

Table S1. Identity of isolated bands from PCR-DGGE fingerprints of mixed communities as recovered from membrane filter of collected samples from the water bottling plant after sequencing of the variable V6-V8 region of 16S rRNA genes

Band <sup>1</sup>	Closest relative	Identity (%)	GenBank accession no. of closest relative
B1	<i>Citrobacter</i> spp.	99	KM187140
B2	<i>Staphylococcus</i> spp.	100	MG162677
B3	<i>Staphylococcus</i> spp.	99	MK942713
B4	<i>Pseudomonas</i> spp.	100	MF144455
B5	<i>Bacillus</i> spp.	97	MK660034
B6	<i>Exiguobacterium</i> spp.	100	KC160692

<sup>1</sup> Bands are indicated in Figure 1



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