

**TITLE: The long journey towards standards for engineering biosystems**

**SUBTITLE: Is the Molecular Biology and the Biotech community ready?**

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Standards are the basis of technology: they allow rigorous description and exact measurement of properties, reliable reproducibility, and a common ‘language’ that enables different communities to work together. Molecular Biology was in part created by physicists; yet, the field did not inherit the focus on the quantitation, the definition of systems boundaries and the robust, unequivocal language that is characteristic of the other natural sciences. However, Synthetic Biology (SynBio) increasingly requires scientific, technical, operational and semantic standards for the field to become a full-fledged engineering discipline with a high level of accuracy in the design, manufacturing, and performance of biological artefacts. Although the benefits of adopting standards are clear, the community is still largely reluctant to accept them, owing to concerns about adoption costs and losses in flexibility.

#### **SUBHEADER: What standards are good for**

In science and technology, the terms *standard* and *standardisation* describe different things: shared semantic and graphical languages for annotating the nature and the properties of systems and their components; the definition of units of relevant properties and parameters along with methods to calculate them; specifications of properties and arrangements for the physical assembly of the components of a system; and unambiguous protocols for the construction of objects. Such standards enable an abstract and precise description of a system with a suitable—also standardized—quantitative language or equivalent methods of representation.

Beyond their important role in the natural sciences, standards were also one of the key drivers for the industrial revolution as they enabled a seamless integration of product design, fabrication of its components and the final assembly—let alone tracing parts and helping to sort out matters of safety and intellectual property. Standards are for instance imperative for designing electronic circuits built from well-defined, universal simple components, such as resistors, diodes, transistors, or for software engineering that uses precompiled modules and functions. Standards enabled the rapid rise of the personal computer industry in the 1980s and 90s by interlinking standard components such as hard disks, memory or keyboards through standardized interfaces and protocols.

From software to nuts and bolts, the concept of a universally usable toolbox of parts to assemble more complex systems is typical for every discipline of engineering: electronics, software, mechanical design, architecture, chemical synthesis, and so on. Standards enable people to work together through interoperability, coordination of labour, reproducibility and reuse of other people's efforts and achievements.

Standards must be reliable, robust and affordable, but, first and foremost, they must be agreed on by their users. Indeed, standardisation—the process of implementing and developing technical standards—requires the consensus of many different parties, such as private and public companies, organizations and policy makers. Standardisation can be driven by public acceptance/market forces (*de facto* standards), directly ordained by law (*de jure* standards) or, most commonly, arise from the combination of legal/technical requirements and recognition by potential operators since, in general, the broader the applicability of a format, the greater its market (Dan, 2019).

#### **SUBHEADER: Standards in the life sciences**

That said, the core standardisation process in many scientific and engineering disciplines took place decades to centuries ago, but it is still in its infancy in the life sciences. Interestingly, it is still a bottleneck for even well-developed technologies: smartphones, for instance, still lack standard key components such as batteries or electric charger cables ([https://ec.europa.eu/info/law/better-regulation/initiatives/ares-2018-6427186\\_en](https://ec.europa.eu/info/law/better-regulation/initiatives/ares-2018-6427186_en)).

In this context, the conceptual frame of synthetic biology aims to making biology easier to engineer by applying principles such as modularity, orthogonality, chain production and reproducibility. Moreover, the rapid advances in wet and computational tools for genome editing, metabolic design and *in silico* modelling are opening new opportunities for genetic programming that could not have been anticipated even just a few years ago, and allow engineers to tackle increasingly complex engineering objectives. The growing demand for scaling up such technologies raises the issue of what is needed to make them work at an industrial scale (Beal et al, 2018). Following the path of other branches of engineering, establishment of standards appears among the key objectives of contemporary SynBio – and eventually of the life sciences as a whole – as a prerequisite

for applications such as bioremediation, biomedicine, bioenergy, novel chemicals, innovative materials and cellular factories.

Although standards in SynBio have contributed to successes such as the synthesis of artemisinin or morphine (both in yeast), the problem of defining common standards is still far from being resolved. The reusability patterns of the iGEM parts database (Vilanova & Porcar, 2014), the context-dependence of biological components (Carr et al., 2017), the variable behavior among strains, genetic stability, or even the contested philosophical analogy between cells and machines are by no means solved issues at this point. However, there is no doubt that even partial progress on standardisation would have major consequences for bioengineering.

One bottleneck is the widespread and incorrect assumption amongst many researchers in the life sciences that standards may increase inter-operability but necessarily limit flexibility—which is obviously important for any creative research. Rather, good standards will increase people’s flexibility and creativity because it will make it easier for them to achieve their scientific objectives. A separate challenge is identifying specific systems and operations that need to be standardized, and then navigating the minefield of personal interests that typically inhibit agreement on a given format or language. As Murray Gell-Mann quipped, “a scientist would rather use someone else's toothbrush than someone else's nomenclature”. Scientists and engineers will adopt standards only when they add value to their efforts to overcome the often steep costs of adoption.

### **SUBHEADER: Standards for engineering biology**

While a number of SynBio standards have already been developed and await adoption by the broader community of users (de Lorenzo & Schmidt, 2018), others touch on core biological questions that are by no means solved from a scientific point of view. There is a legitimate concern that we still need to know more fundamental facts before we can describe engineered biosystems with a formal, unequivocal language. One typical case involves the design of genetic circuits, an archetypal product of SynBio endeavours. Habitual practices include directly transplanting toolkit for building electronic logic gates and related information-processing devices into the biological domain. However,

one must be honest about how far these abstractions and their accompanying theoretical framework reflect biological reality. Boolean logic relies on values that are either true or false. In electronics, this is readily implemented using voltage levels that are separated by a larger amount than the expected noise to faithfully represent the state of the gate. In contrast, biological implementations of circuits tend to have a much higher noise to signal ratio, which makes it difficult to effectively distinguish true and false states and strongly limits the design of logic circuits. One way to alleviate this problem is by redesigning regulatory components to behave more digitally, but ultimately, we may need to revisit information processing in/by biological systems with other formalisms, either existing or yet to be developed, that go beyond Boolean logic (Grozingier et al., 2019).

The same theory/implementation conundrum might be true for biological metrology, one of the main tenets of SynBio. Electronic circuits crucially rely on a clear definition of *potential* and *current*, their description in *volts* and *amperes*, and methods to measure these. By the same token, it is difficult to think about genetic circuits without robust measures of signal transmission through the regulation of gene expression or other core cellular processes. The concepts of RNA *polymerase per second* (PoPS; Endy, 2005) and *ribosome per second* (RiPS) as biological counterparts of *current* were conceptualized early in the history of SynBio. Alas, very little has been done to further develop these units as practicable indicators of genetic circuit performance, perhaps due to the difficulties of measuring them accurately.

These examples showcase how developing standards for biological engineering still requires addressing a number of core scientific and technological gaps that have been left behind in the ongoing frenzy of application-focused development. Yet, such unsolved issues may strike back when the field continues to move from largely academic endeavours towards industrial realisation.

#### **SUBHEADER: Key actors in the standards conversation**

International discussions about SynBio standards, mostly with US and EU stakeholders, have been going on since before 2010. Under the umbrella of the BIOROBOOST Project (<http://standardsinsynbio.eu>), the conversation now incorporates key actors of

SynBio from Europe, North America and Asia. Much of the discussions deal with identifying key challenges for the development, promulgation, and adoption of standards, and identifying stakeholders in academia, industry, research centers and politics.

The most conspicuous technical challenges include standardizing simple biological parts, devices and circuits, chassis, metrology, descriptive languages (including graphical representations), and software tools. But the complexity of the endeavour also asks for the creation of a network of SynBio practitioners that share and evolve these standards together. While this is reminiscent of earlier Computational Modeling in Biology Network (COMBINE, <http://co.mbine.org/>), the focus of these SynBio networks needs to go beyond academic interests to include industry and commerce, and to develop strategies for educating a new generation of synthetic biologists who routinely use standards.

From the regulatory, technical and societal point of view, the challenge is complex. For example, there are practical questions such as the level of detail required in a given biological standard, which can go from light to very deep. As indicated above, *standard* is an umbrella concept, which includes a number of different approaches to harmonization. These range from agreeing on *metrology* units and best practices to measure them, to developing standardized functional chassis – specific, formatted biological hosts for specific applications – to data formats, to safety criteria for approval by regulatory agencies, and to ISO-approved Reports and Technical Specifications.

It is necessary to distinguish between *biological standards* that could be similar to physics and engineering counterparts, such as the PoPS or RiPS units discussed above, and *standard operating procedures* (SOPs), which help users to carry out routine operations with efficiency, consistent quality and performance, and are compliant with regulations. For instance, the composition and preparation of the M9 medium would be an SOP, while the metrics for calculating containment of a given SynBio agent when released in the environment could become a biological *standard*. There are, of course, many grey zones between these two – for instance, formats for enabling communication between unrelated software, cloning methods, CRISPR-based editing, and so on – that will hopefully be solved through conversations between stakeholders in the various

forums just mentioned. The question remains, however, whether the wider community of potential users will see the value of adopting standards in their daily practice. Today, SynBio and Systems Biology practitioners are widely using the Synthetic Biology Open Language SBOL (Galdzicki et al., 2014) and SBOL visual for describing vectors and constructs (Martínez-García et al., 2020), and there is a great consensus on the need to go beyond the state of the art and further advance towards the standardisation of biological systems (Schreiber et al., 2019; de Lorenzo & Schmidt, 2018).

### **SUBHEADER: Stages of adoption**

Is there a take-home lesson from the history of technology adoption that we can learn from for popularizing biological standards? In fact, the trajectory of acceptance in the realm of engineering typically involves several stages: from an innovator phase to adoption by even the most recalcitrant laggards (Fig. 1). Using this frame, it seems that most of SynBio's standards developments are still in the innovators phase.

Many developments, even if critical for the early years in SynBio, never left the innovator state and are now outdated; advances in cloning and DNA synthesis have for instance replaced BioBricks. Others, such as SBOL (Galdzicki *et al.*, 2014) or the Standard European Vector Architecture (SEVA; Martínez-García et al., 2020) are increasingly successful as interim formats in the early adopter stage. Yet, these may or may not become generally adopted depending on success stories and potential alternative scientific and technical solutions. Such progress will be determined by the combination of a bottom-up demand for interoperability and collaboration and a top-down implementation and enforcement by official agencies. Journal editors also have a role to play as well as reviewers of journal articles and grant proposals in insisting on the use of standards to improve reproducibility and reuse. Generally, it is important to realise that standards are ultimately social constructs to represent norms, objects or procedures, and that they become accepted by a group of individuals for practical reasons.

### **SUBHEADER: Low-hanging fruits**

Despite the difficulties, it should be possible to come up with science-based standardisation proposals in SynBio that work across the biological, the digital and the social realms. The already existing ones at hand involve simple biological parts: devices such as promoters and other regulatory nodes and simple circuits – for instance, inverters, basic gates – such as those deposited in the repository of biological parts and other curated collections. The next stage involves definition and adoption of SynBio chassis other than laboratory bacteria or yeast strains. Not every species or strain that can host recombinant DNA can be considered a chassis, and this effort requires establishment of a map of requirements and functional relationships between industrially relevant practical applications and different biological platforms. Finally, standardisation would need to address the issue of metrology through the gene expression flow including, for example, fundamental units and the technologies and references to measure them, as well as computational language and software tools for easing collaborations between different actors. The main efforts to collect such low-hanging fruits would be greatly facilitated by biofoundries with good connections to policy makers with the objective of making the whole endeavour more appealing for the industrial sector.

The academic community cannot be a mere observer of these developments. In fact, there is much to do for endowing biological standards with a solid scientific basis, including the definition of each level of biological complexity amenable to standardisation. But the role in promoting standards is not only technical. There is ample room for networks of practitioners involving industrial players, who can provide information on how biological properties and processes could improve product development, manufacturability and consumer confidence. This could create a framework for identifying and monitoring standardisation requirements and maintaining an evolving list of scientific and industrial priorities. Ideally, such priority lists should also be considered by funding bodies to help in developing and driving adoption of standards. Relevant regulatory bodies should be involved to adapt or ease rules on the management of GMOs and/or SynBio agents. The same academic-industrial networks could also strengthen ongoing public outreach and citizen involvement to help overcoming the negative perception of genetic engineering in general.

In sum, we argue that the promise of SynBio for the benefit of global society and industry will only be met if significant advances are achieved on the standardisation front. To this end, it is not only essential to overcome national/political barriers and particular interests of given research groups, but also to gather key players in a permanent forum with the aim of making biological standards one of the ingredients of the 4<sup>th</sup> Industrial Revolution. Standards in biology will be used provided that they have intrinsic properties such as robustness, ease of use, context-independence, etc. But the key to success is the merger of technical consistency and scientific soundness with legal requirements and consensus among end users. This goes beyond the realm of research and tackles sociological and cultural issues that have been traditionally alien to the conversation. If this can be achieved, the benefits for SynBio and for society at large will be great.

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## **CONFLICT OF INTEREST**

The authors declare that they have no conflict of interest.

## **FIGURE LEGENDS**

**Figure 1:** Illustrative examples of the position of SynBio standards along the technology adoption curve: SynBio standards are largely still in the innovators phase but with a few examples having progressed to the early adopters or early majority

segments. SEVA: Standard European Vector Architecture; SBOL: Synthetic Biology Open Language; MoClo: Modular Cloning

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