

Env #	Env	DNA	Bacteria ID	Relative Abundance
1	Adult Human Gut	GCTACG (RC: CGTAGC)	<i>Ruminococcus</i>	70% (14)
1	Adult Human Gut	GGATCG (RC: CGATCC)	<i>Bifidobacterium</i>	15% (3)
1	Adult Human Gut	CTACGT (RC: ACGTAG)	<i>Corynebacterium</i>	15% (3)
2	Infant Human	GCTACG (RC: CGTAGC)	<i>Ruminococcus</i>	15% (3)
2	Infant Human	GGATCG (RC: CGATCC)	<i>Bifidobacterium</i>	85% (17)
3	Pre-term Infant Human	GGATCG (RC: CGATCC)	<i>Bifidobacterium</i>	5% (1)
3	Pre-term Infant Human	CATAGC (RC: GCTATG)	Enterobacteriaceae	80% (16)
3	Pre-term Infant Human	CCGTCA (RC: TGACGG)	Enterococcaceae	15% (3)
4	Soil	TACTCA (RC: TGAGTA)	Streptomycetaceae	5% (1)
4	Soil	TAGCTA (RC: TAGCTA)	Solibacteraceae	25% (5)
4	Soil	GATTCA (RC: TGAATC)	Syntrophobacteraceae	40% (8)
4	Soil	CAATCG (RC: CGATTG)	Flexibacteraceae	25%(5)
4	Soil	AGACTG (RC: CAGTCT)	Bacillaceae	5% (1)