

Table S2. Microbiological contamination of the leachates in the tested landfill

No.	Number of microorganisms, CFU ml ⁻¹								Sulfate reducing anaerobes
	Psychrophilic bacteria	Mesophilic bacteria	Fungi	Actinomycetes	Haemolytic Staphylococci	<i>Enterobacteriaceae</i>	Mannitol-positive Staphylococci	<i>Pseudomonas fluorescens</i>	
1	M:1.18x10 ⁶	M:7.07x10 ⁵	M:3.67x10 ¹	M:7.67x10 ¹	M:1.07x10 ⁴	M:7.00x10 ³	M:1.47x10 ⁴	M:1.33x10 ¹	M:1.70x10 ³
	SD:9.45x10 ⁴	SD:1.89x10 ⁵	SD:5.77x10 ⁰	SD:5.77x10 ⁰	SD:5.77x10 ²	SD:2.00x10 ³	SD:1.53x10 ³	SD:5.77x10 ⁰	SD:4.58x10 ²
2	M:1.74x10 ⁶	M:8.57x10 ⁵	M:6.67x10 ¹	M:8.67x10 ¹	M:1.27x10 ⁴	M:1.43x10 ⁴	M:1.20x10 ⁴	M:6.67x10 ⁰	M:3.83x10 ³
	SD:4.58x10 ⁴	SD:2.08x10 ⁴	v5.77x10 ⁰	SD:5.77x10 ⁰	SD:1.15x10 ³	SD:4.04x10 ³	SD:2.00x10 ³	SD:5.77x10 ⁰	SD:3.06x10 ²
3	M:2.69x10 ⁶	M:1.18x10 ⁶	M:3.67x10 ¹	M:1.80x10 ²	M:3.33x10 ³	M:7.33x10 ³	M:4.33x10 ³	M:3.00x10 ³	M:6.37x10 ³
	SD:3.00x10 ⁴	SD:2.75x10 ⁵	SD:5.77x10 ⁰	SD:3.00x10 ¹	SD:1.15x10 ³	SD:1.53x10 ³	SD:5.77x10 ²	SD:1.00x10 ³	SD:1.10x10 ³
4	M:5.00x10 ⁴	M:5.33x10 ⁴	M:4.67x10 ¹	M:2.10x10 ²	M:4.33x10 ³	M:7.33x10 ¹	M:7.33x10 ³	M:1.33x10 ¹	M:2.97x10 ³
	SD:1.00x10 ⁴	SD:3.51x10 ⁴	SD:5.77x10 ⁰	SD:3.00x10 ¹	SD:5.77x10 ²	SD:5.77x10 ⁰	SD:1.53x10 ³	SD:5.77x10 ⁰	SD:1.38x10 ³
5	M:3.33x10 ⁴	M:3.37x10 ³	M:8.00x10 ¹	M:8.67x10 ¹	M:6.67x10 ²	M:2.23x10 ²	M:1.67x10 ³	M:1.33x10 ¹	M:5.83x10 ¹
	SD:5.77x10 ³	SD:8.08x10 ²	SD:2.00x10 ¹	SD:3.06x10 ¹	SD:5.77x10 ²	SD:6.51x10 ¹	SD:5.77x10 ²	SD:5.77x10 ⁰	SD:1.42x10 ¹

M - mean; SD - standard deviation