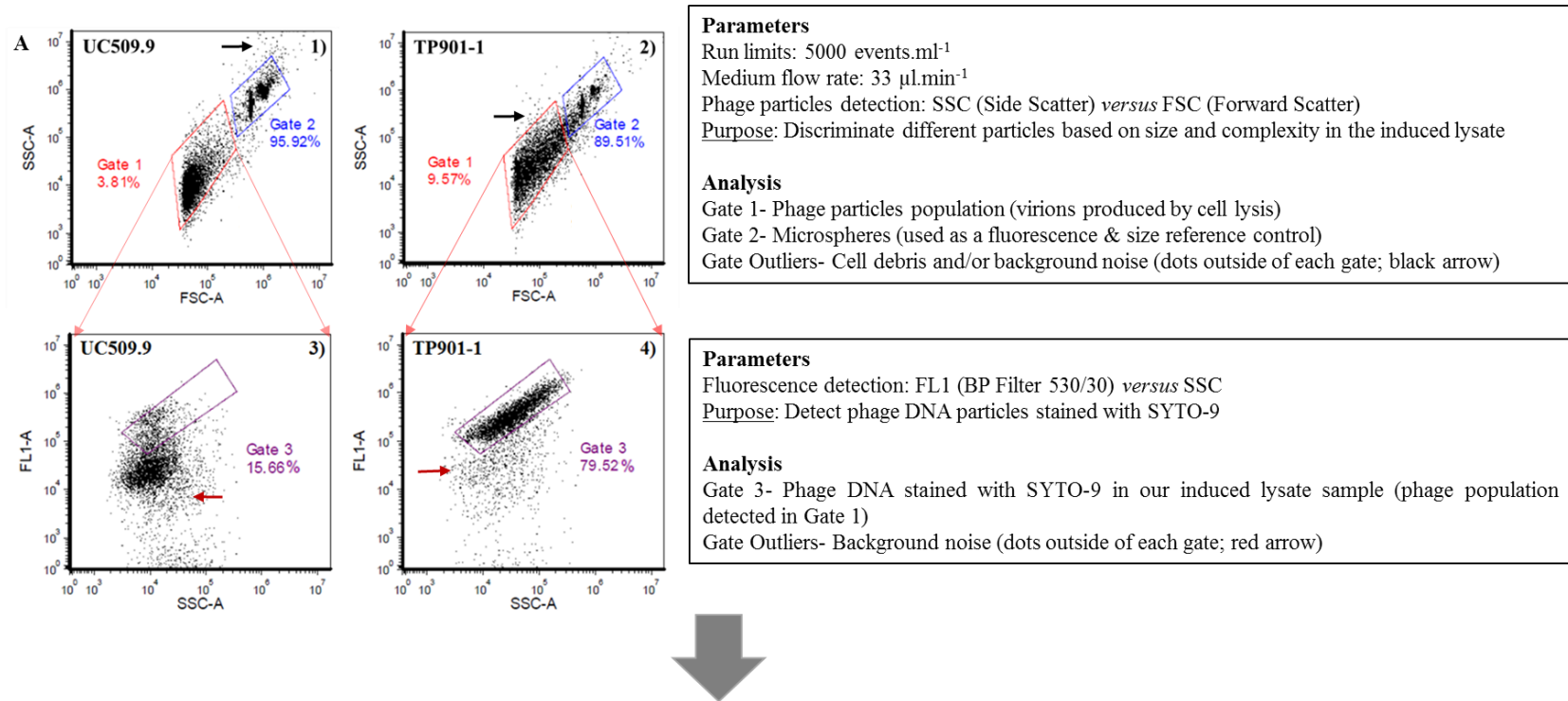


Title	Detecting Lactococcus lactis prophages by Mitomycin C-mediated induction coupled to flow cytometry analysis
Authors	Oliveira, Joana;Mahony, Jennifer;Hanemaaijer, Laurens;Kouwen, Thijs R.;Neve, Horst;MacSharry, John;van Sinderen, Douwe
Publication date	2017
Original Citation	Oliveira, J., Mahony, J., Hanemaaijer, L., Kouwen, T. R. H. M., Neve, H., MacSharry, J. and van Sinderen, D. (2017) 'Detecting Lactococcus lactis prophages by Mitomycin C-mediated induction coupled to flow cytometry analysis', <i>Frontiers in Microbiology</i> , 8, 1343 (11pp). doi: 10.3389/fmicb.2017.01343
Type of publication	Article (peer-reviewed)
Link to publisher's version	http://journal.frontiersin.org/article/10.3389/fmicb.2017.01343/full - 10.3389/fmicb.2017.01343
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		Flow cytometry analysis				
		% of events (SSC <i>versus</i> FSC)			% of fluorescence (FL1 <i>versus</i> SSC)	
<i>L. lactis</i> strains	Features	Sample population	Beads	Cell debris/ noise	Phage particles	Background noise
		Gate 1	Gate 2	Outliers of Gates 1 & 2	Gate 3	Outlier of Gate 3
UC509.9	No prophage released (negative control)	6.73 ± 2.07	92.93 ± 2.12	0.34 ± 0.05	16.33 ± 3.76	83.67 ± 3.76
NZ9000 (TP901-1 <i>erm</i>)	TP901-1 <i>erm</i> prophage (positive control)	10.98 ± 1.13	87.97 ± 1.26	1.05 ± 0.15	80.60 ± 2.39	19.39 ± 2.39

Figure S1. Schematic representation of the flow cytometry parameters, analysis and general results for the two *L. lactis* control strains. BD AccuriTM C6 flow cytometer was used for the implementation of the correct parameters to detect and enumerate phage DNA particles stained with SYTO-9 dye. (A1 and A3): Cytograms of 3 μg.ml⁻¹ MmC-treated *L. lactis* UC509.9 (prophage-free lactococcal strain used as negative control); (A2 and A4): Cytogram of 3 μg.ml⁻¹ MmC-treated *L. lactis* NZ9000 TP901-1*erm* (lactococcal strain harbouring the TP901-1 prophage used as a positive control).