

Journal of Laboratory Automation

<http://jla.sagepub.com/>

Flow Cytometric Enumeration of Bacteria Using TO-PRO[®]-3 Iodide as a Single-Stain Viability Dye

Monique Kerstens, Gaëlle Boulet, Christian Tritsmans, Tessa Horemans, Mario Hellings, Peter Delputte, Louis Maes and Paul Cos

Journal of Laboratory Automation 2014 19: 555 originally published online 14 August 2014

DOI: 10.1177/2211068214546745

The online version of this article can be found at:

<http://jla.sagepub.com/content/19/6/555>

Published by:



<http://www.sagepublications.com>

On behalf of:



Come Transform Research™

Society for Laboratory Automation and Screening

Additional services and information for *Journal of Laboratory Automation* can be found at:

Email Alerts: <http://jla.sagepub.com/cgi/alerts>

Subscriptions: <http://jla.sagepub.com/subscriptions>

Reprints: <http://www.sagepub.com/journalsReprints.nav>

Permissions: <http://www.sagepub.com/journalsPermissions.nav>

>> Version of Record - Nov 20, 2014

OnlineFirst Version of Record - Aug 14, 2014

What is This?

Flow Cytometric Enumeration of Bacteria Using TO-PRO®-3 Iodide as a Single-Stain Viability Dye

Journal of Laboratory Automation
2014, Vol. 19(6) 555–561
© 2014 Society for Laboratory
Automation and Screening
DOI: 10.1177/2211068214546745
jla.sagepub.com



Monique Kerstens¹, Gaëlle Boulet¹, Christian Tritsmans¹, Tessa Horemans¹,
Mario Hellings², Peter Delputte¹, Louis Maes¹, and Paul Cos¹

Abstract

Quantification of bacteria using conventional viable plate counting (VPC) is labor-intensive and time-consuming. Flow cytometry (FCM) can be proposed as a faster alternative. This study aimed to develop a flow cytometric, single-stain approach using TO-PRO®-3 iodide (TP3) for the quantification of *Staphylococcus aureus*, *Escherichia coli*, and *Bacillus subtilis* cells. Live or dead bacterial suspensions were stained with TP3 and analyzed using a FACSCalibur flow cytometer. After optimization of staining parameters and instrument settings, an excellent separation of viable and dead cells was achieved for all species. The quantitative performance of the technique was assessed by analyzing serial dilutions of bacterial suspensions using FCM and VPC. A highly linear correlation ($r^2 > 0.99$) was observed between the colony forming units (CFU)/mL as determined by FCM and by VPC over a concentration range of about 10^4 to 10^8 CFU/mL. As such, FCM quantification of viable bacteria using TP3 can be considered as an accurate and reliable alternative for VPC. The monostain procedure is easy to apply and cost-effective, and it allows bacterial enumeration in a broad variety of samples.

Keywords

flow cytometry, quantification, *Bacillus subtilis*, *Staphylococcus aureus*, *Escherichia coli*

Introduction

The quantification of microorganisms is ubiquitous in microbiology and usually based on the viable plate count (VPC).¹ This time-consuming, labor-intensive, and subjective method only allows enumeration of organisms that readily grow on solid medium, however.^{2–4} Of the alternative techniques under investigation, flow cytometry (FCM) has proven to be fast, precise and accurate. Furthermore, it enables growth-independent quantification and multiparametric analyses.^{4–7}

In a previous article, we reported the use of an FCM single-stain approach to directly quantify viable *Candida albicans* cells.⁸ Viable cells were discriminated from dead cells through a difference in fluorescence intensity using TO-PRO®-3 iodide (TP3), a membrane-impermeable dye that shows a greatly enhanced fluorescence on binding to double-stranded DNA (dsDNA) in bacteria that are dead and thus permeable.

The present study aimed to expand this technique to the enumeration of viable bacteria. We evaluated the influence of the small bacterial size on our FCM approach and assessed whether prokaryotes show the same TP3 staining characteristics. The first part of this study focused on the separation of viable bacteria from the dead subpopulation

and background noise. Second, the quantitative potential of FCM for viable bacteria was assessed in comparison with VPC, with particular focus on the precision of both methods. Four species with different Gram staining were selected because they are mentioned in the European Pharmacopoeia (monographs 2.6.12 and 2.6.13), indicating their importance for the pharmaceutical industry: *Escherichia coli* and *Pseudomonas aeruginosa* as Gram-negative test species, and *Staphylococcus aureus* and *Bacillus subtilis* as Gram-positive species.⁹ Finally, to confirm the applicability of the developed FCM method, the growth of *E. coli* under different experimental conditions was monitored.

¹University of Antwerp, Antwerp, Belgium

²Janssen Research & Development, Division of Janssen Pharmaceutica N.V., Beerse, Belgium

Received May 7, 2014.

Corresponding Author:

Paul Cos, Laboratory of Microbiology, Parasitology and Hygiene (LMPh), S7, Faculty of Pharmaceutical, Biomedical and Veterinary Sciences, University of Antwerp, Universiteitsplein 1, B-2610 Antwerp, Belgium.
Email: paul.cos@uantwerpen.be

Material and Methods

Bacterial Strains and Growth Conditions

E. coli ATCC 8739, *P. aeruginosa* ATCC 15442, *S. aureus* ATCC 5374, and *B. subtilis* ATCC 6633 were maintained on Tryptic Soy Agar (TSA; Lab M, Lancashire, UK) at 37°C and were subcultured every 24 h. A fresh culture was prepared weekly from a cryostock.

Qualitative Analysis

After overnight incubation, a single colony was placed in 10 mL sodium chloride peptone (SCP; 1 g of peptone, 3.6 g KH₂PO₄, 7.2 g Na₂HPO₄·H₂O, and 4.3 g NaCl dissolved in 1 L of ultrapure water with a pH of 7.0 ± 0.2). This suspension was defined as a 100% viable culture. Half of the sample was heat-killed in a water bath at 72 °C for 30 min, resulting in a 100% dead culture. One hundred microliters of this culture was plated on TSA and incubated for 48 h to confirm successful heat killing. A mixture of 50% viable and 50% dead cells was obtained by adding 2 mL of the 100% viable culture to an equal volume of the 100% dead culture. The bacterial suspensions were kept on ice during the experiments.

A FACSCalibur flow cytometer (BD Biosciences, Franklin Lakes, NJ, USA) equipped with a red diode laser ($\lambda_{ex} = 635$ nm) and a band pass filter measuring red fluorescence (FL4; 653–669 nm) was used. The sample flow rate was set at "low" (12 ± 3 µl/min). CountBright absolute counting beads (CB; Molecular Probes®, Eugene, OR, USA) were added to each sample as an internal standard and were counted in the channel detecting green fluorescence [fluorescent channel 1 (FL1)].

Mixed samples were used to optimize FCM separation of the viable and the dead populations by varying final concentrations of TP3 (1 µM–25 nM; Molecular Probes), incubation time (1 min–1 h), and voltages of the FCM channels (forward scatter (FSC), side scatter (SSC), FL1, and FL4) with the parameters for *C. albicans* in mind, as previously described.⁸ The viable and dead cultures were analyzed separately to confirm the localization of the respective populations in the FCM dotplot and to set the gates that delineate these groups.

Bacteria were stained with TP3 ($\lambda_{ex} = 642$ nm – $\lambda_{em} = 661$ nm) by adding 5 µl of the dye to 470 µl sample and 25 µl CB [final TP3 concentration of 200 nM; 10 min incubation time in the dark at room temperature (RT)]. The voltages were set at E01 for FSC, 445 for SSC, 450 for FL1, and 700 for FL4.

Quantitative Analysis

An entire overnight bacterial culture was scraped from agar and suspended in SCP. From this concentrated sample, a 10-fold dilution series was made in SCP. To increase the

number of standards in the linear area of the correlation curve, three twofold and three fivefold dilutions were prepared to cover that concentration range. The dilutions were quantified in triplicate by both VPC and FCM. Sterile SCP was included as a blank. The experiment was repeated on three different days. Correlation between both methods was determined statistically using linear regression analysis (Graphpad prism 4).

Precision Analysis

To compare the precisions of VPC and FCM, one dilution approximating the middle of the linear range of the calibration curve was analyzed 10 times by both VPC and FCM for every bacterial species.⁹ The coefficients of variation (CV) of the viable cell counts obtained by both methods were compared.

Growth Curve

The developed FCM technique was applied to assess the stability of an *E. coli* culture in SCP at RT and on ice. The resulting growth curves were compared to a growth curve of *E. coli* in Tryptic Soy Broth (TSB; Lab M, Lancashire, UK). For this experiment, three 50 mL tubes containing 40 mL of medium were inoculated with 8 × 10⁵ colony forming units/mL (CFU/mL) *E. coli* cells from a fresh culture. Bacterial concentrations were determined by FCM every hour for the first 10 hours and at $t = 12, 24, 30$, and 48 h.

Results

Qualitative Analysis

To optimize the procedure for qualitative analysis, different TP3 concentrations (1 µM–25 nM) and incubation times (1 min–1 h) were explored. A good FCM separation of the viable- and dead-cell populations could be achieved for *B. subtilis* 6633, *S. aureus* 5374, and *E. coli* 8739 (Fig. 1). The best results were obtained using a concentration of 200 nM TP3 followed by 10 min incubation in the dark at RT. The FCM voltages were set at E01 for FSC, 445 for SSC, 450 for FL1, and 700 for FL4. An SSC threshold was set at 125 based on a blank sample to eliminate as much background noise as possible.

Using these settings, it was impossible to distinguish viable from dead *P. aeruginosa* cells (Fig. 2). Varying the concentration of TP3, incubation time or voltages did not improve the separation.

Quantitative Analysis

For *E. coli*, *B. subtilis*, and *S. aureus*, quantitative analyses were performed. A good linear correlation ($r^2 > 0.99$) was observed between 6 × 10³ and 6 × 10⁷ viable cells/mL (VC/mL) (Fig. 3; Table 1). The lower limit of this linear range

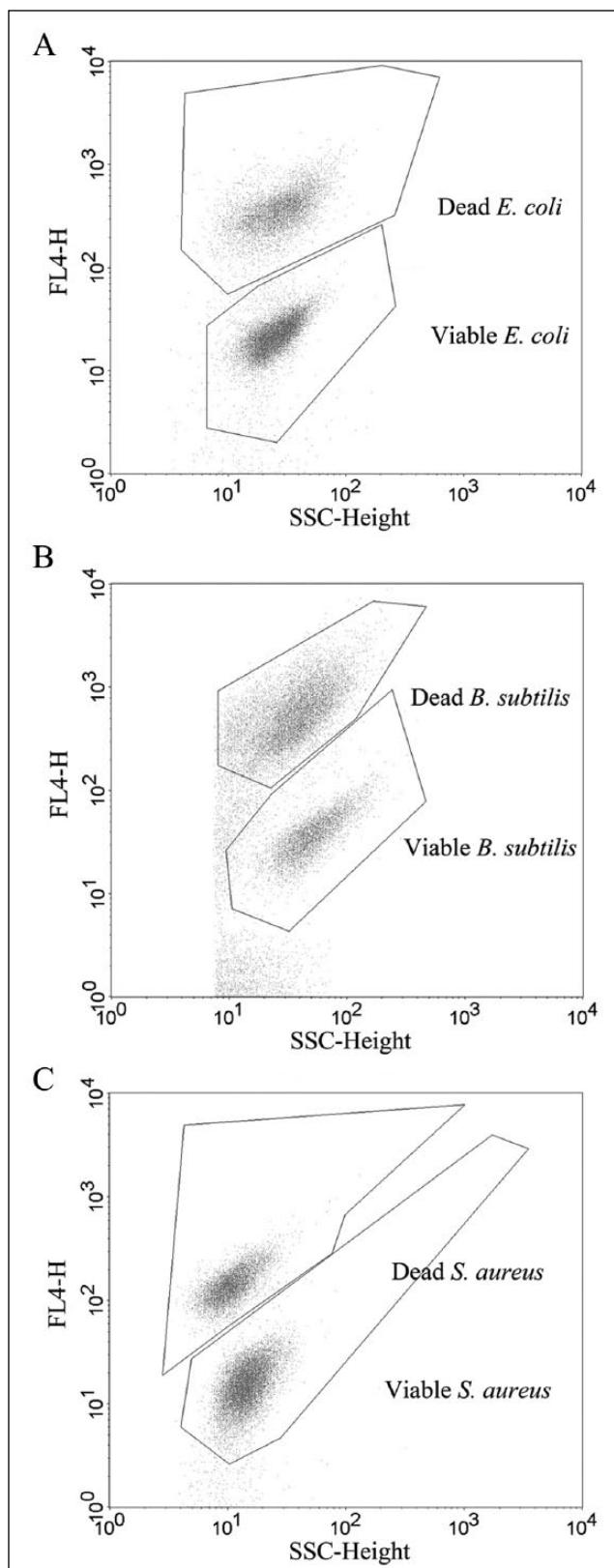


Figure 1. Flow cytometric separation of viable and dead bacterial cells in the side scatter (SSC)–fluorescent channel 4 (FL4) dotplot. (a) *E. coli*. (b) *B. subtilis*. (c) *S. aureus*.

The plots show only the events defined as bacterial cells in the forward scatter–SSC dotplot. Samples contained approximately 50% viable and 50% heat-killed cells. Dead cells show a brighter TO-PRO®-3 iodide fluorescence than viable cells. Hence, the former subpopulation is located higher on the FL4 axis. The respective subpopulations are defined by the polygonal regions. The *B. subtilis* dotplot shows fewer events than the other dotplots due to a lower concentration of microorganisms.

was determined by the limit of detection ($3 \times$ the standard deviation of the blank). The onset of the plateau phase marked the upper end of the range. The excellent correlation between FCM and VPC confirms the feasibility of FCM quantification. For *E. coli* and *S. aureus*, both methods yielded statistically equivalent results, because the 95% confidence interval of the slope included "1" and the interval of the intercept included "0."⁸ Although the slope did not significantly differ from 1, a minimal relative bias was found for *B. subtilis* (intercept > 0).

Precision Analysis

Every fifth sample of the dilution series, corresponding with the middle of the linear range, was analyzed 10 times by VPC and FCM. For each bacterial species, the CV was compared between both methods. FCM consistently yielded a lower CV than VPC, indicating a higher precision. The difference in CV was most pronounced for *E. coli* and *B. subtilis* (Fig. 4).

Growth Curve

Figure 5 shows the growth curves of *E. coli* in SCP (RT and on ice) and in TSB (RT). For SCP on ice, there was no significant increase or decrease in the amount of viable cells present during a period of 48 h. For both media at RT, a characteristic growth curve could be observed. A lag phase of approximately 2 h was followed by an exponential growth, which was the fastest for the TSB sample. A stationary phase was reached after 12 h and continued for at least 36 h. During the stationary phase, more viable cells were present in TSB than in SCP.

Discussion

Quantification of microorganisms is important in a broad range of industries and settings. Within the pharmaceutical and food industry, quantification of bacteria supports quality control of the production process and of the end product.⁴ For basic research purposes, inoculum concentrations need to be determined, whereas in applied research such as preclinical drug development, the amount of viable bacteria after treatment indicates drug efficacy.¹⁰ Clinically, bacterial enumeration can be used for diagnostic purposes and presents an essential element of quality assurance (e.g., to monitor the effectiveness of disinfection procedures).^{11–12}

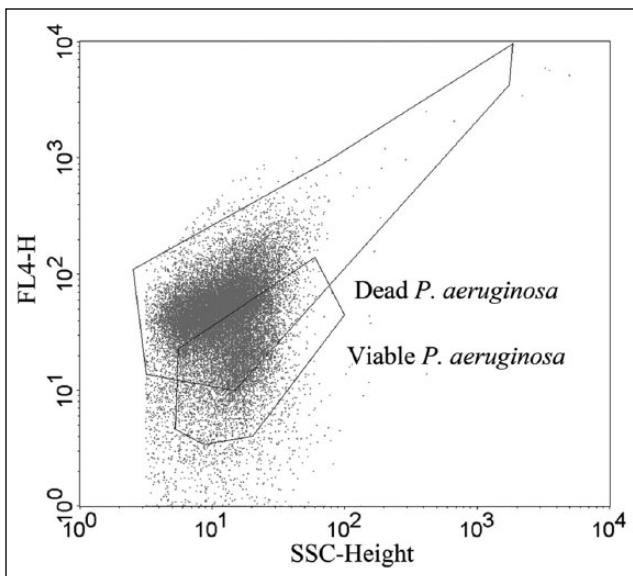


Figure 2. The side scatter (SSC)-fluorescence channel 4 (FL4) dotplot of a mixed *P. aeruginosa* sample. The dotplot clearly shows that the dead subpopulation partially overlaps with the viable subpopulation. The polygonal regions denote the positions of the viable and dead cells as determined by the analysis of a 100% viable-cell and a 100% dead-cell suspension.

Enumeration methods can be based on direct quantification, in which the organisms themselves are counted, or indirect quantification, in which the output parameter correlates with the amount of bacteria. Indirect techniques, such as quantitative PCR (qPCR), impedance, and turbidity measurements, require standard curves, but standards that accurately reflect exact quantities are not always available.¹ Although qPCR has proven to be a valuable method, sample preparation and processing can be relatively time-consuming. Moreover, the presence of inhibitory substances can greatly influence the reaction.^{5,13} For primer design, the organism needs to be identified, which is not always the case for environmental samples.⁷ In addition, depending on the experimental setting, qPCR analysis does not support accurate discrimination between viable and dead bacteria.⁵

Microscopy, VPC, and FCM are examples of direct quantification techniques. Despite its drawbacks, VPC is omnipresent. However, the method allows no real-time data acquisition and is labor-intensive, subjective, and growth-dependent. Considering the estimation that only 1% of all microorganisms is culturable on solid medium, most of them cannot be counted by VPC, nor does the technique allow the detection of viable but nonculturable cells (VBNCs), which are metabolically active but cease to divide under stress conditions, such as starvation and changes in oxygen concentration or temperature.¹⁴ VBNCs can produce metabolites that compromise production processes. Moreover, pathogens can also occur in this state without losing their pathogenic abilities, which clearly highlights the importance of VBNC detection.^{4,15}

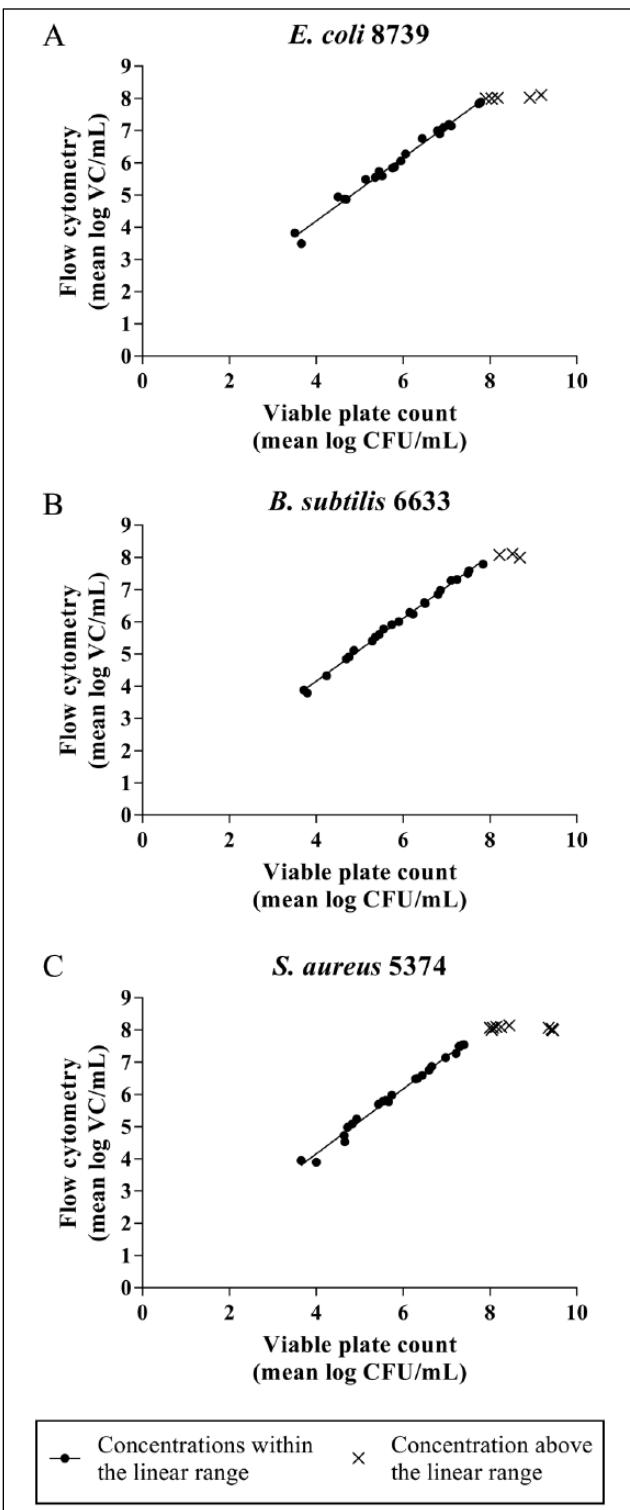


Figure 3. Comparison between the viable count/mL (VC/mL) as determined by flow cytometry and the amount of colony forming units/mL (CFU/mL) by viable plate counting for *E. coli*, *B. subtilis*, and *S. aureus*. (a) *E. coli*. (b) *B. subtilis*. (c) *S. aureus*. The results are displayed as the average of three replicates \pm SEM. Samples with a concentration lower than the limit of detection are not displayed.

Table 1. Outcome Parameters of the Linear Regression Analysis for the Three Bacteria.

Species	<i>E. coli</i>	<i>B. subtilis</i>	<i>S. aureus</i>
Slope*	0.9257–1.029	0.9494–1.001	0.9624–1.050
Intercept*	-0.0106–0.6027	0.0971–0.4074	-0.1348–0.3903
r^2	0.9881	0.9966	0.9909
Lower limit (VC/mL)	3.26×10^3	7.66×10^3	8.26×10^3
Upper limit (VC/mL)	7.65×10^7	6.19×10^7	3.51×10^7

Bacterial suspensions were analyzed in triplicate by viable plate counting (VPC) and by flow cytometry (FCM) on three different days. The logarithm of the average FCM count was plotted against the results of the VPC. This table gives an overview of the results of the linear regression analysis. * 95% confidence interval. VC/mL, viable cells/mL as determined by FCM.

Direct FCM counting poses an interesting alternative to VPC, because of its speed and its objective readout, combined with a high precision and the possibility of investigating different cellular parameters in just one analysis.⁴

This study evaluated FCM-based quantification of TP3-stained bacteria in comparison to VPC as a conventional gold-standard technique. Our previous work showed that TP3 allows the discrimination of viable and dead *C. albicans* cells.⁸ Dead cells showed strong fluorescence, because TP3 can pass the damaged membrane to bind dsDNA, which enhances its fluorescence intensity. Viable cells are stained as well but to a much lesser extent because TP3 binds to the outer cell wall structures. This article proves that the same approach can be applied to bacteria, which differ from yeasts in size, cell wall, and membrane composition.

The staining procedures for *E. coli*, *B. subtilis*, and *S. aureus* proved to be similar to the procedure for yeast cells, which improves the ease of use and enables analysis of mixed samples. Given this uniformity, it seems very likely that the same staining procedure can be applied to other bacterial species as well. Qualitative experiments on various other bacterial species, such as *Lactobacillus rhamnosus* GR-I, *Enterococcus faecalis* (clinical isolate), and *Burkholderia cenocepacia* LMG 16656, have confirmed this assumption (**Suppl. Fig. S1**). Instrument settings for bacterial cells differed from the optimal settings for *C. albicans*, due to the smaller size of the prokaryotes. *P. aeruginosa* was the only exception encountered: A significant overlap between dead and viable cells was found. The fluorescence emitted by TP3-stained viable cells was comparable to that of the other viable bacteria tested, whereas the dead cells were less fluorescent than the other dead bacteria. Our hypothesis is that TP3 penetration into the dead *P. aeruginosa* is impaired due to an intrinsic lower permeability of the outer cell membrane of the cell wall.¹⁶ Compared with other Gram-negative bacteria, *P. aeruginosa* is characterized by an increased resistance to biocides. This phenomenon is attributed to the notably different lipopolysaccharide composition and cation content of the outer membrane, resulting in a stronger interaction between the lipopolysaccharide molecules. Consequently, membrane fluidity is reduced, leading to a decreased permeability.¹⁷

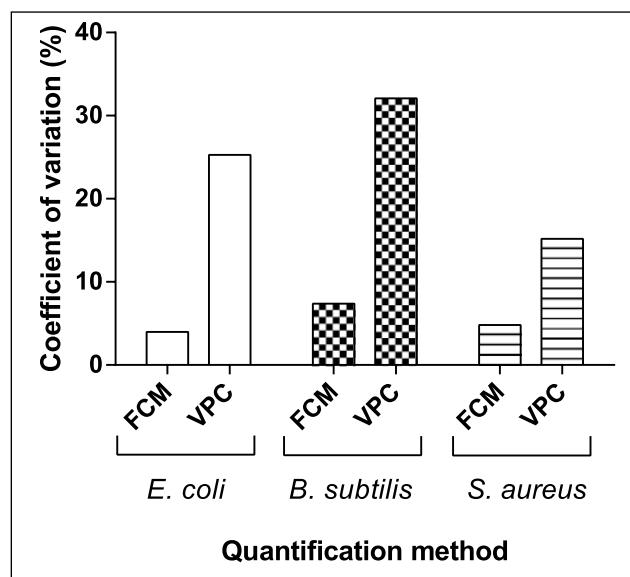


Figure 4. Coefficient of variation (CV) for flow cytometry (FCM) and viable plate count (VPC) quantification as determined from 10 repeated measurements of an identical sample of every bacterial species.

The CV (%) for the FCM quantification of the viable cells/mL is less than 10% for all the bacterial species (*E. coli*: 3.98%; *B. subtilis*: 7.39%; and *S. aureus*: 4.82%), whereas for VPC quantification, the obtained CVs were much higher (*E. coli*: 25.30%; *B. subtilis*: 32.08%; and *S. aureus*: 15.18%).

FCM and VPC quantification correlated very well in the linear range of 6×10^3 to 6×10^7 VC/mL. Below the linear range, background noise interferes with the detection of the bacteria, leading to a high CV and an overestimation of the FCM count. Above the linear range, the FCM detector suffers from oversaturation, but this problem could be overcome by diluting the samples. As to the precision of both methods, CVs were smaller for FCM than for VPC, especially for *E. coli* and *B. subtilis*. Because these species form large colonies, which tend to coalesce, precise VPC becomes more difficult.

In the European Pharmacopoeia, SCP is recommended as commercial buffer to stabilize the amount of microorganisms.⁹ Our FCM method was applied to investigate this assertion. *E. coli* was incubated in TSB and SCP at RT or on ice, and subjected to TP3 staining and FCM. Samples in

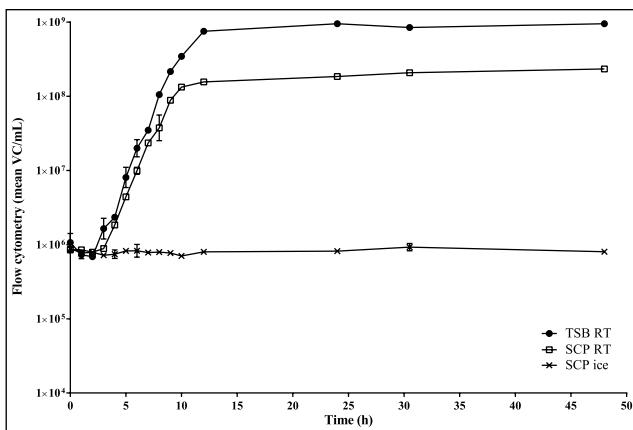


Figure 5. Growth curves of *E. coli* under different conditions: in sodium chloride peptone (SCP) on ice versus in tryptic soy broth (TSB), and in SCP at room temperature (RT). For 2 days, viable cell counts were performed in duplicate by flow cytometry using TO-PRO®-3 iodide. After 48 h, the following cell counts were obtained: 9.51×10^8 viable cells (VC)/mL (TSB RT), 2.34×10^8 VC/mL (SCP RT), and 8.01×10^5 VC/mL (SCP ice).

SPC on ice maintained a constant cell number up to 48 h of incubation. At RT, however, a classical growth pattern could be observed, clearly emphasizing the importance of low temperatures to stabilize cultures during long experiments. This evaluation proved that our FCM technique can be used to assess bacterial growth, although different culture conditions may demand minor experimental modifications. Because cells grown in TSB were somewhat larger than cells grown on solid medium, the gates were readjusted to obtain optimal results.

Because a FACSCalibur is not equipped with a volumetric sensor, CB were added to each sample as an internal standard. Adding the beads to a sample led to a slightly higher fluorescent signal of the viable population owing to the presence of 0.1% Tween 20 in the CB solution. This nonionic surfactant enhances TP3 binding to bacterial cells by increasing membrane fluidity and therefore influences the position of the viable population in the SSC–FL4 dotplot.¹⁸

It must be noted that despite the advantages of FCM, the technique also has its drawbacks compared with VPC. Although the latter is cost-effective and easy to perform, skilled technicians are needed to operate the flow cytometer and interpret the often complex data.^{19–20} Also, flow cytometers can be costly, although in recent years, prices have decreased and affordable bench-top flow cytometers have become commercially available.^{3,21–22}

Our data clearly indicate that FCM using TP3 as a single-stain viability dye allows accurate bacterial quantification. FCM has previously been described as a useful tool for the enumeration of viable bacteria, but most studies combined two or more dyes. The LIVE/DEAD® BacLight™ kit

is one of the most used stain combinations in this context. The kit, marketed by Invitrogen, contains the membrane-impermeable dye propidium iodide to mark dead cells and SYTO®9 to act as a counterstain.^{23–24} The addition of two dyes complicates FCM analysis due to spectral overlap and color compensation issues.^{25–27} Moreover, dual color viability staining saves fewer channels for the investigation of other cellular parameters. Other studies on single-stain FCM rarely focus on bacterial quantification or use indirect quantification approaches.^{28–30} Our TP3 assay enables direct counting of both viable- and dead-cell populations, making the addition of a counterstain redundant, whereas the unique spectral properties of TP3 offer an extra advantage. Because TP3 emission is located in the red end of the spectrum and only red laser light allows excitation, combination with other fluorochromes to assess additional cell parameters with minimal spectral overlap is feasible.³¹ In addition, the uniformity of the staining protocol eliminates the need for species-specific optimization, as is the case with many viability stains, including the LIVE/DEAD® BacLight™ kit.³

In conclusion, TP3 staining of bacteria for FCM quantification offers clear advantages compared with VPC and other FCM enumeration techniques, and it should be further explored in samples of pharmaceutical, environmental, or food origin.

Abbreviations

<i>B. subtilis</i>	= <i>Bacillus subtilis</i>
<i>C. albicans</i>	= <i>Candida albicans</i>
CB	= counting beads
CFU/mL	= colony forming units/mL
CV	= coefficient of variation
dsDNA	= double-stranded DNA
<i>E. coli</i>	= <i>Escherichia coli</i>
FCM	= flow cytometry
FL1	= fluorescence channel 1
FL4	= fluorescence channel 4
FSC	= forward scatter
<i>P. aeruginosa</i>	= <i>Pseudomonas aeruginosa</i>
qPCR	= quantitative PCR
RT	= room temperature
<i>S. aureus</i>	= <i>Staphylococcus aureus</i>
SCP	= sodium chloride peptone
SSC	= side scatter
TP3	= TO-PRO®-3 iodide
TSA	= tryptic soy agar
TSB	= tryptic soy broth
VBNC	= viable but nonculturable
VC/mL	= viable cells/mL
VPC	= viable plate count

Declaration of Conflicting Interests

The authors declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Funding

The authors declared receipt of the following financial support for the research, authorship, and/or publication of this article: This work was supported by the Research Foundation—Flanders (FWO grant 24761) and by the University of Antwerp (BOF/GOA grant 25624). Monique Kerstens is a PhD fellow of the FWO—Flanders.

References

- Jasson, V.; Jacxsens, L.; Luning, P.; et al. Alternative Microbial Methods: An Overview and Selection Criteria. *Food Microbiol.* **2010**, *27* (6), 710–730.
- Corry, J. E.; Jarvis, B.; Passmore, S.; et al. A Critical Review of Measurement Uncertainty in the Enumeration of Food Micro-Organisms. *Food Microbiol.* **2007**, *24* (3), 230–253.
- Davey, H. M. Life, Death, and In-Between: Meanings and Methods in Microbiology. *Appl. Environ. Microbiol.* **2011**, *77* (16), 5571–5576.
- Díaz, M.; Herrero, M.; García, L. A.; et al. Application of Flow Cytometry to Industrial Microbial Bioprocesses. *Biochem. Eng. J.* **2010**, *48* (3), 385–407.
- Khan, M. M. T.; Pyle, B. H.; Camper, A. K. Specific and Rapid Enumeration of Viable but Nonculturable and Viable-Culturable Gram-Negative Bacteria by Using Flow Cytometry. *Appl. Environ. Microbiol.* **2010**, *76* (15), 5088–5096.
- Morgan, C. A.; Bigeni, P.; Herman, N.; et al. Production of Precise Microbiology Standards Using Flow Cytometry and Freeze Drying. *Cytometry* **2004**, *62* (2), 162–168.
- Muller, S.; Nebe-von-Caron, G. Functional Single-Cell Analyses: Flow Cytometry and Cell Sorting of Microbial Populations and Communities. *FEMS Microbiol. Rev.* **2010**, *34* (4), 554–587.
- Kerstens, M.; Boulet, G.; Pintelon, I.; et al. Quantification of *Candida albicans* by Flow Cytometry Using TO-PRO®-3 Iodide as a Single-Stain Viability Dye. *J. Microbiol. Methods* **2013**, *92* (2), 189–191.
- European Directorate for the Quality of Medicines and HealthCare. *European Pharmacopoeia*; Council of Europe: Strasbourg, 2014; 8th ed.
- Broeren, M. A.; Maas, Y.; Retera, E.; et al. Antimicrobial Susceptibility Testing in 90 Min by Bacterial Cell Count Monitoring. *Clin. Microbiol. Infect.* **2013**, *19* (3), 286–291.
- Galvin, S.; Dolan, A.; Cahill, O.; et al. Microbial Monitoring of the Hospital Environment: Why and How? *J. Hosp. Infect.* **2012**, *82* (3), 143–151.
- Kadkhoda, K.; Manickam, K.; Degagne, P.; et al. UF-1000i Flow Cytometry Is an Effective Screening Method for Urine Specimens. *Diagn. Microbiol. Infect. Dis.* **2011**, *69* (2), 130–136.
- Girones, R.; Ferrus, M. A.; Alonso, J. L.; et al. Molecular Detection of Pathogens in Water—the Pros and Cons of Molecular Techniques. *Water Res.* **2010**, *44* (15), 4325–4339.
- Vartoukian, S. R.; Palmer, R. M.; Wade, W. G. Strategies for Culture of “Unculturable” Bacteria. *FEMS Microbiol. Lett.* **2010**, *309* (1), 1–7.
- Oliver, J. D. The Viable but Nonculturable State in Bacteria. *J. Microbiol.* **2005**, *43* (Spec. No.), 93–100.
- Lambert, P. A. Mechanisms of Antibiotic Resistance in *Pseudomonas aeruginosa*. *J. R. Soc. Med.* **2002**, *95* (Suppl. 41), 22–26.
- McDonnell, G. E. *Antisepsis, Disinfection, and Sterilization: Types, Action, and Resistance*; ASM: Washington, DC, 2007.
- Baumann, K.; Adelantado, N.; Lang, C.; et al. Protein Trafficking, Ergosterol Biosynthesis and Membrane Physics Impact Recombinant Protein Secretion in *Pichia pastoris*. *Microb. Cell Fact.* **2011**, *10*, 93.
- Gracias, K. S.; McKillip, J. L. A Review of Conventional Detection and Enumeration Methods for Pathogenic Bacteria in Food. *Can. J. Microbiol.* **2004**, *50* (11), 883–890.
- Reynolds, D. T.; Slade, R. B.; Sykes, N. J.; et al. Detection of *Cryptosporidium* Oocysts in Water: Techniques for Generating Precise Recovery Data. *J. Appl. Microbiol.* **1999**, *87* (6), 804–813.
- O’Donnell, E. A.; Ernst, D. N.; Hingorani, R. Multiparameter Flow Cytometry: Advances in High Resolution Analysis. *Immune Netw.* **2013**, *13* (2), 43–54.
- Mariani, M.; Colombo, F.; Assennato, S. M.; et al. Evaluation of an Easy and Affordable Flow Cytometer for Volumetric Haematopoietic Stem Cell Counting. *Blood Transfus.* **2014**, 1–5.
- Netuschil, L.; Auschill, T. M.; Sculean, A.; et al. Confusion over Live/Dead Stainings for the Detection of Vital Microorganisms in Oral Biofilms—Which Stain Is Suitable? *BMC Oral Health* **2014**, *14*, 2.
- Berney, M.; Hammes, F.; Bosshard, F.; et al. Assessment and Interpretation of Bacterial Viability by Using the LIVE/DEAD BacLight Kit in Combination with Flow Cytometry. *Appl. Environ. Microbiol.* **2007**, *73* (10), 3283–3290.
- Foladori, P.; Bruni, L.; Tamburini, S.; et al. Direct Quantification of Bacterial Biomass in Influent, Effluent and Activated Sludge of Wastewater Treatment Plants by Using Flow Cytometry. *Water Res.* **2010**, *44* (13), 3807–3818.
- Kramer, M.; Obermajer, N.; Bogovic Matijasic, B.; et al. Quantification of Live and Dead Probiotic Bacteria in Lyophilised Product by Real-Time PCR and by Flow Cytometry. *Appl. Microbiol. Biotechnol.* **2009**, *84* (6), 1137–1147.
- Tawakoli, P. N.; Al-Ahmad, A.; Hoth-Hannig, W.; et al. Comparison of Different Live/Dead Stainings for Detection and Quantification of Adherent Microorganisms in the Initial Oral Biofilm. *Clin. Oral Investig.* **2013**, *17* (3), 841–850.
- Budde, B. B.; Rasch, M. A Comparative Study on the Use of Flow Cytometry and Colony Forming Units for Assessment of the Antibacterial Effect of Bacteriocins. *Int. J. Food Microbiol.* **2001**, *63* (1–2), 65–72.
- Joung, Y. H.; Kim, H. R.; Lee, M. K.; et al. Fluconazole Susceptibility Testing of *Candida* Species by Flow Cytometry. *J. Infect.* **2007**, *54* (5), 504–508.
- Lapinsky, S. E.; Glencross, D.; Car, N. G.; et al. Quantification and Assessment of Viability of *Pneumocystis carinii* Organisms by Flow Cytometry. *J. Clin. Microbiol.* **1991**, *29* (5), 911–915.
- Bink, K.; Walch, A.; Feuchtinger, A.; et al. TO-PRO-3 Is an Optimal Fluorescent Dye for Nuclear Counterstaining in Dual-Colour FISH on Paraffin Sections. *Histochem. Cell Biol.* **2001**, *115* (4), 293–299.