

# Modular biofactories at the cellular level

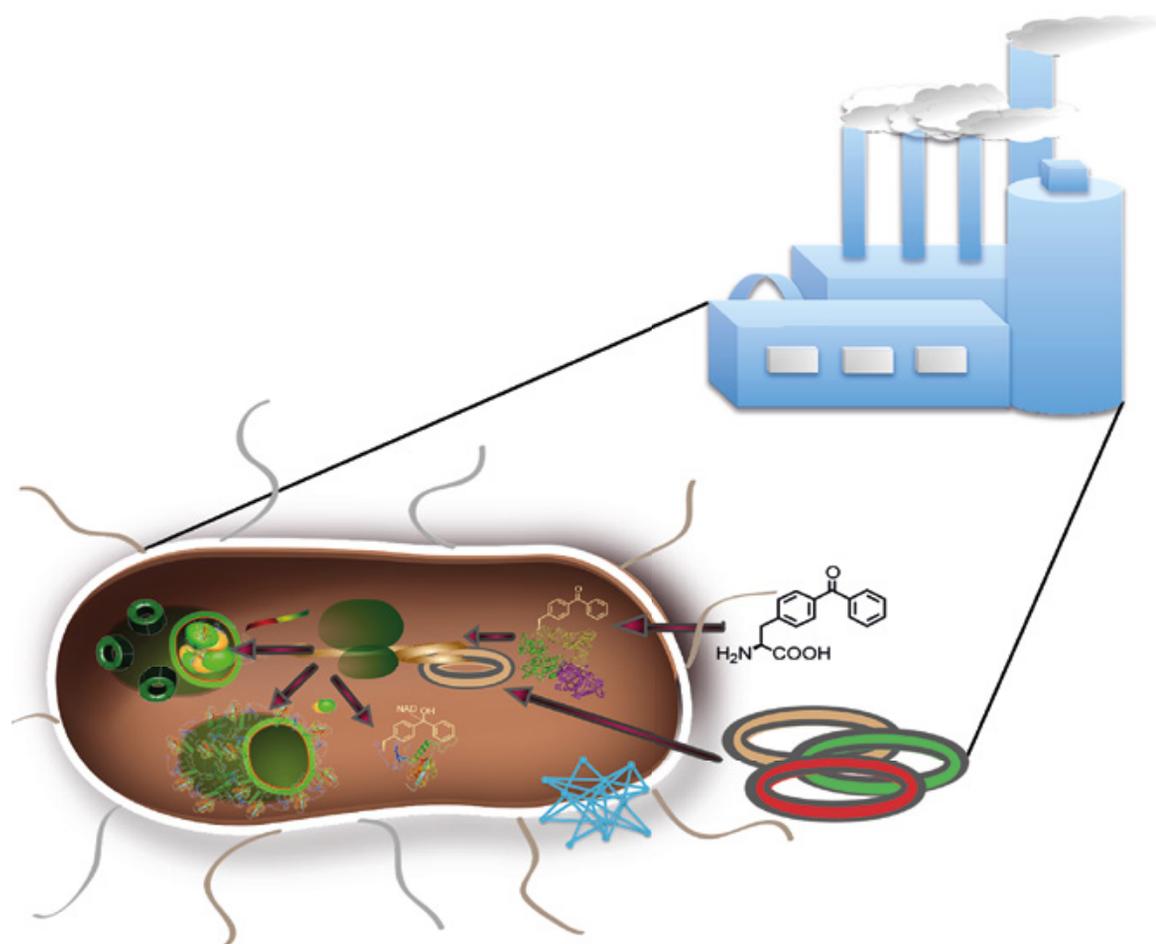
A new chapter in the book of life  
for a sustainable bioeconomy

Dr. Stefan Schiller  
Center for Biological Systems Analysis, University of Freiburg, Germany

Despite his love for complex molecular architectures, this „dyed-in-the-wool“ bio-organic chemist has never embraced the conventional segregation of synthetic polymers and biological macromolecules. All molecules are composed of atoms, after all. Why make an artificial distinction? Why not simply combine these domains together? Accordingly, the Schiller lab is developing new functional units or „modules“ in cells by a combination of approaches from molecular biology and from conventional synthetic chemistry.

This modularized extension of the cell with new, mutually-compatible elements such as de novo organelles, redesigned enzymes, transporters and switches, will permit future researchers to enlarge the functional spectrum of the cell. Not only to manufacture sustainable chemical basic raw materials that conserve resources but also to produce fuels, medicines and much more. These are indispensable prerequisites for the successful transformation of our economy into a sustainable and resilient bioeconomy.

Manipulating complex (cell-based) molecular systems might sound like a futuristic construction-kit concept but is solidly grounded in an understanding of molecules' chemical and physical properties alike, as well as their potential reaction and interactions – in short, the building blocks of matter. Indeed, work on complex systems fairly challenges us to do the same with cells. Alongside the interesting possibilities that result from the linking of conventional synthetic chemistry with cell biosynthesis – namely the synthesis of the most complex



**Fig. 1** The functional extension of cell functionality by means of new molecular modules is a key starting-point for a functional approach to studying cell functions and their application for the biosynthesis of molecules previously harvested by other procedures from non-renewable sources. To this end, new compartments/organelles are equipped with new synthesis pathways by methods including combination with enzyme complexes, molecular switches and the incorporation of unnatural/non-canonical amino acids.

and yet most highly-defined molecules that we know of – this symbiosis has another charm all of its own. The transference of new chemical synthesis products into the long-term cycle of natural material and energy flows is achieved by the module-like extension or substitution of cells, and grants access to raw materials (basic chemical substances), food products, energy sources and pharmaceuticals in a sustainable way that conserves resources. As we focus on developing biologics production systems to face new challenges, the performance limits of previous systems/cells must be made fit for purpose from scratch, i.e. at the molecular level. One significant aspect here is the extension of organisms' synthetic potential in terms of the biological accessibility to new compounds/ molecules/products; another is the integration of potentially disruptive factors into the system's performance profile and adaptive capabilities.

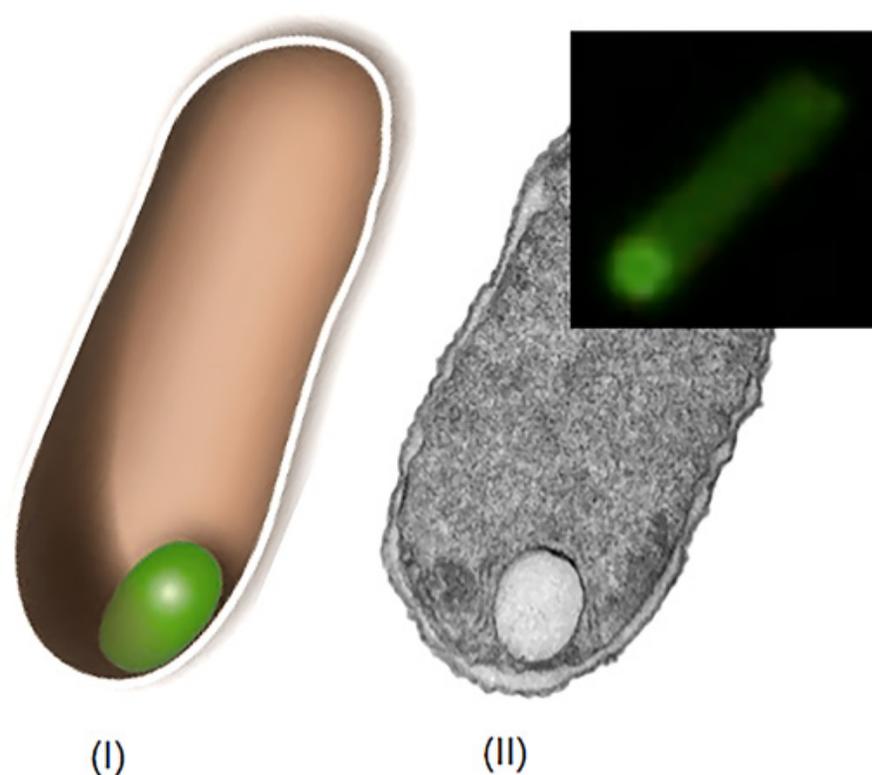
## Sustainability and resilience: from human ideas to living molecules

Achieving a modular understanding of the complex molecular system of the cell, introducing new modules, controlling interfaces, implementing new reactions in self-contained, self-forming reaction spaces within the cell: all are goals being pursued by the Stefan Schiller lab in the Center for Biological Systems Analysis at the Albert Ludwig University in Freiburg – and the applications are legion! This not only requires the development of mutually-compatible “modules” for the cells, such as new organelles, transporters, switches, enzymes, genes, etc. (fig. 1) – collectively termed “synthetic biology” – but also requires their investigation on a system-wide scale to determine their influence on the remaining cell components. This makes the integration of approaches from systems biology – metabolomics, proteomics and genomics – as important as the creation of mathematical models via methods such as metabolic flux analysis. The development of complex molecules and systems, coupled with the analysis and modeling of systemic influences, is a fundamental building block for the molecular science of the future. And not solely for bioproduction but also for a highly-developed understanding of the function of the body in a medicinal context.

One major advantage of such bio-based systems for the bio-economic model is the fact that they possess a series of bio-inherent properties, which are of fundamental importance for sustainable and resilient processes.

These include the natural dynamic properties of such molecular systems, their capacity for “self-healing” and adaptation, self-replication (and hence scalability), and, last but not least, their durability or resilience.

Over the long term, this will require a broad-based network of research teams and institutions working in close collaboration, capable of developing effective methods for the safe application of such microorganisms (biologic biosafety). Key techniques have already been established here, whereby cells remain alive only by targeted dosing with specific substances, for example, and are also unable to bypass this with simple mutations. Also known as “genetic containment strategies”, such approaches cannot be surmounted by spontaneous mutation or horizontal gene transfer. The conceptual basis is the redesign of essential enzymes with the aid of an extended genetic code, rendered non-viable in the environment by the specific incorporation of unnatural



**Fig. 2** Diagram of a bacterial cell (left) with a de novo-constructed compartment highlighted in green. The greyscale image to the right shows the electron micrograph of this new compartment. In the black enlarged detail from the fluorescence micrograph, we can see the cell and the modified compartment with the green fluorescent protein (GFP) as a round structure at the bottom left.



Researchers in the Schiller lab working on „synthetic“ cell modules are: Dr. Andreas Schreiber (top row right) and Dr. Matthias C. Huber (center row right); bottom row, left to right: Mildred Kramer, Biljana Maksimovic, Dr. Chunyan Yao, Stefan Schiller (standing); center row, left to right: Andreas Grabow, Dr. Wiltrud Wild, Nehrukumar Mathaiyan, Lisa Boos, Dr. Matthias C. Huber; top row, left to right: Cordula Hege, Dr. Andreas Schreiber [www.biotechonic.de](http://www.biotechonic.de)

**Stefan M. Schiller** studied chemistry at Gießen (Mainz, Germany) and the University of Massachusetts, majoring in macromolecular chemistry and biochemistry. For his doctorate in biomimetic membrane systems he worked till 2003 at the Max Planck Institute for Polymer Research in Mainz. Research positions in Israel and the USA (Stanford & IBM Research Center Almaden, San Jose) then followed. During a postdoc position at the Scripps Research Institute, La Jolla (USA), he completed research in the field of chemical and synthetic biology. He has been a principal investigator at the University of Freiburg

since 2008, first as a Junior Fellow at the Freiburg Institute of Advanced Studies (FRIAS) and then (since 2014) with his own lab at the Center for Biological Systems Analysis. In 2014, he received the BMBF „Next Generation of Biotechnological Methods“ research prize for his work on universal modular production organisms. His research focuses on the development of complex, functional molecular systems and architectures for both in vivo and in vitro applications, achieved by combining chemical and biological methods in conjunction with nano-/biotechnology.

amino acids in the essential enzymes – in *Escherichia coli*, for example. Control over the cells is achieved by the fact that metabolism is now dependent on dosing with unnatural amino acids that the biosystem cannot itself synthesize.

Our lab is drawing on new bio-based and chemically enhanced methods of synthesis to research new molecules for a wide variety of applications. The primary focus of current research, however, is on novel cell compartmentalization and the implementation of new synthesis pathways by designing and arranging enzyme cascades, for example. New methods for the construction of complex blueprints for proteins permit functions to be programmed by sequence design. If dynamic processes, controllable by small molecules or physical stimuli such as light or temperature, are then combined at the cellular level, this permits the execution of complex interventions (also logic-based and therefore programmable).

## Extending the cell as a mini-factory

One option permitted by the combination of conventional organic synthesis and biosynthesis is the synthesis of unnatural amino acids in the lab. These amino acids are then supplied to the cell for incorporation in proteins by a modified biosynthesis apparatus. This process of extending the genetic code with unnatural or non-canonical amino acids facilitates the selective furnishing of proteins – including cell proteins – with new chemical functions. Examples of this include fluorescently-labeled de novo organelles (see fig. 2) and enzymes whose cofactors have been redesigned so that they are tightly bound to the enzymes. This paves the way for new applications for such enzymes, both in technical procedures and within the cell.

The first such applications comprise highly-elastic, biocompatible protein materials for technical and biomedical applications (such as tissue replacement), protein-based detergents/amphiphiles for de novo organelles/compartments in the cell (fig. 2) and for drug formulations. Predefined protein “donuts” support the LEGO-like arrangement of nano-objects in a process of self-organization akin to mineralization, which supplies the biohybrid materials with new physical properties (visual qualities, magnetic attributes, etc.). Combination with photosynthetic complexes, on the other hand, permits the conversion of light energy into electrical energy based on a 10 nm molecular architecture. These are just a few examples of the kind of potential

offered by the synthesis and manipulation of complex bio-based molecular systems. This is where biochemistry impinges on nanotechnology and materials science – and vice versa.

These approaches form part of the BMBF’s Biotechnology 2020+ initiative and are a key pillar within bio-economic strategy. Achieving resilience in technical and organic systems, by solving the problem of maintaining a chaotic system in its stable phase space, is possible only by implementing dynamism, adaptability, renewal and self-healing – that is to say, biomimetic principles – from the molecular to the macroscopic level. This also implies the enhanced mimicry or simple “co-installation” of biological concepts.

■ [stefan.schiller@frias.uni-freiburg.de](mailto:stefan.schiller@frias.uni-freiburg.de)

### Bibliography

Huber, M. C. et al. (2015) *Nature Materials*, 14, 125–132  
Schreiber, A. et al. (2015) *Nature Communications*, in press, DOI: 10.1038/ncomms7705  
Huber, M. C. et al. (2014) *Biomaterials*, 35, 8767–8779  
Schiller, S. M., „Protein Tectons in Synthetic Biology: The expansion of cellular functionality combining chemical biology of small organic molecules with protein tectons – unnatural amino acids, protein based biohybrid materials & de novo organelles“, in: *Synthetic Biology*, ed. Giese, B., von Gleich, A., Pade, C., Wigger, H. (Springer, 2014, ISBN 978-3-319-02782-1)  
Schreiber, A., Schiller, S. M. „Nanobiotechnology of Protein Compartments: Steps towards Nanofactories“ *Bioinspired, Biomimetic and Nanobiomaterials (ICE Virtual Library)* 2013, 4, 2, 157–164

*“For the first time ever, we have succeeded in forming a new organelle in the cell and equipping it with functions on the basis of rationally designed protein building blocks. This is a fundamentally new medicine.”*

Dr. Stefan Schiller