

# Advanced Applications Synthetic Biology

Suvin Sundararajan  
Bio Bootcamp III  
January 23, 2026

*Scanning electron micrograph images of cellularly engineered HUVECs*  
Harvard University • May 1



HOW TO GROW (ALMOST) ANYTHING

# Agenda

# Agenda

Part 1: Developments in the Field

(10 minutes)

# Agenda

**Part 1: Developments in the Field** (10 minutes)

**Part 2: Role of Automation and AI** (10 minutes)

# Agenda

- |  |              |
|--|--------------|
| <b>Part 1:</b> Developments in the Field | (10 minutes) |
| <b>Part 2:</b> Role of Automation and AI | (10 minutes) |
| <b>Part 3:</b> Live Demonstrations       | (15 min)     |

# Agenda

- |  |              |
|--|--------------|
| <b>Part 1:</b> Developments in the Field | (10 minutes) |
| <b>Part 2:</b> Role of Automation and AI | (10 minutes) |
| <b>Part 3:</b> Live Demonstrations       | (15 min)     |
| ○ <i>RF Diffusion, AlphaFold</i>         |              |

# Agenda

**Part 1:** Developments in the Field (10 minutes)

**Part 2:** Role of Automation and AI (10 minutes)

**Part 3:** Live Demonstrations (15 min)

- *RF Diffusion, AlphaFold*
- *Lab Automation (Time Pending)*

# Agenda

**Part 1:** Developments in the Field (10 minutes)

**Part 2:** Role of Automation and AI (10 minutes)

**Part 3:** Live Demonstrations (15 min)

- *RF Diffusion, AlphaFold*
- *Lab Automation (Time Pending)*

**Part 4:** Case Study (15 min)

# Part 1: Developments in the Field

**How can we improve experimental discovery to advance society and improve wellbeing?**

# 1. Protein Prediction and Analysis Milestones

# 1. Protein Prediction and Analysis Milestones

- **2021: Structure prediction becomes usable (AlphaFold2)**
  - Accurate protein structures at scale shift biology to model-informed decisions.

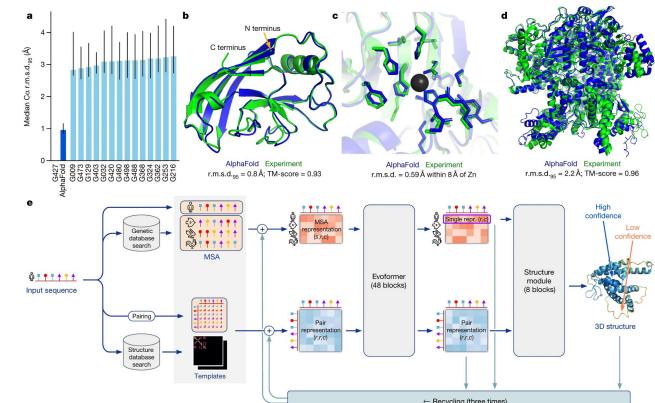
Article | [Open access](#) | Published: 15 July 2021

## Highly accurate protein structure prediction with AlphaFold

John Jumper , Richard Evans, Alexander Pritzel, Tim Green, Michael Figurnov, Olaf Ronneberger, Kathryn Tunyasuvunakool, Russ Bates, Augustin Žídek, Anna Potapenko, Alex Bridgland, Clemens Meyer, Simon A. A. Kohl, Andrew J. Ballard, Andrew Cowie, Bernardino Romera-Paredes, Stanislav Nikolov, Rishabh Jain, Jonas Adler, Trevor Back, Stig Petersen, David Reiman, Ellen Clancy, Michal Zeliński, ... Demis Hassabis  + Show authors

*Nature* **596**, 583–589 (2021) | [Cite this article](#)

1.96M Accesses | 20k Citations | 4010 Altmetric | [Metrics](#)



# 1. Protein Prediction and Analysis Milestones

- **2021: Structure prediction becomes usable (AlphaFold2)**
  - Accurate protein structures at scale shift biology to model-informed decisions.
- **2022: Protein language models scale the “sequence → insight” loop (ESMFold / ESM-2)**
  - Faster, cheaper inference and massive sequence-space coverage enable large metagenomic structure atlases and numerous hypotheses.

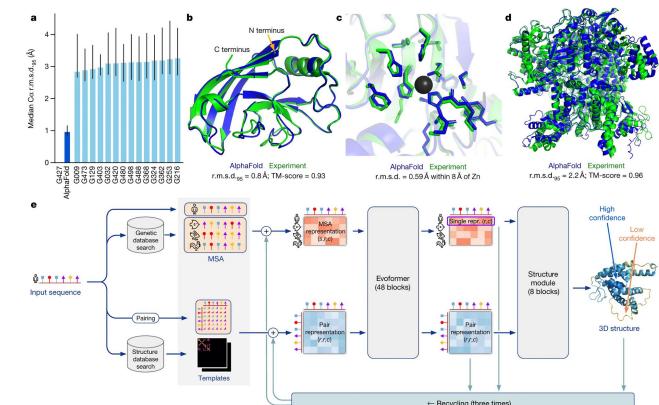
Article | [Open access](#) | Published: 15 July 2021

## Highly accurate protein structure prediction with AlphaFold

John Jumper , Richard Evans, Alexander Pritzel, Tim Green, Michael Figurnov, Olaf Ronneberger, Kathryn Tunyasuvunakool, Russ Bates, Augustin Žídek, Anna Potapenko, Alex Bridgland, Clemens Meyer, Simon A. Kohl, Andrew J. Ballard, Andrew Cowie, Bernardino Romera-Paredes, Stanislav Nikolov, Rishabh Jain, Jonas Adler, Trevor Back, Stig Petersen, David Reiman, Ellen Clancy, Michał Zieliński, ... Demis Hassabis  + Show authors

*Nature* **596**, 583–589 (2021) | [Cite this article](#)

1.96M Accesses | 20k Citations | 4010 Altmetric | [Metrics](#)



# 1. Protein Prediction and Analysis Milestones

- **2021: Structure prediction becomes usable (AlphaFold2)**
  - Accurate protein structures at scale shift biology to model-informed decisions.
- **2022: Protein language models scale the “sequence → insight” loop (ESMFold / ESM-2)**
  - Faster, cheaper inference and massive sequence-space coverage enable large metagenomic structure atlases and numerous hypotheses.
- **2023: Generative design arrives (RFdiffusion)**
  - Models begin proposing new sequences/molecules, not just predicting nature.

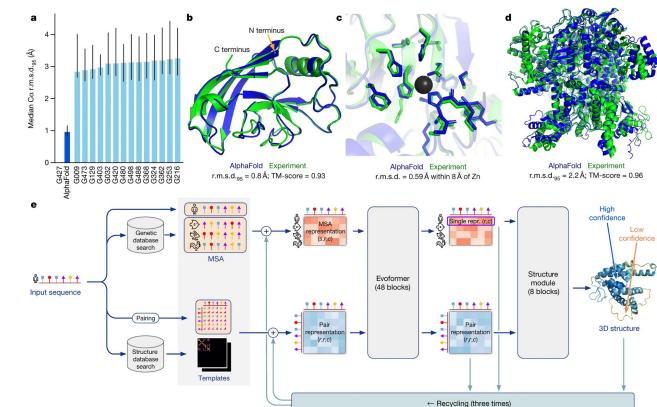
Article | [Open access](#) | Published: 15 July 2021

## Highly accurate protein structure prediction with AlphaFold

John Jumper , Richard Evans, Alexander Pritzel, Tim Green, Michael Figurnov, Olaf Ronneberger, Kathryn Tunyasuvunakool, Russ Bates, Augustin Žídek, Anna Potapenko, Alex Bridgland, Clemens Meyer, Simon A. A. Kohl, Andrew J. Ballard, Andrew Cowie, Bernardino Romera-Paredes, Stanislav Nikolov, Rishabh Jain, Jonas Adler, Trevor Back, Stig Petersen, David Reiman, Ellen Clancy, Michał Zieliński, ... Demis Hassabis  + Show authors

*Nature* **596**, 583–589 (2021) | [Cite this article](#)

1.96M Accesses | 20k Citations | 4010 Altmetric | [Metrics](#)



# 1. Protein Prediction and Analysis Milestones

- **2021: Structure prediction becomes usable (AlphaFold2)**
  - Accurate protein structures at scale shift biology to model-informed decisions.
- **2022: Protein language models scale the “sequence → insight” loop (ESMFold / ESM-2)**
  - Faster, cheaper inference and massive sequence-space coverage enable large metagenomic structure atlases and numerous hypotheses.
- **2023: Generative design arrives (RFdiffusion)**
  - Models begin proposing new sequences/molecules, not just predicting nature.
- **2024: Interaction-centric modeling (AlphaFold 3)**
  - Focus shifts from "folded shape" to binding interactions (complexes, ligands, RNA/DNA).

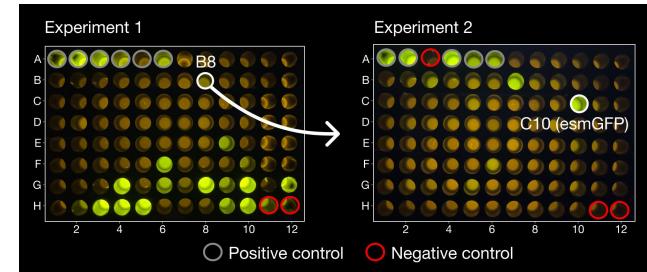
Article | [Open access](#) | Published: 15 July 2021

## Highly accurate protein structure prediction with AlphaFold

John Jumper , Richard Evans, Alexander Pritzel, Tim Green, Michael Figurnov, Olaf Ronneberger, Kathryn Tunyasuvunakool, Russ Bates, Augustin Žídek, Anna Potapenko, Alex Bridgland, Clemens Meyer, Simon A. A. Kohl, Andrew J. Ballard, Andrew Cowie, Bernardino Romera-Paredes, Stanislav Nikolov, Rishabh Jain, Jonas Adler, Trevor Back, Stig Petersen, David Reiman, Ellen Clancy, Michał Zieliński, ... Demis Hassabis  + Show authors

*Nature* **596**, 583–589 (2021) | [Cite this article](#)

1.96M Accesses | 20k Citations | 4010 Altmetric | [Metrics](#)



# 1. Protein Prediction and Analysis Milestones

- **2021: Structure prediction becomes usable (AlphaFold2)**
  - Accurate protein structures at scale shift biology to model-informed decisions.
- **2022: Protein language models scale the “sequence → insight” loop (ESMFold / ESM-2)**
  - Faster, cheaper inference and massive sequence-space coverage enable large metagenomic structure atlases and numerous hypotheses.
- **2023: Generative design arrives (RFdiffusion)**
  - Models begin proposing new sequences/molecules, not just predicting nature.
- **2024: Interaction-centric modeling (AlphaFold 3)**
  - Focus shifts from "folded shape" to binding interactions (complexes, ligands, RNA/DNA).
- **2025: Truly novel functional proteins and target-conditioned design emerge (ESM3 → esmGFP)**
  - Foundation/diffusion models generate novel, functional proteins, making target-conditioned design practical and moving protein engineering toward an iterative workflow.

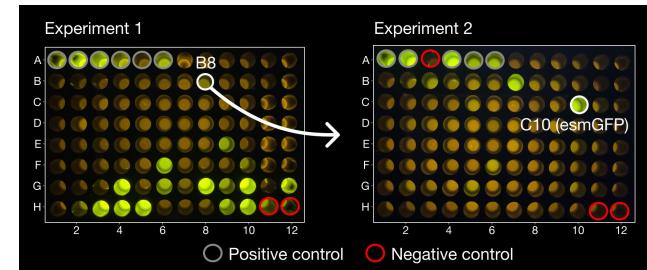
Article | [Open access](#) | Published: 15 July 2021

## Highly accurate protein structure prediction with AlphaFold

John Jumper , Richard Evans, Alexander Pritzel, Tim Green, Michael Figurnov, Olaf Ronneberger, Kathryn Tunyasuvunakool, Russ Bates, Augustin Žídek, Anna Potapenko, Alex Bridgland, Clemens Meyer, Simon A. A. Kohl, Andrew J. Ballard, Andrew Cowie, Bernardino Romera-Paredes, Stanislav Nikolov, Rishabh Jain, Jonas Adler, Trevor Back, Stig Petersen, David Reiman, Ellen Clancy, Michał Zieliński, ... Demis Hassabis  + Show authors

*Nature* **596**, 583–589 (2021) | [Cite this article](#)

1.96M Accesses | 20k Citations | 4010 Altmetric | [Metrics](#)



## 2. GLP-1 (Ozempic) Launch

## 2. GLP-1 (Ozempic) Launch



## 2. GLP-1 (Ozempic) Launch



### What Is Ozempic and Why Is It Getting So Much Attention?

More people are turning to a diabetes medication to induce weight loss — but experts say it's not a miracle drug.

[Share full article](#) 775



Getty Images

## 2. GLP-1 (Ozempic) Launch



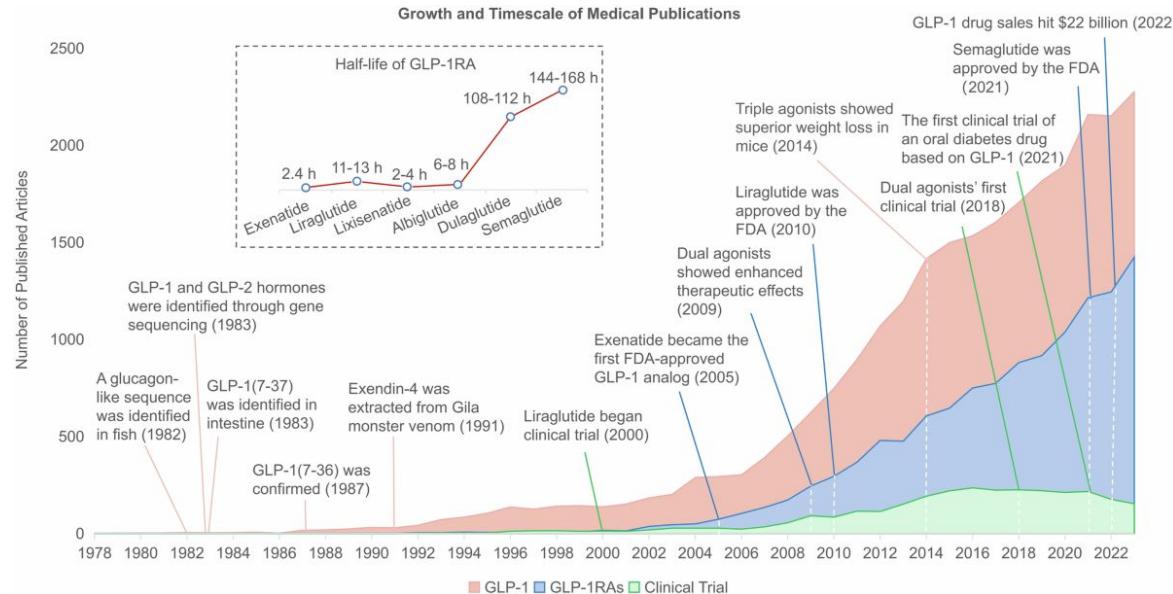
### What Is Ozempic and Why Is It Getting So Much Attention?

More people are turning to a diabetes medication to induce weight loss — but experts say it's not a miracle drug.

[Share full article](#) 775



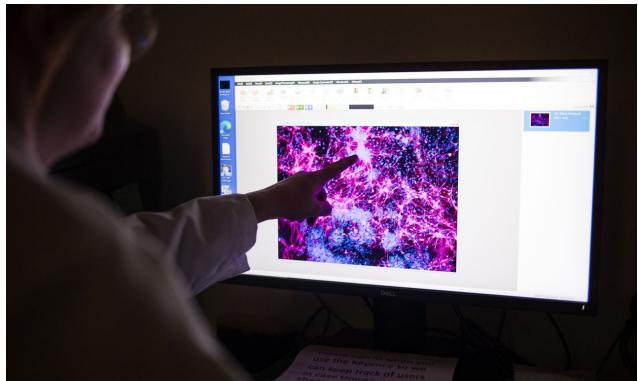
Getty Images



### 3. Personalized CRISPR Therapy

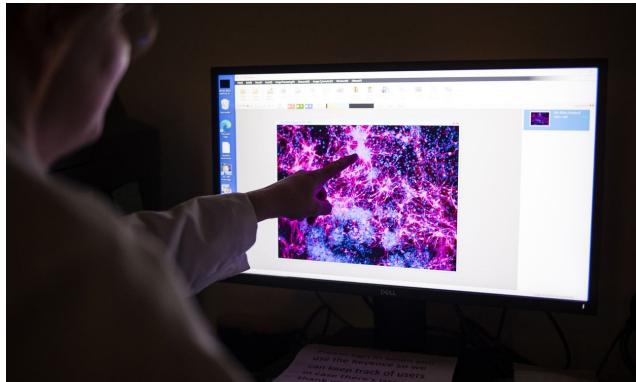
### 3. Personalized CRISPR Therapy

- **Traditional Model**
  - One editor applied to one disease program over many years, treating many patients.



### 3. Personalized CRISPR Therapy

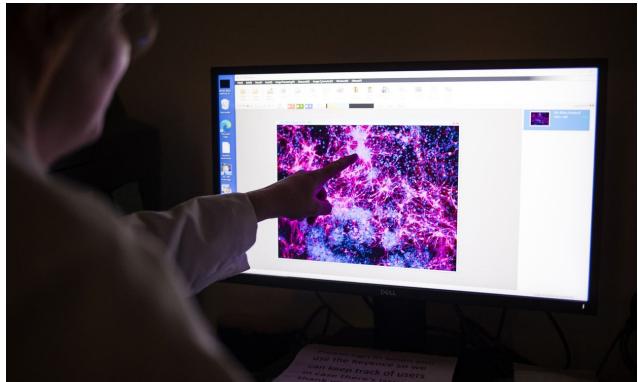
- **Traditional Model**
  - One editor applied to one disease program over many years, treating many patients.
- **New Frontier (N-of-1)**
  - Patient-specific editing for ultra-rare mutations (e.g., bespoke therapy for a child by CHOP/Penn).



<https://www.chop.edu/news/worlds-first-patient-treated-personalized-crispr-gene-editing-therapy-childrens-hospital>

### 3. Personalized CRISPR Therapy

- **Traditional Model**
  - One editor applied to one disease program over many years, treating many patients.
- **New Frontier (N-of-1)**
  - Patient-specific editing for ultra-rare mutations (e.g., bespoke therapy for a child by CHOP/Penn).
- **Shift in Value**
  - The "product" is now the capacity to rapidly generate safe, specific edits. Value is placed on speed and correctness, not just novelty.

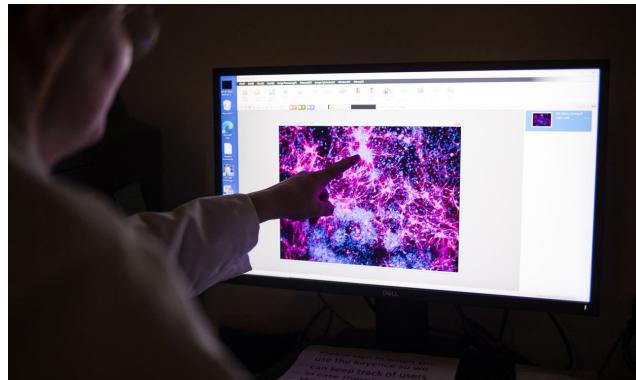


<https://www.chop.edu/news/worlds-first-patient-treated-personalized-crispr-gene-editing-therapy-childrens-hospital>

### 3. Personalized CRISPR Therapy

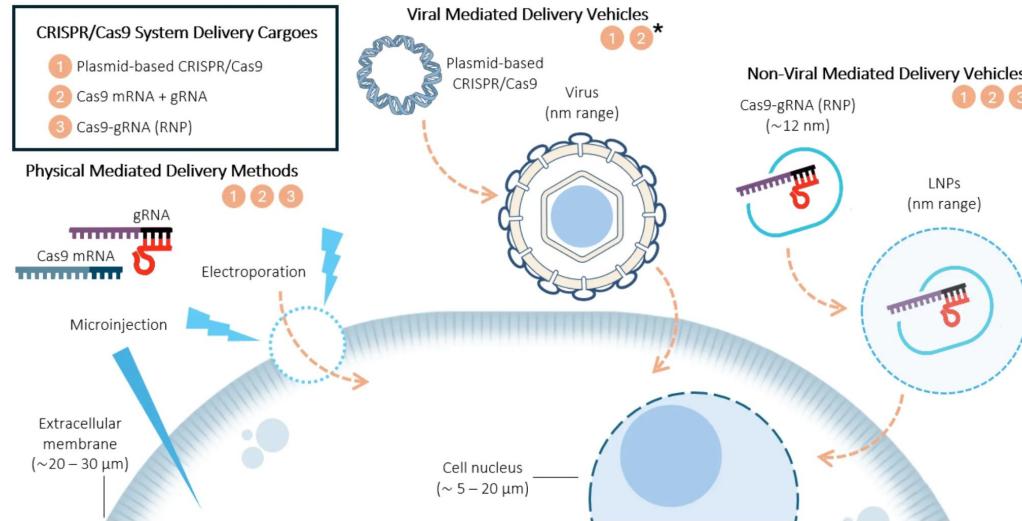
- **Traditional Model**
  - One editor applied to one disease program over many years, treating many patients.
- **New Frontier (N-of-1)**
  - Patient-specific editing for ultra-rare mutations (e.g., bespoke therapy for a child by CHOP/Penn).
- **Shift in Value**
  - The "product" is now the capacity to rapidly generate safe, specific edits. Value is placed on speed and correctness, not just novelty.

***Gene editing is becoming a more repeatable process, reframing medicine as an engineering discipline where biology is increasingly programmable.***



<https://www.chop.edu/news/worlds-first-patient-treated-personalized-crispr-gene-editing-therapy-childrens-hospital>

### 3. Personalized CRISPR Therapy - Challenges

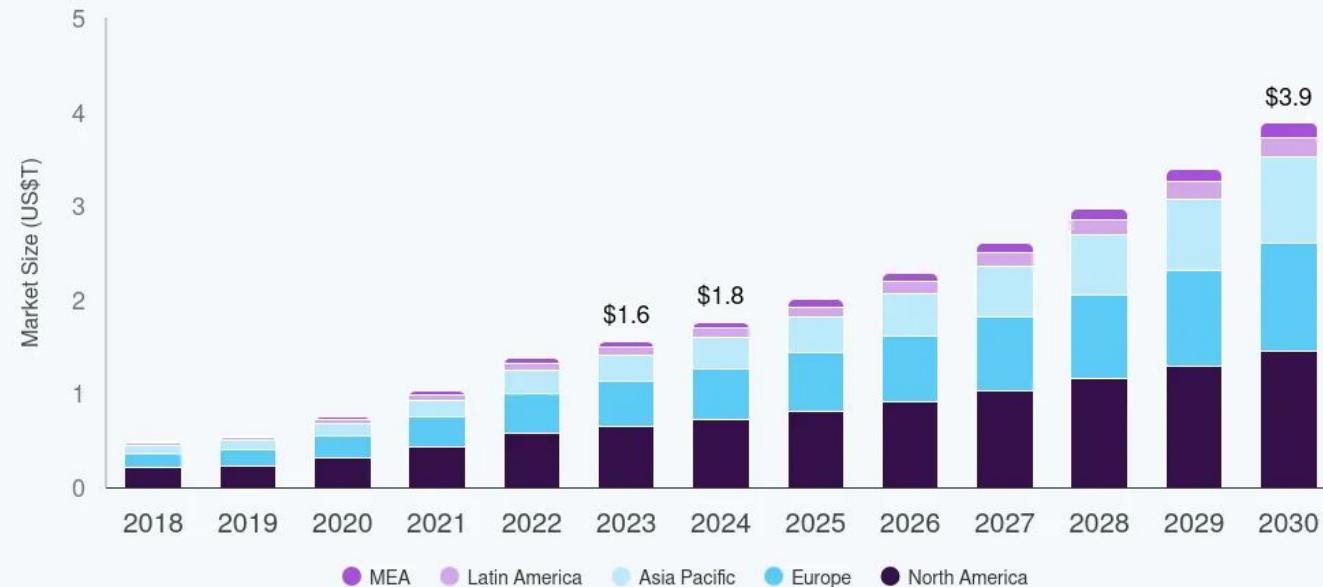


<https://www.synthego.com/blog/delivery-crispr-cas9/>

# 4. Market Growth

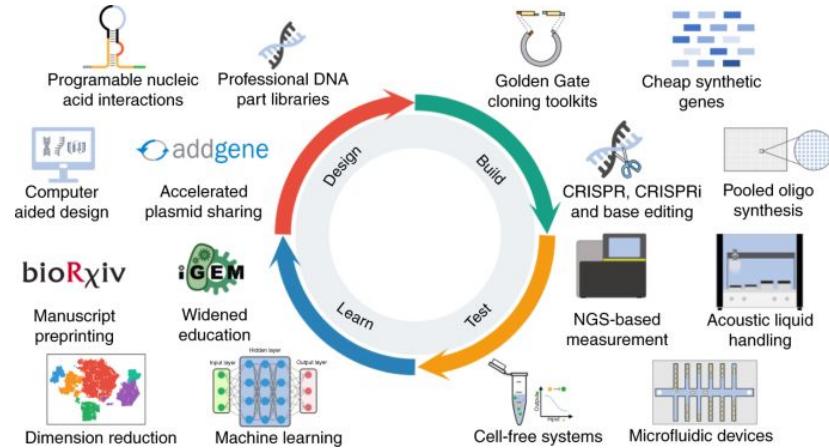
## Biotechnology Market

Size, by Region, 2018 - 2030



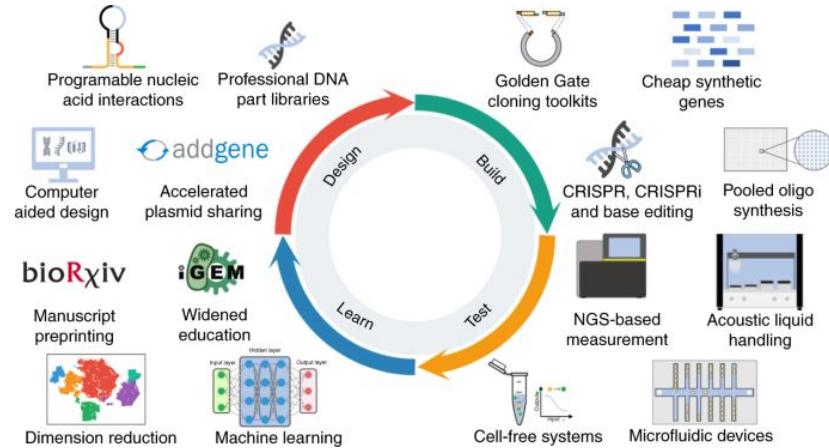
# Future Developments

- Biology is shifting from running single experiments to coordinated campaigns of experiments, Bayesian optimization, and active learning.



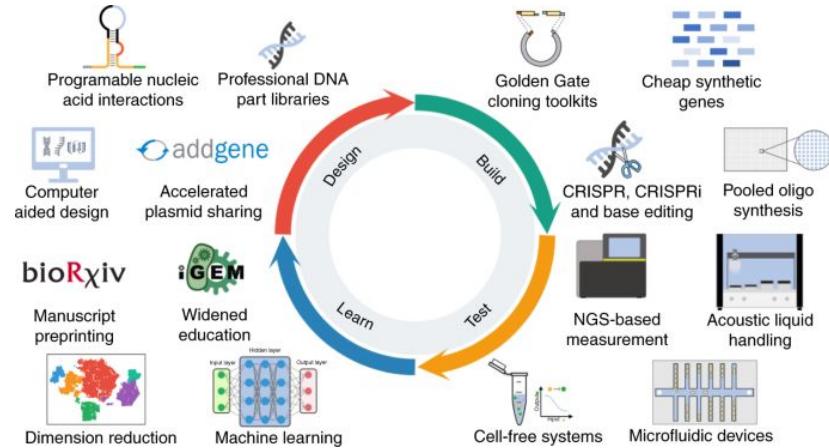
# Future Developments

- Biology is shifting from running single experiments to coordinated campaigns of experiments, Bayesian optimization, and active learning.
- **This shift is happening now because models are useful, lab robots are accessible, and assays are high-throughput, so iteration speed has become the limiting factor.**



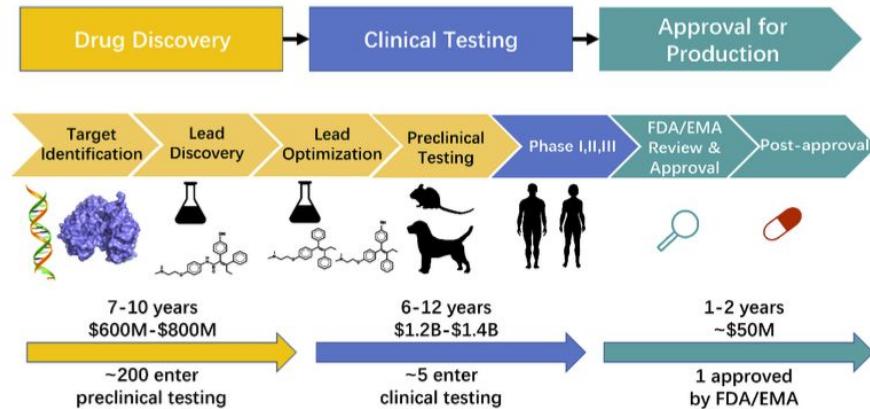
# Future Developments

- Biology is shifting from running single experiments to coordinated campaigns of experiments, Bayesian optimization, and active learning.
- This shift is happening now because models are useful, lab robots are accessible, and assays are high-throughput, so iteration speed has become the limiting factor.
- **The core workflow is a repeatable loop: design the next set of conditions, execute them, measure outcomes, learn from the data, and choose the next run—on a daily or weekly cadence rather than quarterly.**



# Future Developments

- Biology is shifting from running single experiments to coordinated campaigns of experiments, Bayesian optimization, and active learning.
- This shift is happening now because models are useful, lab robots are accessible, and assays are high-throughput, so iteration speed has become the limiting factor.
- **The core workflow is a repeatable loop: design the next set of conditions, execute them, measure outcomes, learn from the data, and choose the next run—on a daily or weekly cadence rather than quarterly.**



## Part 2: Role of Automation and AI

# What is the most common tool for synthetic biologists?

# What is the most common tool for synthetic biologists?



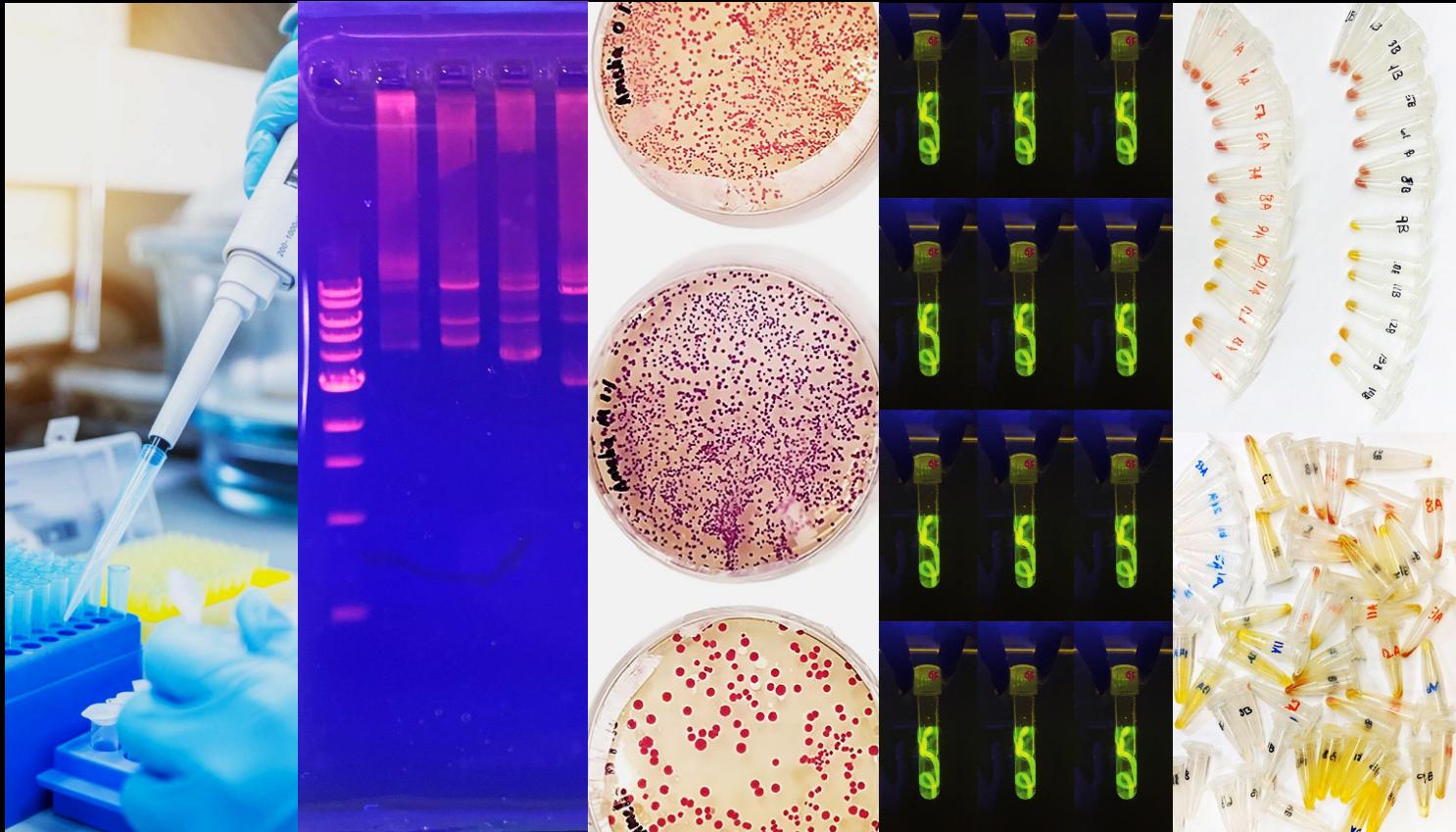
# How can we improve this?



=



# Sneak Peak



# HOW TO GROW (ALMOST) ANYTHING

# Sneak Peak

How to Grow (Almost) Anything 2025

Lab Protocol: Bioproduction of Beta-Carotene and Lycopene

## Protocol | Part 1: Overnight Cultures

Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- LB with pre-added antibiotic (chloramphenicol)
- Pipette set, serological pipettes and pipette tips
- Incubation room
- Culture tubes

You'll be testing two genes, two media combinations, one input, and two temperature conditions shown above. You'll need to set up **16 unique overnight cultures** in given culture tubes, alongside duplicates for each, and two cultures with just media (so a total of 34 cultures in total)

Condition #	Plasmid	Culture temp	Growth Medium
1 and 2	pAC-LYC	30°C, 37°C	Luria Broth (LB)
3 and 4	pAC-LYC	30°C, 37°C	LB + fructose
5 and 6	pAC-LYC	30°C, 37°C	2YT
7 and 8	pAC-LYC	30°C, 37°C	2YT + fructose
9 and 10	pAC-BETA	30°C, 37°C	LB
11 and 12	pAC-BETA	30°C, 37°C	LB + fructose
13 and 14	pAC-BETA	30°C, 37°C	2YT
15 and 16	pAC-BETA	30°C, 37°C	2YT + fructose

1. For each conditions above, set up overnight cultures that contain the following
  - a. 5 mL of the specified media (already supplemented with antibiotic)
  - b. 10 uL of *E. coli* from the starter culture with the specified plasmid
2. Grow cultures for 24 hours in the circular roller drum in the appropriate warm room

How to Grow (Almost) Anything 2025

Lab Protocol: The Chromophore Color Cloning Quest

Edited December 29, 2024

## Protocol | Part 1: PCR

Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- Ice bucket
- Phusion High-Fidelity PCR Master Mix
- Primers (working stocks, at 10 uM)
- UltraPure Water
- PCR tubes
- Thermocycler
- P20 pipette and 10uL tips
- P200 pipette and 200uL tips

### Setup the following PCR reactions:

Universal DNA Fragment (Univ Fwd and Univ Primers)				
Reaction	Reagent	Stock Conc.	Desired Conc.	Volume
All	Mini Prepped mUAV	Check Stock	Typically 10 ng	X uL
All	Univ Forward Primer	10 uM	0.5 uM	1.25 uL
All	Univ Reverse Primer	10 uM	0.5 uM	1.25 uL
All	Phusion HF PCR Mix	2X	1x	12.5 uL
All	Nuclease-free water	n/a	n/a	10 - X uL
All	Total Volume			25 uL

# Sneak Peak

How to Grow (Almost) Anything 2025  
Lab Protocol: Bioproduction of Beta-Carotene and Lycopene

## Protocol | Part 1: Overnight Cultures

Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- LB with pre-added antibiotic (chloramphenicol)
- Pipette set, serological pipettes and pipette tips
- Incubation room
- Culture tubes

You'll be testing two genes, two media combinations, one input, and two temperature conditions shown above. You'll need to set up **16 unique overnight cultures** in given culture tubes, alongside duplicates for each, and two cultures with just media (so a total of 34 cultures in total)

Condition #	Plasmid	Culture temp	Growth Medium
1 and 2	pAC-LYC	30°C, 37°C	Luria Broth (LB)
3 and 4	pAC-LYC	30°C, 37°C	LB + fructose
5 and 6	pAC-LYC	30°C, 37°C	2YT
7 and 8	pAC-LYC	30°C, 37°C	2YT + fructose
9 and 10	pAC-BETA	30°C, 37°C	LB
11 and 12	pAC-BETA	30°C, 37°C	LB + fructose
13 and 14	pAC-BETA	30°C, 37°C	2YT
15 and 16	pAC-BETA	30°C, 37°C	2YT + fructose

1. For each conditions above, set up overnight cultures that contain the following
  - a. 5 mL of the specified media (already supplemented with antibiotic)
  - b. 10 uL of *E. coli* from the starter culture with the specified plasmid
2. Grow cultures for 24 hours in the circular roller drum in the appropriate warm room

How to Grow (Almost) Anything 2025  
Lab Protocol: The Chromophore Color Cloning Quest

Edited December 29, 2024

## Protocol | Part 1: PCR

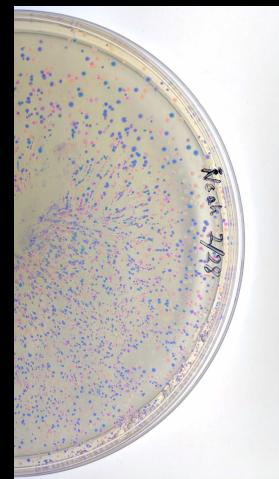
Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- Ice bucket
- Phusion High-Fidelity PCR Master Mix
- Primers (working stocks, at 10 uM)
- UltraPure Water
- PCR tubes
- Thermocycler
- P20 pipette and 10uL tips
- P200 pipette and 200uL tips

### Setup the following PCR reactions:

Universal DNA Fragment (Univ Fwd and Univ Primers)				
Reaction	Reagent	Stock Conc.	Desired Conc.	Volume
All	Mini Prepped mUAV	Check Stock	Typically 10 ng	X uL
All	Univ Forward Primer	10 uM	0.5 uM	1.25 uL
All	Univ Reverse Primer	10 uM	0.5 uM	1.25 uL
All	Phusion HF PCR Mix	2X	1x	12.5 uL
All	Nuclease-free water	n/a	n/a	10 - X uL
All	Total Volume			25 uL



# Sneak Peak

How to Grow (Almost) Anything 2025  
Lab Protocol: Bioproduction of Beta-Carotene and Lycopene

## Protocol | Part 1: Overnight Cultures

Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- LB with pre-added antibiotic (chloramphenicol)
- Pipette set, serological pipettes and pipette tips
- Incubation room
- Culture tubes

You'll be testing two genes, two media combinations, one input, and two temperature conditions shown above. You'll need to set up **16 unique overnight cultures** in given culture tubes, alongside duplicates for each, and two cultures with just media (so a total of 34 cultures in total)

Condition #	Plasmid	Culture temp	Growth Medium
1 and 2	pAC-LYC	30°C, 37°C	Luria Broth (LB)
3 and 4	pAC-LYC	30°C, 37°C	LB + fructose
5 and 6	pAC-LYC	30°C, 37°C	2YT
7 and 8	pAC-LYC	30°C, 37°C	2YT + fructose
9 and 10	pAC-BETA	30°C, 37°C	LB
11 and 12	pAC-BETA	30°C, 37°C	LB + fructose
13 and 14	pAC-BETA	30°C, 37°C	2YT
15 and 16	pAC-BETA	30°C, 37°C	2YT + fructose

1. For each conditions above, set up overnight cultures that contain the following
  - a. 5 mL of the specified media (already supplemented with antibiotic)
  - b. 10  $\mu$ L of *E. coli* from the starter culture with the specified plasmid
2. Grow cultures for 24 hours in the circular roller drum in the appropriate warm room

How to Grow (Almost) Anything 2025  
Lab Protocol: The Chromophore Color Cloning Quest

Edited December 29, 2024

## Protocol | Part 1: PCR

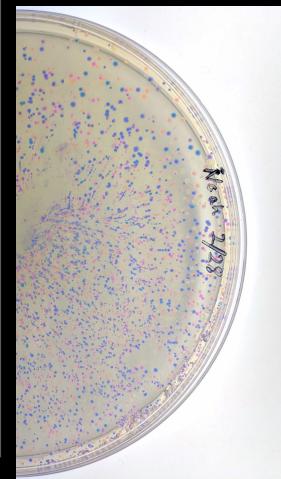
Time Estimate: 30 Minutes, 24 Hour Incubation

### Media, Equipment and Consumables

- Ice bucket
- Phusion High-Fidelity PCR Master Mix
- Primers (working stocks, at 10  $\mu$ M)
- UltraPure Water
- PCR tubes
- Thermocycler
- P20 pipette and 10 $\mu$ L tips
- P200 pipette and 200 $\mu$ L tips

### Setup the following PCR reactions:

Universal DNA Fragment (Univ Fwd and Univ Primers)				
Reaction	Reagent	Stock Conc.	Desired Conc.	Volume
All	Mini Prepped mUAV	Check Stock	Typically 10 ng	X $\mu$ L
All	Univ Forward Primer	10 $\mu$ M	0.5 $\mu$ M	1.25 $\mu$ L
All	Univ Reverse Primer	10 $\mu$ M	0.5 $\mu$ M	1.25 $\mu$ L
All	Phusion HF PCR Mix	2X	1X	12.5 $\mu$ L
All	Nuclease-free water	n/a	n/a	10 - X $\mu$ L
All	Total Volume			25 $\mu$ L



# Lab Work is Constrained by Physical Capacity



68-089 (HTGAA Lab Space)  ~15.5 ft lab bench

 ~5 ft space

Total Bench Width: ~101.5 ft

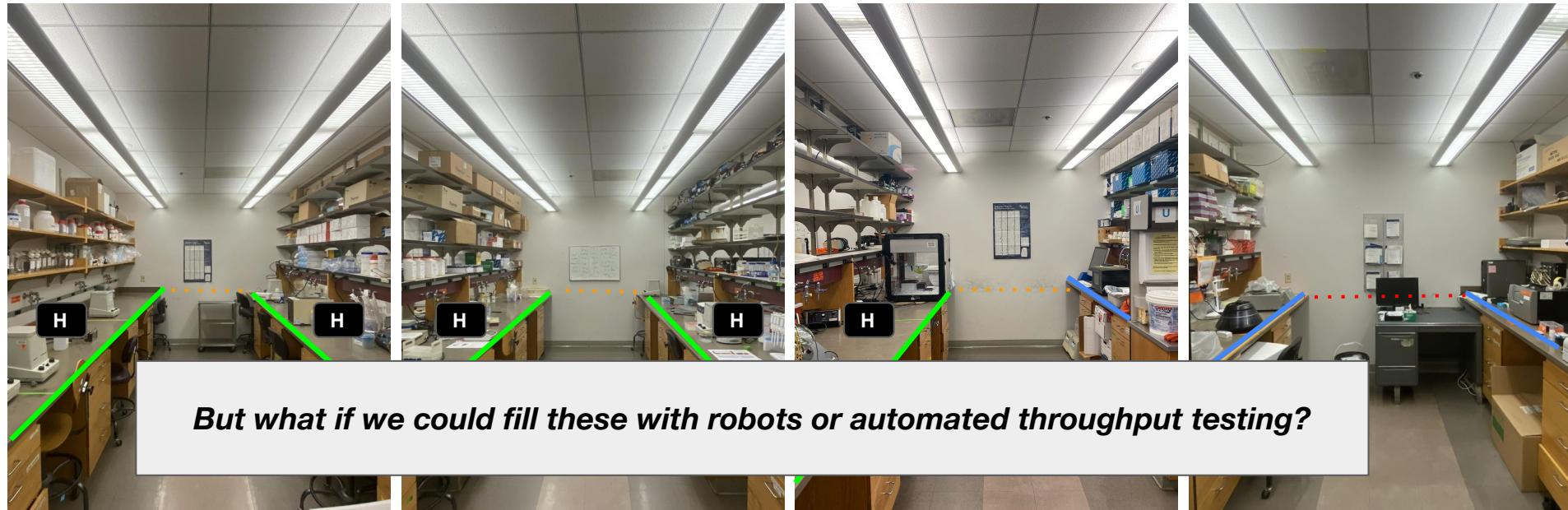
 ~8 ft lab bench

 ~5.5 ft space

Bench Depth: ~2.5 ft

NOTE: HTGAA Currently uses the first 5 lab benches in this room, denoted by 

# Lab Work is Constrained by Physical Capacity



***But what if we could fill these with robots or automated throughput testing?***

68-089 (HTGAA Lab Space)  ~15.5 ft lab bench

 ~5 ft space

Total Bench Width: ~101.5 ft

 ~8 ft lab bench

 ~5.5 ft space

Bench Depth: ~2.5 ft

NOTE: HTGAA Currently uses the first 5 lab benches in this room, denoted by 

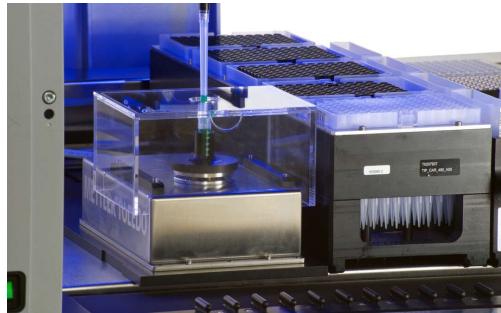
# Lab Modules



# Lab Modules



Opentrons Thermocycler module



Opentrons heater shaker



On-deck plate reader

# Lab Devices

>\$300k



Hamilton STAR

<\$10k



Opentrons OT-2\*

>\$300k

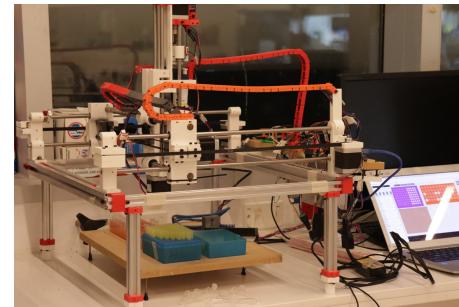


Tecan Freedom EVO

>\$300k



Hamilton Vantage



Open source robot

Many more...

# AI Scientist and Discovery

nature chemical engineering



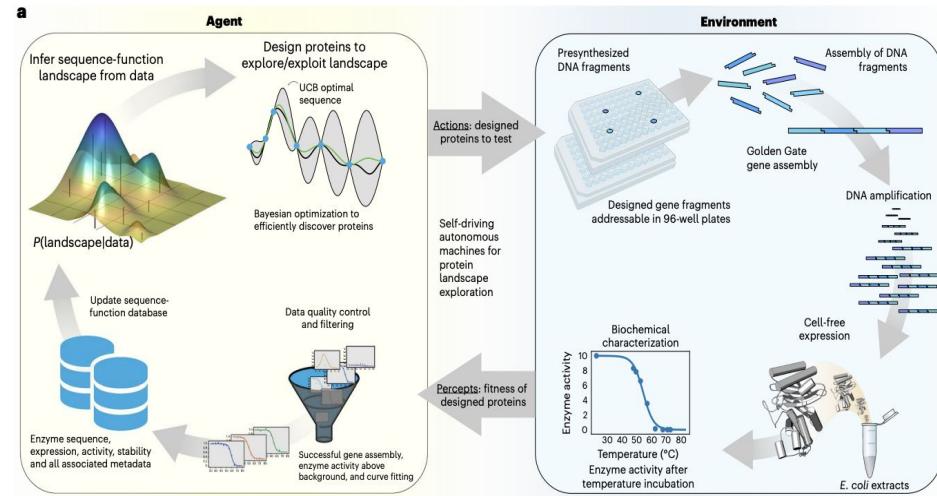
Article

<https://doi.org/10.1038/s44286-023-00002-4>

## Self-driving laboratories to autonomously navigate the protein fitness landscape

Received: 20 July 2023

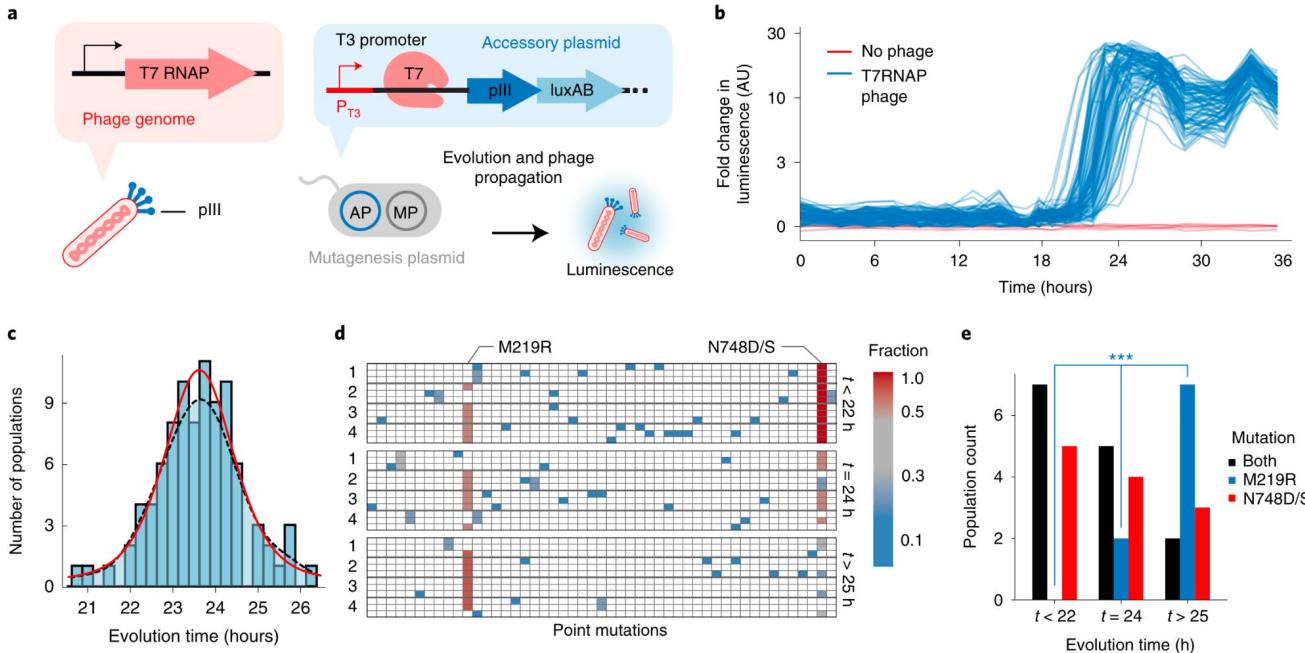
Jacob T. Rapp<sup>1</sup>, Bennett J. Bremer<sup>1</sup> & Philip A. Romero<sup>1,2</sup>



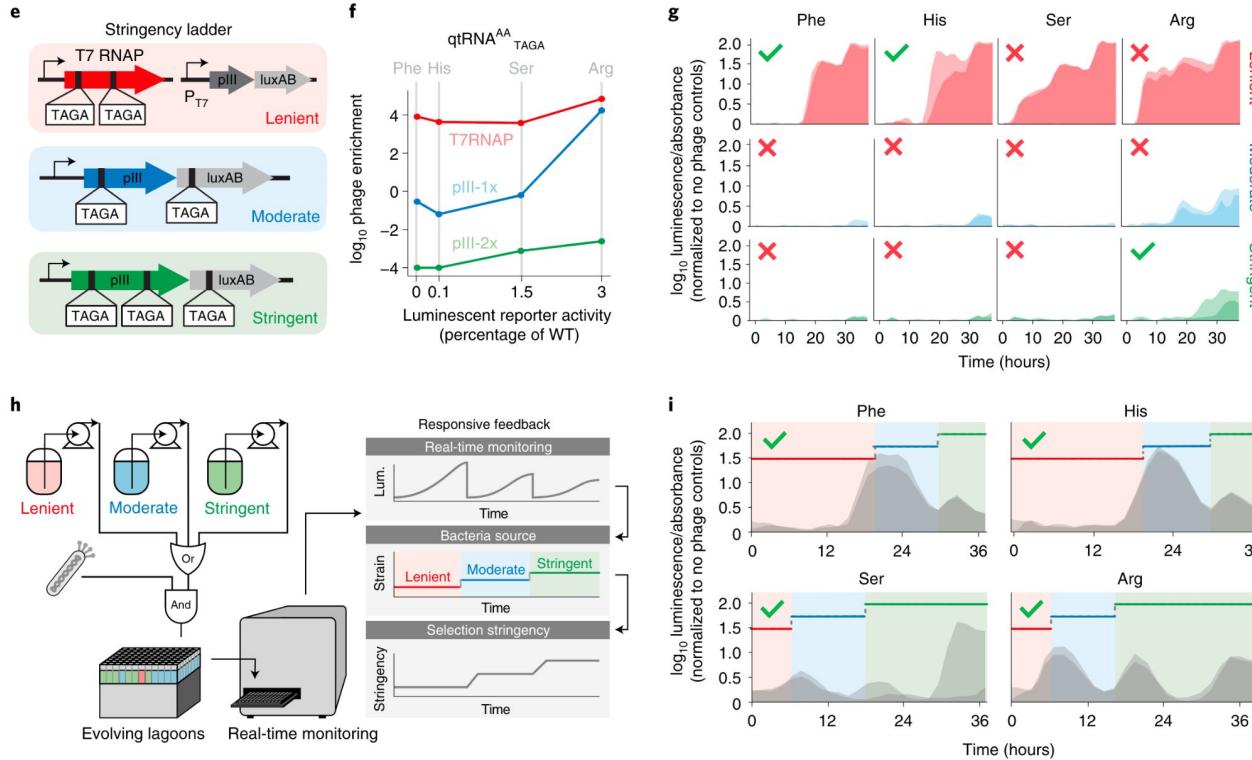
# Automation-enabled real-time monitoring and feedback

## Example: Continuous phage development

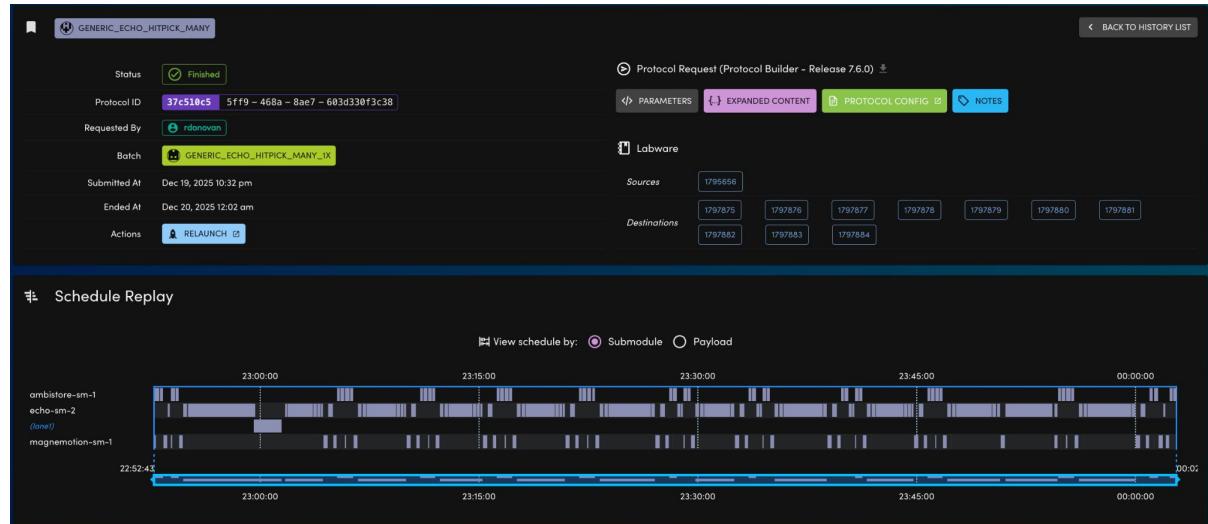
Can be used to automate conditions and DNA constructs for expression



# Automation-enabled real-time monitoring and feedback



# Automation-enabled real-time monitoring and feedback



Esther, Suvin, Ronan conducting fluorescent artwork preparation with a laboratory scheduler

## Part 3: Demonstrations

## Demo 1: RF Diffusion, AlphaFold

## Demo 2: Lab Automation

## Part 4: Case Study