

Microbes' heterogeneousness – a focus issue on cytometric technologies in microbial single cell analytics ((Forum contribution))

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Microbes' heterogeneousness – a focus issue on cytometric technologies in microbial single cell analytics ((Forum contribution))

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microbial single cell analytics	1.1.1	Deleted: Commentary¶ ¶
➢ http://doi.wiley.com/10.1002/cyto.a.20724	$\frac{1}{2}$	Formatted: Left, Line spacing: single
	, ,	Deleted: about
Cytometry of microorganisms was introduced about 30		Formatted: Bullets and Number
years ago [1]. Since those early times single cell		Fornaccea: Ballets and Namber
techniques have found their way into many fields of		
application for microorganisms ranging from health to		
biotechnology. As we now understand, more than 98%	netry	
[2] of the microbes in our environment cannot be kept in	K	
culture technologies that can precisely characterize	and a	
them, yet these microbes are of imminent importance.	*	Deleted: . Y y
Flow and image cytometry are the key technologies	↓ [*]	
herein. High resolution Omics-approaches become	1990 - Carlor	
add-ons for understanding intra- and interrelationships		
as well as for analyzing performances in bacterial		
production cultures or in natural communities [3, 4].		Exemption Fonty 11 nt
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The February 2009 issue of Cytometry Part A focuses		Deleted: ¶
on the cytometry of microorganisms and provides a rich	4	Formatted: Font: Italic
source of the newest approaches to the stoichiometric	~	
analysis of millions of individual microbial cells. This		
focus issue on microbiology contains a selection of manuscripts based on presentations		
at the 4 th International Conference on the Analysis of Microbial Cells at the Single Cell		
Level which was organised on behalf of the Microbial Physiology Section of the		
European Federation of Biotechnology, and held in Bad Schandau, Germany in 2008		
(http://qbab.dbs.aber.ac.uk/sc2008).	- < 7 -	Field Code Changed
((quote))		Formatted: Hyperlink, Font:
Newest approaches to the stoichiometric analysis of millions of individual microbial cells		(Default) Arial, English (U.K.)
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The issue contains concepts on how pathogens causing illness and death in humans,	-<5	Deleted: ¶
plants and animals may be sensed quickly and reliably. To address these problems,		Deleted: February
sophisticated single cell based techniques such as microfluidic surface plasmon		
resonance (SPR) are being developed to detect pathogens even at very low		
concentrations [5]. These negative features, however, should be considered to be one		
of the lesser impacts that microorganisms have on the existence of human society as		
microorganisms also have many beneficial effects.		
For millennia microorganisms have been exploited in traditional biotechnologies such as		Deleted: ¶
brewing and bread making. A growing range of microorganisms are used in modern		
biotechnology industries for the synthesis of valuable materials such as		
pharmaceuticals, antibiotics, enzymes, metabolites and various other compounds. As		
was shown in the special issue, maintenance of viable and productive populations		
within the industrial environment is of paramount importance to the success and		
efficiency of these industries and these processes can be optimized by the use of flow		
cytometric analyses [6]. Understanding bacterial physiology, growth behaviour and		

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survival strategies will open new ways to use, exploit or overcome bacteria on the basis of their inherent characteristics.

Another matter that is often overlooked when we study microorganisms in the laboratory is the fact that many of them do not normally grow as planktonic single cells in nature. Structured communities known as biofilms are the most common state of many bacteria within the environment and this distinction has important consequences for the physiology and behaviour of the individual cells within these communities. Cells respond to local environments by placing stimuli which may arise both from, for example, nutritional local 'hot spots' but also from signalling from neighbouring cells. Biofilm structures offer protection against extrinsic stressors such as predators or toxic chemicals via the production of impenetrable extracellular matrices. Such multicellular biofilms are ubiquitous and play a well accepted role in stabilizing infection in human disease. As the special issue showed, scientists are on the way to resolve the interactions between individuals of *Pseudomonas* spp. in order to understand infection processes [7].



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The heterogeneity of cells within a population might be considered as an important factor in maintaining stability of a variety of systems. It is known from phylogenetic diversity analyses that an increase in species numbers leads to a concomitant increase in the stability of a system [8]. Additionally, heterogeneity of cells with regard to productivity within populations should be seen not only in terms of its impact on bioprocess yields, but also in terms of the revelation of information about cell response to a complex interaction of environmental stimuli. Currently our understanding of how and why an individual microbial cell reacts to its environment is often poorly understood. A recent approach that will improve our understanding of the mechanisms and triggers involved in heterogeneity of response is a single cell bioreactor [9]. Such devices will facilitate the acquisition of functional and structural information about individual cells. In addition it will permit analysis of the single cell fluxome — the quantification of metabolite turnover in the single cell, at least for larger microorganisms such as yeast cells. Of equal importance is the need to develop methods to administer the huge single cell data profiles in a form which can be understood and efficiently accessed by the user

community – whether they are academics, clinicians or industrial scientists – a topic which was therefore intensively discussed at the 4th Conference in Bad Schandau. For example, in modelling applications, such data must be accessed and used to elucidate the behaviour of a cell in a community or population to improve our understanding of populations leading to process optimization and permitting accurate predictions [10].

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Mathematical modelling helps to understand the behaviour of biochemical reaction networks, or the highly coordinated response of single cell populations to changing (micro)environmental conditions.

To gain a deeper understanding of these topics take a close look at the February Issue of Cytometry Part A which is exclusively dedicated to the world of the single microbial cell.

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Cytometry PARTA

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special issue Cytometry of Microbes

4th International Conference on Analysis of Microbial Cells at the Single Cell Level Germany May 22nd -25th, 2008



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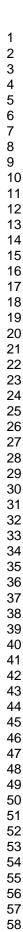
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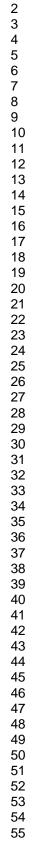
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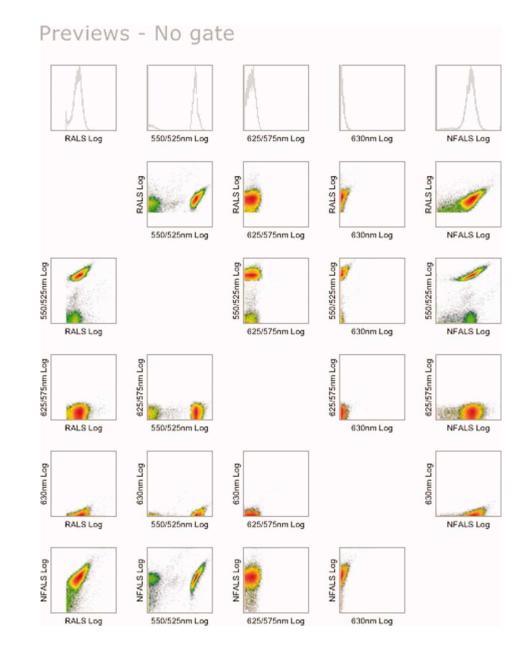
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