

Env #	Env	DNA	Bacteria ID	Relative Abundance
1	Infant Human	GCTACG (RC: CGTAGC)	<i>Ruminococcus</i>	15% (3)
1	Infant Human	GGATCG (RC: CGATCC)	<i>Bifidobacterium</i>	85% (17)
2	Pre-term Infant Human	GGATCG (RC: CGATCC)	<i>Bifidobacterium</i>	5% (1)
2	Pre-term Infant Human	CATAGC (RC: GCTATG)	Enterobacteriaceae	80% (16)
2	Pre-term Infant Human	CCGTCA (RC: TGACGG)	Enterococcaceae	15% (3)
3	Soil	TACTCA (RC: TGAGTA)	Streptomycetaceae	5% (1)
3	Soil	TAGCTA (RC: TAGCTA)	Solibacteraceae	25% (5)
3	Soil	GATTCA (RC: TGAATC)	Syntrophobacteraceae	40% (8)
3	Soil	CAATCG (RC: CGATTG)	Flexibacteraceae	25%(5)
3	Soil	AGACTG (RC: CAGTCT)	Bacillaceae	5% (1)
4	Plant Roots	TACTCA (RC: TGAGTA)	Streptomycetaceae	60% (12)
4	Plant Roots	GATTCA (RC: TGAATC)	Syntrophobacteraceae	25% (5)
4	Plant Roots	CAATCG (RC: CGATTG)	Flexibacteraceae	15% (3)