

**Table 6. Genera detected in 1,221 16S rDNA clones from normal human skin**

Genus <sup>#</sup>	SubA (n=208)	SubB (n=204)	SubC (n=202)	SubD (n=204)	SubE (n=203)	SubF (n=200)	Total (n=1221)
<i>Acetobacteraceae</i> *	2						2
<i>Acidovorax</i>		2			2	1	5
<b><i>Acinetobacter</i></b>	3	14	14	10	1	3	45
<i>Actinomyces</i>	1		4		3	4	12
<i>Actinomycetales</i> *	2						2
<i>Actinomycetales</i> AY770698 *	4		2		7	6	19
<i>Alkanindiges</i>		5					5
<i>Amaricoccus</i>	1						1
<i>Anaerococcus</i>	2	2		2	1	4	11
<i>Aquabacterium</i>			2				2
<i>Atopobium</i>	2			1			3
<i>Bacteria</i> AJ619064 *	8						8
<i>Bacteria</i> *	1						1
<i>Bacteroidales</i> AF385513 *					2		2
<i>Bdellovibrio</i>			4				4
<i>Betaproteobacteria</i> AY360547 *		3					3
<i>Bradyrhizobiaceae</i> U87763 *		2					2
<i>Brevibacterium</i>				3			3
<i>Brevundimonas</i>	3						3
<i>Burkholderiales</i> DQ016727 *	1	3					4
<i>Carnobacterium</i>		10					10
<i>Caulobacteraceae</i> AJ459874 *					1		1
<i>Chitinophaga</i>	1						1
<i>Comamonadaceae</i> *	4						4
<b><i>Corynebacterium</i></b>	3	15	53	91	43	27	232
<i>Cyanobacteria</i> *			2				2
<i>Deinococcus</i>	35						35
<i>Dermacoccus</i>	9		1				10
<i>Dialister</i>				2			2
<i>Diaphorobacter</i>	1	2	2		2		7
<i>Enhydrobacter</i>	9	10	15				34
<i>Enterobacter</i>			1				1
<i>Enterococcus</i>			2				2
<i>Eremococcus</i>				1			1
<i>Facklamia</i>				5	2		7

Genus <sup>#</sup>	SubA (n=208)	SubB (n=204)	SubC (n=202)	SubD (n=204)	SubE (n=203)	SubF (n=200)	Total (n=1221)
<b><i>Finegoldia</i></b>	2	6	1	2	3	2	16
<i>Flavobacteriaceae</i> AF502204 *	1						1
<i>Flexibacteraceae</i> *	1						1
<i>Gammaproteobacteria</i> AY922146 *						1	1
<i>Gardnerella</i>			3				3
<i>Gemella</i>				1	2		3
<i>Gordonia</i>	11						11
<i>Granulicatella</i>		3					3
<i>Haemophilus</i>	1	2	2			2	7
<i>Hymenobacter</i>					2		2
<i>Hyphomicrobium</i>			1				1
<i>Roseateles</i>		10	1		1		12
<i>Janibacter</i>	1						1
<i>Klebsiella</i>		4					4
<i>Kocuria</i>	2	3	6				11
<i>Lactobacillus</i>		9	3		2		14
<i>Leuconostoc</i>			1				1
<i>Methylobacterium</i>	4						4
<i>Methylophilus</i>		7					7
<i>Micrococcus</i>		5				1	6
<i>Microcunatus</i>				1			1
<i>Mobiluncus</i>	1						1
<i>Mycobacterium</i>	3						3
<i>Nakamurella</i>	2						2
<i>Neisseria</i>	1						1
<i>Neisseriaceae</i> AY225604 *		17			1		18
<i>Nostocoida</i>	1						1
<i>Paracoccus</i>	1						1
<i>Pasteurellaceae</i> AJ290758 *					1		1
<i>Pasteurellaceae</i> AY005034 *					1		1
<i>Pedomicrobium</i>		4					4
<i>Peptoniphilus</i>			1	1	2		4
<i>Peptostreptococcus</i>				1			1
<i>Porphyromonas</i>				2			2
<i>Prevotella</i>	1			3	5	1	10
<b><i>Propionibacterium</i></b>	22	25	25	30	47	119	268
<i>Pseudomonadaceae</i> *		4					4
<i>Pseudomonas</i>		19	6		6	2	33
<i>Rhizobiales</i> *	1						1

<b>Genus<sup>#</sup></b>	<b>SubA (n=208)</b>	<b>SubB (n=204)</b>	<b>SubC (n=202)</b>	<b>SubD (n=204)</b>	<b>SubE (n=203)</b>	<b>SubF (n=200)</b>	<b>Total (n=1221)</b>
<i>Rhizobiales</i> *			2				2
<i>Rhizobiales AF358012</i> *			1				1
<i>Rhodococcus</i>	12						12
<i>Roseomonas</i>				1			1
<i>Rothia</i>	13		1	3	4	1	22
<i>Selenomonas</i>					4		4
<i>Serratia</i>					2		2
<i>Sphingopyxis</i>	3					1	4
<b><i>Staphylococcus</i></b>	7	6	34	38	32	19	136
<i>Stenotrophomonas</i>					1		1
<b><i>Streptococcus</i></b>	16	12	12	4	22	5	71
<i>Tetrasphaera</i>	1						1
<i>Tsakamurella</i>	3						3
<i>Thermomicrobia</i>	1						1
<i>Veillonella</i>	1			1	1	1	4
<i>Xanthomonadaceae</i> *	3						3
<i>Zimmermannella</i>				1			1

<sup>#</sup> Bold lettering indicates that the genus was found in all six subjects tested.

\*This sequence could not be classified at the genus level.