

DOI: https://doi.org/10.1093/synbio/ysad011 Advance access publication date: 27 May 2023 Synthetic Biology News

An artificial protein translation language makes bacteria resistant to viruses

David M. Truong^{[]1,2,*}

¹Department of Biomedical Engineering, NYU Tandon School of Engineering, Brooklyn, NY, USA ²Department of Pathology, NYU Langone Health, New York, NY, USA

* Corresponding author: E-mail: David.Truong@nyu.edu

American statesman Benjamin Franklin once stated, '... in this world nothing can be said to be certain, except death and taxes'. Had he been a microbiologist, he would have known about another near certainty: that viruses and the organisms they infect, will continue waging cycles of genetic conflict where the host builds immunity against a virus and the virus finds a way around it. But what if we had a way to break that cycle?

In a recent paper by Nyerges *et al.* from George Church's lab at Harvard Medical School in the USA, the authors engineered the bacterium *Escherichia coli* to be resistant to bacteriophages (bacterial viruses) by using the phage's reliance on host machinery against them—viruses rely almost entirely on their host's protein machinery to reproduce, as they cannot replicate on their own (1). To achieve this, they generated a new artificial genetic 'language' by which messenger RNAs (mRNAs) are read and translated into proteins. Importantly, the *E. coli* host neither needs nor uses this artificial language to make proteins, but crucially, the bacteriophage does. The system might not only prohibit phage infections but also enable 'biocontainment' of the engineered bacteria to prevent escape into the wild.

The genetic code transcribed into mRNAs, which code for proteins, is divided into a three-nucleotide language called codons, which is universal to life as well as viruses. Codons are interpreted by molecules called transfer RNAs (tRNAs), which are bound to specific amino acids. There is redundancy in this language, as there are only 20 amino acids, but 64 possible codons. Because some amino acids have multiple codon/tRNA options, the field of synthetic genomics has long sought to repurpose excess codons for new designer functions, such as inputting non-natural amino acids into proteins (2).

Synthetic genomics practitioners can generate organisms with new designer genomes using advanced DNA synthesis and genetic engineering, and it was a long-standing question whether phage infections could be prevented by recoding entire genomes. Indeed, prior work by the laboratory of Jason Chin at the Medical Research Council Laboratory of Molecular Biology in Cambridge, UK, fully recoded the *E. coli* genome to make the strain Syn61 Δ 3, in which they genome-wide repurposed two codons specifying serine and one stop sequence (3). In doing so, invading viruses now improperly make proteins, and the same team later concluded that Syn61 Δ 3 was resistant to viruses (4). The current study by Nyerges *et al.* found instead that Syn61 Δ 3 was only resistant to some bacteriophages but was susceptible to many other types.

It turns out that many viruses also carry their own tRNAs and thus come ready with the old language when infecting Syn61 Δ 3. To overcome this loophole, Nyerges focused on changing the tRNAs themselves. They hacked the viruses' own tRNA-serine pair to now be a tRNA-leucine pair. When they placed this new tRNA into Syn61 Δ 3 (which no longer has the codons this tRNA recognizes), it caused invading bacteriophages (which do use it) to improperly incorporate leucine instead of serine into their proteins, like typing on a Greek keyboard thinking it was an English one. Using another tRNA, they also addicted Syn61 Δ 3 to a non-natural amino acid called BipA, thereby preventing its growth in the wild without human help. Concurrent to this, Chin's group demonstrated a similar strategy, but they instead used weaker *E. coli* tRNAs (5), which the Nyerges study suggests viruses could overcome.

Host organisms like *E.* coli are routinely used for producing biologics and other industrial products. Bacteriophage infections are a common problem in biomanufacturing (6). Genomically recoded organisms like these might prevent virus contamination, saving millions of dollars lost each year. The work also serves as a basis for greater challenges like the Genome-Project Write Consortiums' goal to make virus-resistant mammalian cell lines (7). If successful, perhaps one day we might be rid of our unwanted relationship with viruses once and for all.

Conflict of interest statement. None declared.

References

 Nyerges, A., Vinke, S., Flynn, R., Owen, S.V., Rand, E.A., Budnik, B., Keen, E., Narasimhan, K., Marchand, J.A., Baas-Thomas, M. *et al.* (2023) A swapped genetic code prevents viral infections and gene transfer. Nature, **615**, 720–727.

Submitted: 22 May 2023; Received: 22 May 2023; Accepted: 26 May 2023

© The Author(s) 2023. Published by Oxford University Press.

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (https://creativecommons.org/licenses/by/4.0/), which permits unrestricted reuse, distribution, and reproduction in any medium, provided the original work is properly cited.

- 2. Gesteland, R.F., Weiss, R.B. and Atkins, J.F. (1992) Recoding: reprogrammed genetic decoding. *Science*, **257**, 1640–1641.
- 3. Fredens, J., Wang, K., de la Torre, D., Funke, L.F.H., Robertson, W.E., Christova, Y., Chia, T., Schmied, W.H., Dunkelmann, D.L., Beranek, V. et al. (2019) Total synthesis of *Escherichia coli* with a recoded genome. *Nature*, **569**, 514–518.
- 4. Robertson, W.E., Funke, L.F.H., de la Torre, D., Fredens, J., Elliott, T.S., Spinck, M., Christova, Y., Cervettini, D., Boge, F.L., Liu, K.C. et al. (2021) Sense codon reassignment enables viral resistance and encoded polymer synthesis. *Science*, **372**, 1057–1062.
- 5. Zurcher, J.F., Robertson, W.E., Kappes, T., Petris, G., Elliott, T.S., Salmond, G.P.C. and Chin, J.W. (2022) Refactored genetic codes enable bidirectional genetic isolation. *Science*, **378**, 516–523.
- 6. Barone, P.W., Wiebe, M.E., Leung, J.C., Hussein, I.T.M., Keumurian, F.J., Bouressa, J., Brussel, A., Chen, D., Chong, M., Dehghani, H. et al. (2020) Viral contamination in biologic manufacture and implications for emerging therapies. Nat. Biotechnol., 38, 563–572.
- 7. Dolgin, E. (2018) Scientists downsize bold plan to make human genome from scratch. *Nature*, **557**, 16–17.