The Synthetic Microbiology Caucus: from abstract ideas to turning microbes into cellular machines and back

Huang, Wei E.; Nikel, Pablo Ivan

Published in:
Microbial Biotechnology

Link to article, DOI:
10.1111/1751-7915.13337

Publication date:
2019

Document Version
Publisher's PDF, also known as Version of record

Link back to DTU Orbit

Citation (APA):

General rights
Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the legal requirements associated with these rights.

- Users may download and print one copy of any publication from the public portal for the purpose of private study or research.
- You may not further distribute the material or use it for any profit-making activity or commercial gain
- You may freely distribute the URL identifying the publication in the public portal

If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.
Editorial

The Synthetic Microbiology Caucus: from abstract ideas to turning microbes into cellular machines and back

Wei E. Huang1 and Pablo I. Nikel2
1Department of Engineering Science, University of Oxford, Parks Road, Oxford, OX1 3PJ, UK.
2The Novo Nordisk Foundation Center for Biosustainability, Technical University of Denmark, 2800 Lyngby, Denmark.

The core tenet of synthetic biology is the application of engineering principles such as standardization, modularity and rational design to accelerate the design-build-test loop for reprogramming biological system by endowing them with novel tasks (Endy, 2005; de Lorenzo and Danchin, 2008; Church et al., 2014; Badenhorst and Bornscheuer, 2018; Kohman et al., 2018; de Lorenzo et al., 2018). Since its very inception as a field, synthetic biology has enabled researchers from different disciplines to extend and re-think genetic manipulations as the rational design and engineering of cells. Against this background, (micro)organisms can be regarded as programmable cellular machines—which can be modified by manipulating the cells’ software (DNA/RNA), hardware (physical cell components), and the processes encompassing processing and regulation of nucleic acids and, even more importantly, metabolism (which, together, make up for the operation system; Rampley et al., 2017; Danchin, 2009). Moreover, in the context of contemporary synthetic biology practice, the hardware of cells can be purposefully defined by means of genome design and engineering, for which systems biology plays an important role. Synthetic biology and systems biology are viewed, in this sense, as two sides of the same coin. To fulfil this overarching engineering purpose, the ever-expanding synthetic biology toolbox allows for the modification or tuning of gene network connectivities in a precise and in-place manner. Decomposing, manipulating and re-assembling the factors governing the hardware, software and operation system of the cell is crucial to understand its functioning in toto, e.g. for re-factoring—the process by which a set of related genes are removed from their native regulatory context and placed under synthetic expression control.

As an enabling-technology discipline, synthetic biology continues to provide sophisticated tools to precisely and rationally manipulate genetic programs, e.g. by adopting specific gene circuits, genetic toggle switches, amplifiers, sensors, (alternative) memories and oscillators. Yet, synthetic biology is still far from achieving its full potential in fields as diverse as healthcare, environmental protection, energy, agriculture, bio-computing and efficient production of advanced chemicals and materials from renewable sources. Several challenges still remain along the road, e.g. undefined and incompatible bioparts, unpredictable gene circuit design, difficult-to-understand (let alone controlling) biological complexity and the appearance of phenotypic heterogeneities within isogenic populations (Kwok, 2010; Stephanopoulos, 2012; Vilanova et al., 2015; Wu et al., 2016). Furthermore, cells are living systems and, as such, they are constantly evolving (Bull and Barrick, 2017). Because of this unavoidable circumstance of every biological system, the introduction of designed gene circuits will likely interfere and cross-talk with the gene networks defined by the indigenous genome. Such interactions between gene circuits and the host genome will influence the performance of gene expression and even crash with the functions of the implanted genetic devices. Moreover, since synthetic biology ultimately aims at programming cells that can execute the implanted functions in a predictable fashion, the adoption of specific, formatted hosts (biological chassis), endowed with different properties depending on the application envisioned, is still a matter in dispute (Calero and Nikel, 2018). Finally, several issues surrounding the use of genetically modified (micro)organisms, including intellectual property protection and open access to biological materials (Carbonell et al., 2016; Kahl et al., 2018), and the need of establishing efficient biocontainment strategies are also barriers in implementing synthetic biology, especially in practical (field) applications (Rambley et al., 2017; Whitford et al., 2018).

What is the state of affairs and the way ahead to tackle these challenges in the field? A combination of...
top-down and bottom-up approaches is used in synthetic biology, and the rational removal of genes and entire genomic regions has helped understanding the functions of genes and instruct the cell design in synthetic biology (Hutchison et al., 2016). The minimal genome project has helped to interrogate the essential genes that defines life, and the assembly of individual yeast chromosomes has been key to understand genetic regulations in a model microorganism. Looking at the hardware afresh has enabled to gain insight in structural organization of the bacterial cell, and a recent milestone in this sense has been the in vitro reconstitution of the (long neglected) bacterial cytoskeleton (Ramm and Schwille, 2018). Thus, understanding the wiring and functioning of the microbial cell at the structural level is a stepping stone in which synthetic biology can play a decisive role.

In this sense, the metabolic operation system has been placed under the spotlight after years of undeserved ostracism (Nielsen, 2017), and the implementation of synthetic metabolisms has resulted in the design and construction of specialized cell factories that can utilize a range of substrates to produce complex, added-value molecules (Smanski et al., 2016; Aslan et al., 2017; Nikel and de Lorenzo, 2018). These approaches rely on emerging analytics technologies for high-resolution mapping of macromolecules and metabolites (Wang et al., 2016) supported by multi-omic analyses and big data (Danchin et al., 2018). The use of xenonucleic acids is another fascinating aspect that continues to help expanding the repertoire of new-to-Nature cell functionalities (Schmidt et al., 2018); Furthermore, microbes usually (co-)exist in communities, instead of isolated, single species. Some of the next major challenges would move from engineering single species towards the design of truly synthetic and programmable microbial communities, which would be able to perform more complex functionalities such as degradation of recalcitrant chemicals (Pelz et al., 1999)—taking full advantage of the division of metabolic labour (Brenner et al., 2008; Thommes et al., 2018), a feature that characterizes natural microbial communities (Nikel et al., 2014). Establishing integrated community systems with high-level functionality would be the next step in this direction by understanding the factors governing assembly and maintenance of stable microbial consortia and synthetic microbiomes (Pham et al., 2017). Besides the clear applied angle of these approaches, synthetic biology now offers the tools and enabling technologies to explore both cellular and metabolic interactions of bacteria with other organisms beyond microbes, such as phages and potential hosts (both as symbionts and parasites). Understanding these processes from an engineering perspective will shed light on the bacterial lifestyle in different environments, the consequences of which will be relevant for fields as diverse as agriculture and medicine. Last, but certainly not least, most of the examples currently handy deal with the bacterial cell as a whole, but the adoption of cell-free systems is also gaining momentum in the synthetic biology arena (Hodgman and Jewett, 2012).

Although the detailed enumeration of all the challenges that the synthetic biology community is facing would be burdensome, we believe that these aspects to constitute the next grand challenge for the field. An exciting brainstorming with Profs. Víctor de Lorenzo (CNB-CSIC, Spain), Philippe Marlène (GénoPole, France) and Kenneth Timmis lead to the inception of the Synthetic Biology Caucus: a novel section of Microbial Biotechnology, which will provide an agora for sharing, documenting and discussing fresh ideas—from the entirely abstract ones to completely applied biotechnology designs—and to attract critical and constructive comments from the members of the synthetic biology community. Contributors to the section will communicate their thoughts in a dynamic format to identify key (either original or long standing) research questions that could prompt projects and experiments. The drafting of grant proposals represents a specific case in which many of the arguments that are elaborated or vented during the process have an intellectual value in themselves and would deserve to be shared and discussed in the community. The section will be run by Pablo I. Nikel (The Novo Nordisk Foundation Center for Biosustainability, Denmark) and Wei Huang (Oxford University, United Kingdom). Pablo’s expertise is focused on metabolic engineering and synthetic metabolism in bacteria; Wei’s competences are related to biosensors and bioenergy, and single cell omics. The editors will work together with the members of the community to move forward the concepts of synthetic biology and to promote delivery of the many promises that the field entails. Although we will invite potential contributors to submit their articles to this section, we strongly encourage all synthetic biologists from both fundamental and applied fields to contribute ideas, suggestions and arguments to the Caucus: this will be an exciting opportunity to foster open communication and scientific brainstorming with a wide reach out in the community!

References