

1 **Reconfiguring the challenge of biological complexity as a resource for bio-design**

2 Erika Szymanski

3

4 Department of English

5 Colorado State University

6 359 Willard O. Eddy Hall

7 Fort Collins, CO 80523-1773

8 Erika.szymanski@colostate.edu

9

10 James Henriksen

11

12 Natural Resource Ecology Laboratory

13 Colorado State University

14 Fort Collins, CO 80523-1499

15 James.henriksen@colostate.edu

16

17

18 **Abstract:** Biological complexity is widely seen as the central, intractable challenge of engineering biology. Yet this
19 challenge has been constructed through the field's dominant metaphors. Alternative ways of thinking—latent in
20 progressive experimental approaches, but rarely articulated as such—could instead position complexity as
21 engineering biology's greatest resource. We outline how assumptions about engineered microorganisms have
22 been built into the field, carried by entrenched metaphors, even as contemporary methods move beyond them.
23 We suggest that alternative metaphors would better align engineering biology's conceptual infrastructure with
24 the field's move away from conventionally engineering-inspired methods toward "biology-centric" ones.
25 Innovating new conceptual frameworks would also enable better aligning scientific work with higher-level
26 conversations about that work. Such innovation—thinking about how engineering microbes might be more like
27 user-centered design than like programming a computer or building a car—could highlight complexity as a
28 resource to leverage, not a problem to erase or negate.

29
30 **Importance:** Biological complexity—the many-parted, highly interconnected, highly responsive character of living
31 things—is widely seen as the greatest challenge of engineering biology, because living things respond to
32 engineering interventions in hard-to-predict ways. That challenge, we argue, is a product of the metaphors that
33 frame how engineering biology is understood. Different metaphors could instead make it possible to leverage
34 biological complexity as engineering biology's greatest resource, to be productively employed rather than
35 avoided or erased. Organism-centric approaches are gaining ground as alternatives to (more) conventionally
36 engineering-modeled modes of synthetic biology. Employing language that consciously embraces this shift is
37 likely to inspire new experimental approaches to designing with biology that are otherwise difficult to imagine.

38
39 **Perspective**

40
41 Engineering biology is suffering from a lack of imagination, at least in the language used to conduct it.
42 From its inception, biological complexity has been engineering biology's central challenge¹. Biological systems
43 involve seemingly uncountable, highly interconnected components, many of which remain poorly described and
44 which are not necessarily fully decomposable². Engineering biology involves abstracting these complex systems
45 into machine-like, discrete, interchangeable parts^{3,3}. Yet projects are routinely stymied when biological parts
46 interact with context, affecting function in unplanned ways^{4,5}—in other words, when organismal behavior
47 exceeds the machine analogy.

48 Recent experimental developments have diverged from its foundational (non-living) engineering
49 analogies, becoming more "host"- or "organism"-centric, adding more biology back into the picture^{4,5}.
50 Meanwhile, the conceptual infrastructure for designing and building with biology has lagged behind, largely
51 continuing to construct complexity as an engineering challenge through analogizing living things to cars and
52 computers. We suggest that a different set of metaphors or conceptual tools for imagining what microbes (and
53 other cells) are like might reconfigure complexity as engineering biology's greatest resource, not its greatest
54 barrier.

55 Synthetic biology has been built on the back of two analogies: first, that cells are machines operated by
56 genomes; second, that if genomes are information-storage molecules written in genetic code, scientists should be
57 able to program cells in the same way that they program computers. The conceptual and physical infrastructure
58 of the field is so tightly linked to these analogies that they have become assumptions embedded in the way things
59 get done.

60 "Host-aware" strategies have begun to shift away from a strict application of these analogies,
61 incorporating more biological context into engineering approaches. Some aim to alleviate the metabolic burden
62 of genetic constructs by decoupling heterologous gene expression from autochthonous processes⁶; others fine-
63 tune engineered pathways in light of a larger picture of resource-flux within the cell⁷. Yet while some contextual

64 variables can be quantified and modeled, all but the most deliberately constructed biological systems involve far
65 too many potentially relevant parts and connections to comprehensively characterize. The residual
66 undifferentiated details are routinely glossed as “complexity,” especially when they interrupt or impede design
67 goals.

68 Because of complexity, rational approaches to bio-design become resource-intensive cycles of tinkering
69 with a genetic assembly until it works. The frequent disconnect between the predicted and observed function of
70 DNA designs has led some to say that engineering biology is not “really engineering,” on the basis of a limited
71 vision of engineering that has permeated much of the field^{9,10}. That disconnect has also been blamed for why
72 bioengineering chronically lags behind the developmental trajectory anticipated for it on the basis of cognate
73 trajectories of engineering computers. Decoupling biological parts into (hopefully) standardized modules has
74 enabled relatively simple designs that intersect with a cell’s dense, recursive regulatory networks at a limited
75 number of points. However, as design become more ambitious, so too does the likelihood of encounters with
76 complexity exceeding the practical capacity to tinker—even when orthogonal control systems are designed to
77 avoid interfering with cellular processes^{3,10}. Moreover, high-level descriptions in inherited engineering terms hide
78 tinkering under the umbrella of the ubiquitous design-build-test (DBTL) cycle, feeding unrealistic expectations
79 about what is scientifically plausible.

80 A promising bottom-up approach to addressing the challenge of complexity involves building synthetic
81 living systems from well-described parts, limiting the number of parts in the system to the minimum needed to
82 achieve a specific function^{13,14}. An alternative, top-down approach involves constructing minimal cells by
83 beginning with existing organisms and then eliminating genetic (and perhaps other) elements found to be
84 unnecessary under specific conditions. A third strategy involves tailoring parts-based assemblies to account for
85 cellular conditions, using modeling or directed evolution.

86 A limitation of all three approaches is that they are likely to construct cells adept at doing one thing under
87 narrowly defined conditions^{12,13}; in contrast, precisely because they are complex and thus responsive and
88 adaptable, microbes and mammalian cells are adept at doing many things under shifting conditions. A second
89 limitation is that they are difficult to achieve. The comprehensive whole-cell model that would ideally anchor the
90 third strategy, in particular, remains out of reach; simplifications must be made. Again, because of complexity,
91 identifying what can be safely eliminated from a model—or a cell—without disabling its utility, disrupting
92 essential cell functions, or causing other genes to become essential or deleterious in turn is tricky. A third
93 difficulty is that, as minimal genome projects have illustrated, a large number of genes are required to sustain life,
94 even under highly controlled conditions, but for unidentified reasons.

95 Some organism-centric approaches, such as directed evolution strategies, employ complex cellular
96 responsiveness as a design strategy rather than an interruption to design—delegating cycles of trial-and-error to
97 “evolution,” or, we could say, to complex cell-environment interactions. Our proposition is that engineering
98 biology would benefit from building on this opening by making the tension between the field’s foundational
99 machine analogies and contemporary organism-centric approaches explicit, and by innovating alternatives.
100 Experimenting with metaphors that do not reproduce the assumption that cells look or should look like
101 computers and cars, we suggest, is likely to invite additional strategies for employing complexity as a resource
102 rather than a problem to be overcome.

103 Synthetic biology’s central analogies—expressed in such ubiquitous conceptual infrastructure as the DBTL
104 cycle—configure biological systems as *imperfect* machines because their complexity gets in the way of predictable
105 sequence-function modularity. What happens if, instead of imagining cells as imperfect machines and trying to
106 make them simpler, engineering biology involves imagining cells as being really good cells? Multi-part, redundant,
107 recursive, interacting functional systems enable cells to grow, reproduce, and maintain tightly regulated, finely

108 tuned responses to environmental change. Their responsive and self-amplifying capacities are a major part of why
109 biological systems are useful technologies in the first place. Engineering biology is exciting precisely because
110 machine analogies are imperfect. Cells are imperfect machines, but life is great at being alive.

111 Machine analogies limit the range of conceivable bio-design strategies by embedding several
112 assumptions: 1) that biological systems should be made increasingly passive and controllable, 2) that unplanned
113 biological responses constitute undesirable interference in design, and 3) that engineering biology is lagging
114 behind conventional engineering along an established trajectory for how the field is supposed to develop.
115 Because these assumptions are carried along with “dead” analogies—analogies easily employed without
116 recognizing them as analogies—they become less visible as choices that could be made differently. Their
117 influence remains visible even, for example, in justifications of directed evolution as a stop-gap measure en route
118 to better rational design.

119 Alternative analogies might enable engineering biology to better leverage what could be called cellular
120 expertise, accounting for and working with their responsiveness rather than trying to engineer it away. Where
121 machine analogies suggest disassembling complex networks into decoupled parts, such that they are better
122 defined but no longer responsive, organism-centric analogies suggest working with complex systems’ capacity to
123 respond to change. This shift builds on extant movements in the field to reconfigure biological complexity as a
124 valuable resource rather than an intractable challenge.

125 We see such organism-centric perspectives as being grounded in three principles:

126 **Engineering biology differs from engineering automobiles or semiconductors**, not because living and non-living
127 systems fundamentally differ, but because organisms are not the product of human design practices that make
128 establishing predictive principles for structure-function relationships easy. Biological systems may therefore be
129 said to “know” things that researchers do not, and may respond to change in ways that researchers would not
130 have anticipated and may not be able to intentionally recapitulate.

131 **Achieving design goals is more useful than making biology into a particular kind of engineering discipline.**
132 Making biology into an engineering discipline prioritizes increasing control over biological systems and reducing
133 their complexity by decomposing networks into discrete parts. This is a very different goal than trying to design,
134 build, and implement useful biological technologies—the direction in which directed evolution and some other
135 host-aware strategies are refocusing the field. This direction might be better served by building effective working
136 relationships with biological systems, with less focus on control and more focus on outcomes.

137 **Intervening in biological systems is about communication.** In 1934, the biologist and proto-cybernetician Jakob
138 von Uexküll suggested that all living things inhabit their own *umwelt* or lifeworld, comprised of the phenomena
139 that an organism can sense and effect¹⁶. Organisms communicate with each other when and only when their
140 respective *umwelten* overlap. Organisms can expand their *umwelten* via what von Uexküll called prostheses, or
141 (broadly defined) technologies; scientists, for example, expand their *umwelten* with DNA sequencing, while
142 bacteria expand their *umwelten* with horizontal gene transfer. We envision engineering biology as being about
143 expanding the overlaps in *umwelten* among scientists and the organisms with which they work so that they can
144 share overlapping goals and effectively communicate toward achieving them.

145 To exemplify what such an approach might entail, we reconsider the DBTL cycle as applied to designing
146 microbial consortia. Following the DBTL cycle customarily means imagining and constructing a design in silico,
147 building that design from synthetic or extracted and amplified DNA, loading the assembly into a biological system
148 to test its function, and learning from what does and does not work to inform a better design. In microbial
149 consortia engineering, this process may be repeated at several levels of hierarchy to customize microbial strains
150 that are then assembled into a synthetic community, or that are introduced into an existing community.

151 An organism-centric frame that explicitly accounts for microbial responsiveness might reconfigure the
152 DBTL cycle as the Listen-Parse-Respond (LPR) cycle: “listen” to the microbe or microbial consortia, “parse”
153 relationships among microbial communications and researcher goals, and “respond” with an informed
154 intervention to continue the conversation. Genetic material has been analogized to encrypted human or encoded
155 computer languages to make identifying and interpreting genetic “words” analogous to making sense of human
156 language use, and to develop new techniques through that analogy^{17, 18, 19}. We extend this analogy by imagining
157 genetic statements as dialogue or discursive resources that microbes use to negotiate environments and that
158 microbes and humans can use to communicate with each other. Consequently, LPR workflows might resemble
159 other negotiated communication scenarios, and might encourage more diverse “conversational” strategies in
160 organism-centric experiments. Microbial consortia design might, for example, be described in terms of:

- 161 • **User-centered (participatory) design:** Researchers configure design goals in terms of a problem that can be
162 shared with the microbial community required to enact it, such as how a microbial community with particular
163 characteristics can thrive under particular conditions, and then invite (microbial) users to participate in the
164 design of a solution to the problem. Directed evolution experiments can be seen as participatory design
165 experiments in which scientists equip microbes with a technology (one or more novel genetic statements),
166 ask microbes to use that technology to solve a design problem in the form of a challenging environment,
167 parse the responses of the most successful, and respond with an additional challenge that advances toward a
168 functional design that becomes, effectively, a shared goal (see, e.g.,²⁰⁻²²).
- 169 • **Marketing:** Researchers aim to convince a microbial community to adopt specific practices with novel
170 (genetic) resources. To do so persuasively, they need to evaluate and account for microbial responses
171 regarding product, place (context), price (metabolic cost), and promotion (delivery and incentive to retain and
172 continue using the genetic construct) (see, e.g.,⁴). Reconfiguring typical experimental parameters through this
173 frame may enable making more deliberate and varied use of microbial responsiveness as valuable data rather
174 than a barrier to enacting a design.
- 175 • **Public engagement:** Researchers aim to dialogue and negotiate with microbial stakeholders to identify a
176 communally acceptable route toward a technoscientific aim. Through this frame, crafting enrichment culture
177 conditions could be seen as a parallel to providing public spaces for mutually beneficial activities.

178 These ideas may sound dangerously anthropomorphic, but we could just as easily say that engineering metaphors
179 are dangerously mechanomorphic. Scientific reasoning is intrinsically analogical, because to apprehend and make
180 sense of as-yet unknown phenomena, we must have some idea of what they are *like*. Metaphors such as the DBTL
181 cycle structurally embed analogical reasoning in language, such that doing science without metaphors is
182 impossible¹⁹. Problems therefore arise not because a metaphor is employed, but because the metaphor and the
183 assumptions it carries may be unhelpful for a particular purpose, and because it becomes invisible as a metaphor
184 that describes some but not all of a phenomenon’s characteristics. Focusing on machine-like capacities may be
185 less useful than focusing on microbes’ responsive organism-like capacities for achieving bio-design goals that
186 involve contextual dependencies. While the necessity of bioengineering standards is often articulated, a diverse
187 set of approaches to bio-design should expand the long-term resilience of the field and the scope of what it can
188 attempt, in contrast to locking all projects into the same underpinning analogy.

189 Numerous philosophy of biology papers detail how organisms are not machines so that they can describe
190 why engineering biology fails^{22,23}. We are far more interested in how engineering biology succeeds. We are,
191 additionally, not concerned with the ethics of whether organisms *should* be analogized to machines. Instead, we
192 are interested in how to develop successful co-working strategies with organism-technologies in light of their
193 distinctive capacities. Leveraging responsiveness and complexity requires rethinking the analogies that underpin

194 bio-design. Doing so will not be a panacea for bio-design challenges. However, re-examining institutionalized
195 assumptions through alternative paradigms may prompt new design strategies in a clogged space.

196 Biotechnology's most practical and versatile successes are, arguably, furthest from idealized "real"
197 engineering, from recent successes with directed evolution to dynamic self-adapting microbial communities that
198 power wastewater treatment plants²⁴, spontaneous sourdough bread ferments²⁵, and microbial ecosystem
199 contributions to sustainable agriculture²⁶. Biological complexity is a strength in these applications in that
200 microbes—individually and communally—resiliently adapt to changing circumstances while maintaining a
201 functional identity. As Wei and Endy have argued in describing where modularity fails in constructing living
202 systems from non-living parts, researchers (re)make systems in the image of what they expect them to be². Given
203 the diversity of technical approaches now available, engineering biologists have choices: to erase complexity to
204 make cells stupider, or to develop strategies to work with their intelligent complexity. We think that the latter is,
205 at the very least, equally promising.

206

207

208 **References**

- 209 1. Endy, D. Foundations for engineering biology. *Nature* **438**, 449–453 (2005).
- 210 2. Wei, E. & Endy, D. *Experimental tests of functional molecular regeneration via a standard framework for*
211 *coordinating synthetic cell building*. 2021.03.03.433818
- 212 <https://www.biorxiv.org/content/10.1101/2021.03.03.433818v1> (2021) doi:10.1101/2021.03.03.433818.
- 213 3. Elowitz, M. B. & Leibler, S. A synthetic oscillatory network of transcriptional regulators. *Nature* **403**, 335–338
214 (2000).
- 215 4. Boo, A., Ellis, T. & Stan, G.-B. Host-aware synthetic biology. *Curr. Opin. Syst. Biol.* **14**, 66–72 (2019).
- 216 5. Darlington, A. & Bates, D. G. Host-aware modelling of a synthetic genetic oscillator. *Annu. Int. Conf. IEEE Eng.*
217 *Med. Biol. Soc. IEEE Eng. Med. Biol. Soc. Annu. Int. Conf.* **2016**, 1463–1466 (2016).
- 218 6. Darlington, A. P. S., Kim, J., Jiménez, J. I. & Bates, D. G. Dynamic allocation of orthogonal ribosomes facilitates
219 uncoupling of co-expressed genes. *Nat. Commun.* **9**, 695 (2018).
- 220 7. Gutiérrez Mena, J., Kumar, S. & Khammash, M. Dynamic cybergenetic control of bacterial co-culture
221 composition via optogenetic feedback. *Nat. Commun.* **13**, 4808 (2022).
- 222 8. Arkin, A. Setting the standard in synthetic biology. *Nat. Biotechnol.* **26**, 771–774 (2008).
- 223 9. Davies, J. A. Real-World Synthetic Biology: Is It Founded on an Engineering Approach, and Should It Be? *Life* **9**,
224 (2019).
- 225 10. Glass, J. I. *et al.* Essential genes of a minimal bacterium. *Proc. Natl. Acad. Sci.* **103**, 425–430 (2006).
- 226 11. Purnick, P. E. M. & Weiss, R. The second wave of synthetic biology: from modules to systems. *Nat. Rev. Mol.*
227 *Cell Biol.* **10**, 410–422 (2009).
- 228 12. Hutchison, C. A. *et al.* Design and synthesis of a minimal bacterial genome. *Science* **351**, aad6253 (2016).
- 229 13. Kaminski Strauss, S. *et al.* Evolthon: A community endeavor to evolve lab evolution. *PLoS Biol.* **17**, e3000182
230 (2019).
- 231 14. von Uexküll, J. *A Foray into the Worlds of Animals and Humans with A Theory of Meaning*. (University of
232 Minnesota Press).
- 233 15. Searls, D. B. The Linguistics of DNA. *Am. Sci.* **80**, 579–591 (1992).

234 16. Searls, D. B. The language of genes. *Nature* (2002) doi:10.1038/nature01255.

235 17. Kim, Y.-A. & Przytycka, T. M. The language of a virus. *Science* **371**, 233–234 (2021).

236 18. Gelfand, M. S. Genetic language: metaphore or analogy? *Biosystems* **30**, 277–288 (1993).

237 19. Szymanski, E. & Scher, E. Models for DNA Design Tools: The Trouble with Metaphors Is That They Don't Go
238 Away. *ACS Synth. Biol.* acssynbio.9b00302 (2019) doi:10.1021/acssynbio.9b00302.

239 20. Szymanski, E. & Calvert, J. Designing with living systems in the synthetic yeast project. *Nat. Commun.* **9**,
240 (2018).

241 21. Sandberg, T. E., Salazar, M. J., Weng, L. L., Palsson, B. O. & Feist, A. M. The emergence of adaptive laboratory
242 evolution as an efficient tool for biological discovery and industrial biotechnology. *Metab. Eng.* **56**, 1–16
243 (2019).

244 22. LaCroix, R. A. Automation, Optimization, and Characterization of Adaptive Laboratory Evolution. (University
245 of California, San Diego, 2016).

246 23. Lowney, C. Rethinking the Machine Metaphor Since Descartes: On the Irreducibility of Bodies, Minds, and
247 Meanings. *Bull. Sci. Technol. Soc.* **31**, 179–192 (2011).

248 24. Nicholson, D. J. Organisms ≠ Machines. *Stud. Hist. Philos. Sci. Part C Stud. Hist. Philos. Biol. Biomed. Sci.* **44**,
249 669–678 (2013).

250 25. Nicholson, D. J. The machine conception of the organism in development and evolution: A critical analysis.
251 *Stud. Hist. Philos. Sci. Part C Stud. Hist. Philos. Biol. Biomed. Sci.* **48**, 162–174 (2014).

252 26. van den Belt, H. Playing God in Frankenstein's Footsteps: Synthetic Biology and the Meaning of Life.
253 *Nanoethics* **3**, 257–268 (2009).

254 27. Saha, S. *et al.* Microbial Symbiosis: A Network towards Biomethanation. *Trends Microbiol.* **28**, 968–984
255 (2020).

256 28. Jacoby, R., Peukert, M., Succurro, A., Koprivova, A. & Kopriva, S. The Role of Soil Microorganisms in Plant
257 Mineral Nutrition—Current Knowledge and Future Directions. *Front. Plant Sci.* **8**, (2017).

258 29. Krzywoszynska, A. Nonhuman Labor and the Making of Resources Making Soils a Resource through Microbial
259 Labor. *Environ. Humanit.* **12**, 227–249 (2020).

