

Table 10. New taxonomic units identified from resampling of four subjects

Category	Description	No. of clones	No. of subjects
Phylum	<i>TM7</i>	6	2
	<i>Fusobacteria</i>	1	1
	Total	7	
Genus	<i>Flavobacteriaceae DQ337018[#]</i>	10	3
	<i>TM7</i>	6	2
	<i>Wautersia</i>	2	2
	<i>Chryseobacterium</i>	7	1
	<i>Flexibacteraceae[#]</i>	7	1
	<i>Sphingomonas</i>	4	1
	<i>Cyanobacteria[#]</i>	3	1
	<i>Abiotrophia</i>	2	1
	<i>Clostridiales AF287775[#]</i>	2	1
	<i>Flavobacteriaceae[#]</i>	2	1
	<i>Flexibacteraceae[#]</i>	2	1
	<i>Rhizobium</i>	2	1
	<i>Arsenicicoccus</i>	1	1
	<i>Comamonas</i>	1	1
	<i>Devosia</i>	1	1
	<i>Enterobacteriaceae[#]</i>	1	1
	<i>Flexibacteraceae^a</i>	1	1
	<i>Fusobacterium</i>	1	1
	<i>Incertae sedis 5 DQ297980[#]</i>	1	1
	<i>Janthinobacterium</i>	1	1
	<i>Lachnospira</i>	1	1
	<i>Lysobacter</i>	1	1
	<i>Microbacterium</i>	1	1
	<i>Neisseriaceae[#]</i>	1	1
	<i>Novosphingobium</i>	1	1
	<i>Ralstonia</i>	1	1
	<i>Shigella</i>	1	1
	<i>Xanthomonadaceae AB008508[#]</i>	1	1
	Total	63	
SLOTU	<i>Flavobacteriaceae DQ337018</i>	10	3
	<i>Streptococcus AF429766</i>	8	3

Category	Description	No. of clones	No. of subjects
	<i>Wautersia metallidurans</i>	2	2
	<i>Acinetobacter calcoaceticus</i>	12	1
	<i>Tsukamurella pulmonis</i>	10	1
	<i>Flexibacteraceae</i> AM000023 (85%)*	7	1
	<i>Chryseobacterium</i> DQ314741	5	1
	<i>Acinetobacter</i> DQ337025	4	1
	<i>Alkanindiges</i> AY251390 (94%)*	3	1
	<i>Cyanobacteria</i> AY853673 (83%)*	3	1
	<i>Abiotrophia defectiva</i>	2	1
	<i>Anaerococcus</i> Y07841 (95%)*	2	1
	<i>Anaerococcus</i> AY958826	2	1
	<i>Aquabacterium</i> AY212659	2	1
	<i>Chryseobacterium</i> AJ619057 (96%)*	2	1
	<i>Clostridiales</i> AF287775	2	1
	<i>Corynebacterium thomssenii</i>	2	1
	<i>Deinococcus</i> AY743261 (93%)*	2	1
	<i>Deinococcus grandis</i>	2	1
	<i>Deinococcus radiodurans</i>	2	1
	<i>Flavobacteriaceae</i> DQ256356 (94%)*	2	1
	<i>Flexibacteraceae</i> AM000023 (89%)*	2	1
	<i>Granulicatella adiacens</i>	2	1
	<i>Haemophilus</i> AY005033	2	1
	<i>Rhizobium giardinii</i>	2	1
	<i>Sphingomonas oligophenolica</i>	2	1
	<i>Sphingomonas taejonensis</i>	2	1
	<i>Staphylococcus lugdunensis</i>	2	1
	<i>Staphylococcus pasteurii</i>	2	1
	TM7 DQ248299 (95%)	2	1
	TM7 phylum sp. oral clone DR034	2	1
	<i>Acinetobacter parvus</i>	1	1
	<i>Actinomyces odontolyticus</i>	1	1
	<i>Anaerococcus</i> AY542229	1	1
	<i>Arsenicococcus bolidensis</i>	1	1
	<i>Bdellovibrio</i> DQ413094 (90%)*	1	1
	<i>Comamonas denitrificans</i>	1	1
	<i>Devosia</i> AJ863377	1	1
	<i>Enterobacter aerogenes</i>	1	1
	<i>Flexibacteraceae</i> AM000023 (91%)*	1	1

Category	Description	No. of clones	No. of subjects
	<i>Fusobacterium</i> AY278617	1	1
	<i>Incertae sedis</i> DQ297980	1	1
	<i>Janthinobacterium lividum</i>	1	1
	<i>Lachnospira</i> AY977894	1	1
	<i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i>	1	1
	<i>Lysobacter</i> AY074793 (94%)*	1	1
	<i>Methylobacterium</i> Z23159	1	1
	<i>Microbacterium lacticum</i>	1	1
	<i>Neisseria</i> AJ239301	1	1
	<i>Neisseriaceae</i> AY225604 (91%)*	1	1
	<i>Novosphingobium subterraneum</i>	1	1
	<i>Paracoccus aminovorans</i>	1	1
	<i>Propionibacterium</i> AB042290	1	1
	<i>Pseudomonas costantinii</i>	1	1
	<i>Pseudomonas monteilii</i>	1	1
	<i>Ralstonia insidiosa</i>	1	1
	<i>Shigella dysenteriae</i>	1	1
	<i>Sphingomonas</i> AJ555475	1	1
	<i>Sphingomonas yabuuchiae</i>	1	1
	<i>Streptococcus australis</i>	1	1
	<i>Streptococcus</i> AY959092	1	1
	<i>Streptococcus pseudopneumoniae</i>	1	1
	TM7 AY349415	1	1
	TM7 AB195911 (94%)*	1	1
	<i>Xanthomonadaceae</i> AB008508	1	1
	Total	140	

#This sequence could not be classified at the genus level.

*Number in parentheses represents similarity to closest sequence in GenBank.