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Kingdom	Phylum	Class	Order	Family	Genus	Relative Abundance (%)
Bacteria	Bacteroidetes	Bacteroidia	Bacteroidales	Bacteroidaceae	<i>Bacteroides</i>	56.09
Bacteria	Bacteroidetes	Bacteroidia	Bacteroidales	[Paraprevotellaceae]	<i>Paraprevotella</i>	12.44
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	---	5.81
Bacteria	Bacteroidetes	Bacteroidia	Bacteroidales	Porphyromonadaceae	<i>Parabacteroides</i>	4.68
Bacteria	Firmicutes	Clostridia	Clostridiales	Veillonellaceae	<i>Phascolarctobacterium</i>	4.59
Bacteria	Firmicutes	Clostridia	Clostridiales	Ruminococcaceae	---	2.68
Bacteria	Firmicutes	Clostridia	Clostridiales	Ruminococcaceae	<i>Faecalibacterium</i>	1.36
Bacteria	Firmicutes	Clostridia	Clostridiales	Clostridiaceae	<i>Clostridium</i>	1.23
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	[<i>Ruminococcus</i>]	1.18
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Lachnospira</i>	1.18
Bacteria	Proteobacteria	Deltaproteobacteria	Desulfovibrionales	Desulfovibrionaceae	<i>Bilophila</i>	1.09
Bacteria	Firmicutes	Erysipelotrichi	Erysipelotrichales	Erysipelotrichaceae	---	0.86
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Blautia</i>	0.73
Bacteria	Bacteroidetes	Bacteroidia	Bacteroidales	S24-7	---	0.73
Bacteria	Firmicutes	Clostridia	Clostridiales	Ruminococcaceae	<i>Oscillospira</i>	0.68
Bacteria	Proteobacteria	Gammaproteobacteria	Enterobacteriales	Enterobacteriaceae	---	0.55
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Anaerostipes</i>	0.50
Bacteria	Firmicutes	Erysipelotrichi	Erysipelotrichales	Erysipelotrichaceae	[<i>Eubacterium</i>]	0.45
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Dorea</i>	0.45
Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Coproccoccus</i>	0.45
Bacteria	Proteobacteria	Betaproteobacteria	Burkholderiales	Alcaligenaceae	<i>Achromobacter</i>	0.41
Bacteria	Firmicutes	Clostridia	Clostridiales	Ruminococcaceae	<i>Ruminococcus</i>	0.41
Bacteria	Proteobacteria	Betaproteobacteria	Burkholderiales	Alcaligenaceae	<i>Sutterella</i>	0.23

	Bacteria	Fusobacteria	Fusobacteriia	Fusobacteriales	Fusobacteriaceae	<i>Fusobacterium</i>	0.23
	Bacteria	Firmicutes	Clostridia	Clostridiales	Lachnospiraceae	<i>Roseburia</i>	0.18
	Bacteria	Firmicutes	Clostridia	Clostridiales	---	---	0.18
	Bacteria	Cyanobacteria	Chloroplast	Streptophyta	---	---	0.18
	Bacteria	Verrucomicrobia	Verrucomicrobiae	Verrucomicrobiales	Verrucomicrobiaceae	<i>Akkermansia</i>	0.09
	Bacteria	Firmicutes	Erysipelotrichi	Erysipelotrichales	Erysipelotrichaceae	<i>Holdemania</i>	0.09
	Bacteria	Firmicutes	Clostridia	Clostridiales	Peptostreptococcaceae	---	0.09
	Bacteria	Bacteroidetes	Bacteroidia	Bacteroidales	Rikenellaceae	---	0.09
	Bacteria	Proteobacteria	Alphaproteobacteria	Rhizobiales	Brucellaceae	<i>Ochrobactrum</i>	0.05
	Bacteria	Actinobacteria	Coriobacteriia	Coriobacteriales	Coriobacteriaceae	---	0.05