



PMC full text: [BMC Microbiol. 2009; 9: 102.](#)

Published online 2009 May 20. doi: [10.1186/1471-2180-9-102](#)

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## Table 1

Detailed composition of the neovaginal microflora of 30 swab samples, as determined by culture and tDNA-PCR based identification.

Cultured species	n	Cultured species	n
<b>Actinobacteria</b>		<b>Firmicutes</b>	
<i>Actinobaculum massiliense</i>	2	<i>Anaerococcus hydrogenalis</i>	1
<i>Actinobaculum schaalii</i>	1	<i>Anaerococcus tetradius</i>	1
<i>Actinomyces meyeri</i>	6	<i>Anaerococcus vaginalis</i>	3
<i>Actinomyces neuui</i>	2	<i>Bacillus</i> sp.	1
<i>Actinomyces radingae</i>	1	<i>Clostridium perfringens</i>	1
<i>Actinomyces</i> sp.	2	<i>Enterococcus faecalis</i>	13
<i>Actinomyces turicensis</i>	1	<i>Enterococcus</i> sp.	1
<i>Actinomyces urogenitalis</i>	2	<i>Facklamia hominis</i>	1
<i>Arcanobacterium bernardiae</i>	1	<i>Finegoldia magna</i>	7
<i>Arcanobacterium pyogenes</i> like	1	<i>Lactobacillus casei</i>	1
<i>Atopobium vaginae</i>	2	<i>Peptoniphilus indolicus</i>	6
<i>Bifidobacterium bifidum</i>	1	<i>Peptoniphilus lacrimalis</i>	6
<i>Bifidobacterium longum</i>	1	<i>Peptoniphilus</i> sp.	6
<i>Corynebacterium aurimucosum</i>	2	<i>Staphylococcus aureus</i>	3
<i>Corynebacterium glucuronolyticum</i>	1	<i>Staphylococcus capitis</i>	1
<i>Corynebacterium pseudogenitalium</i>	1	<i>Staphylococcus cohnii</i>	1
<i>Corynebacterium</i> sp.	13	<i>Staphylococcus epidermidis</i>	19
<i>Gardnerella vaginalis</i>	1	<i>Staphylococcus haemolyticus</i>	8
<i>Mobiluncus curtisii</i>	10	<i>Staphylococcus hominis</i>	3
<i>Olsenella uli</i>	1	<i>Staphylococcus lugdunensis</i>	3
<i>Slackia exigua</i>	2	<i>Staphylococcus pettenkoferi</i>	3
<i>Varibaculum cambriense</i>	7	<i>Staphylococcus simulans</i>	1
		<i>Staphylococcus</i> sp.	6
<b>Bacteroidetes</b>		<i>Staphylococcus warneri</i>	2
<i>Bacteroides coagulans</i>	8	<i>Streptococcus agalactiae</i>	4
<i>Bacteroides ureolyticus</i>	10	<i>Streptococcus anginosus</i> group	16
<i>Porphyromonas somerae</i>	6	<i>Streptococcus dysgalactiae</i>	1
<i>Prevotella bivia</i>	1	<i>Streptococcus oralis</i>	1

<b>Cultured species</b>	<b>n</b>	<b>Cultured species</b>	<b>n</b>
<i>Prevotella corporis</i>	4	<i>Streptococcus</i> sp.	4
<i>Prevotella disiens</i>	1		
<i>Prevotella</i> sp.	1	<b>Possible novel species and genera*</b>	
		TSWGenotypeA <i>Betaproteobacterium</i> [FM945400]	4
<b>Fusobacteria</b>		TSWGenotypeB <i>Porphyromonas</i> sp. [FM945401]	1
<i>Fusobacterium nucleatum</i>	1	TSWGenotypeC <i>Bacteroidetes</i> [FM945402]	3
<i>Fusobacterium</i> sp.	2	TSWGenotypeD <i>Clostridia</i> [FM945403]	5
		TSWGenotypeE <i>Clostridia</i> [FM945404]	2
<b>Proteobacteria</b>		TSWGenotypeF <i>Clostridia</i> [FM945405]	1
<i>Acinetobacter</i> sp.	1	TSWGenotypeG <i>Clostridia</i> [FM945406]	1
<i>Alcaligenes faecalis-like</i>	1	TSWGenotypeH <i>Bacilli</i> [FM945407]	2
<i>Escherichia coli</i>	7	TSWGenotypeI <i>Brevibacterium</i> sp. [FM945408]	2
<i>Klebsiella pneumoniae</i>	1		
<i>Proteus mirabilis</i>	1		

\* accession number between brackets