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# Accelerated Engineering of Non-model Microbes

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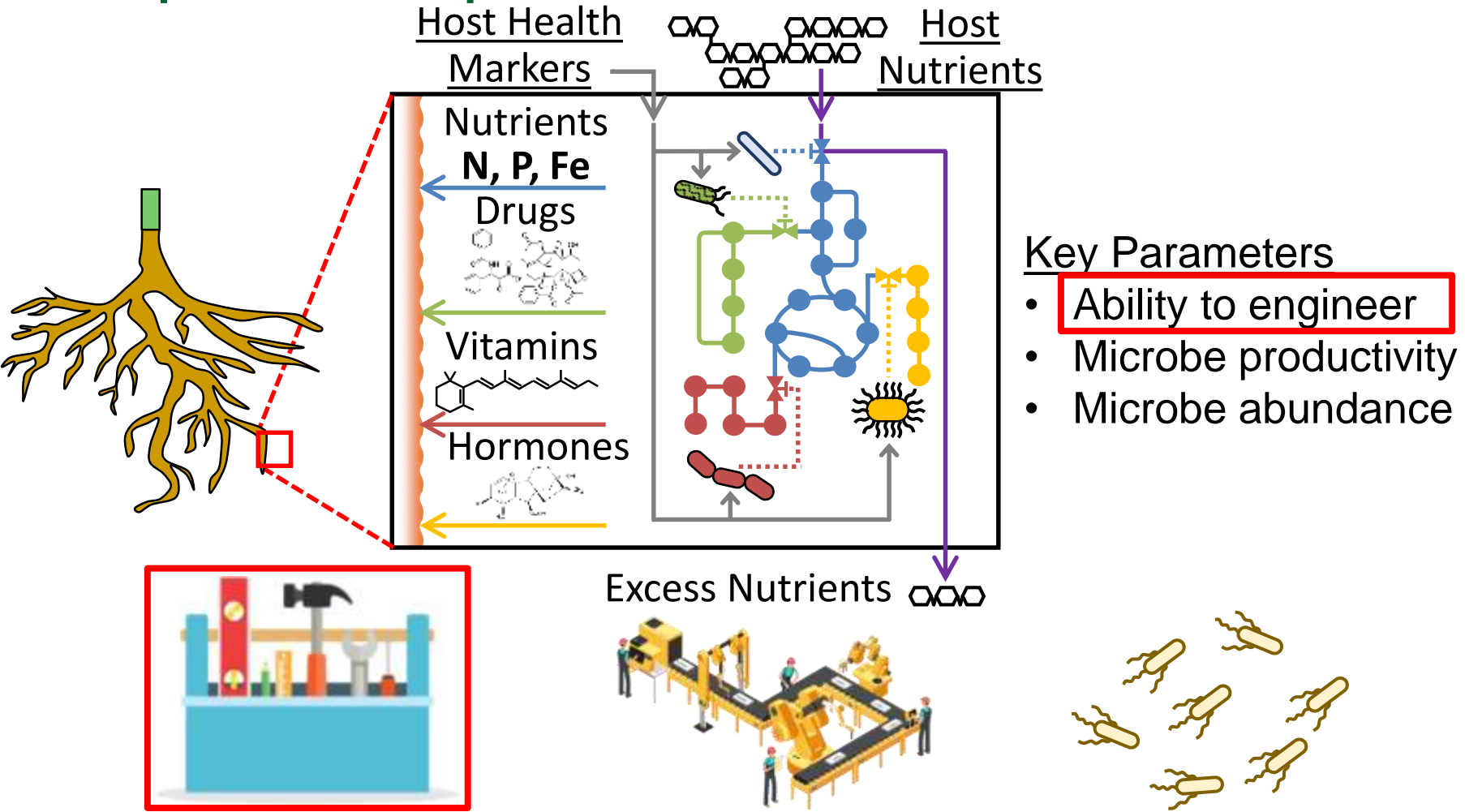
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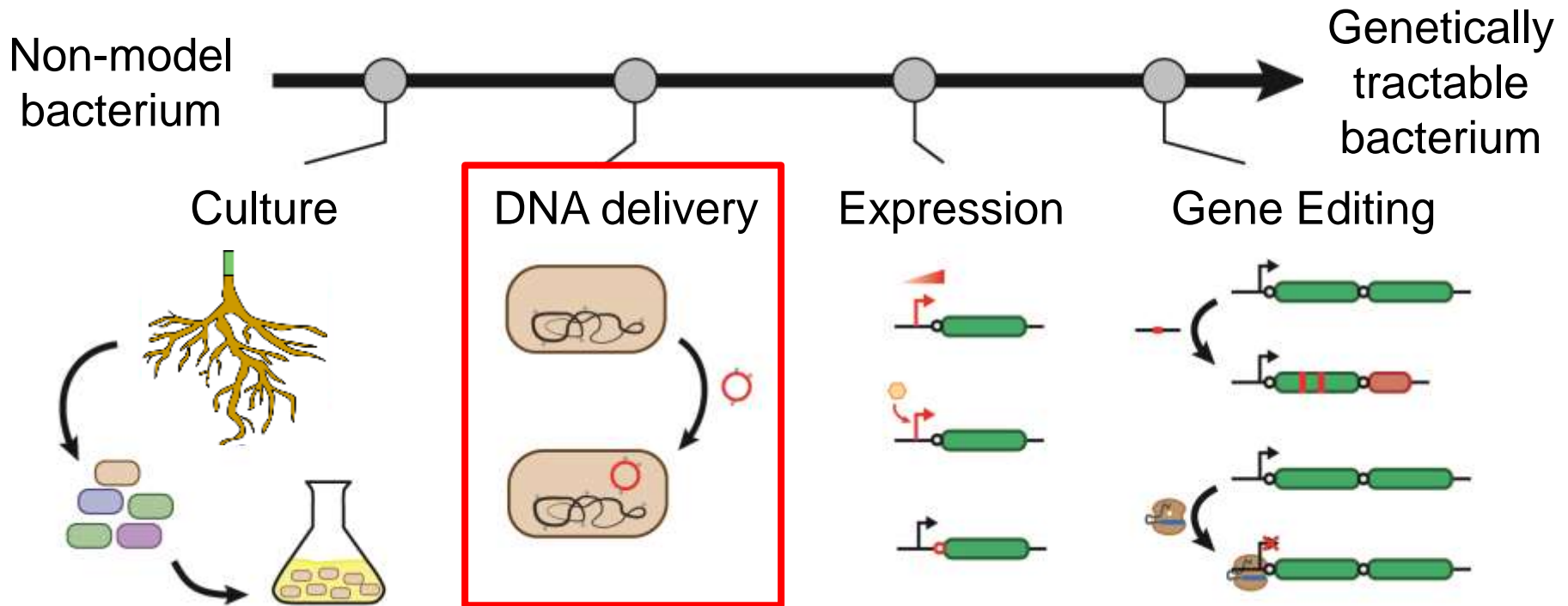
[www.crooklab.net](http://www.crooklab.net)

# Engineering the microbiota to improve crop health



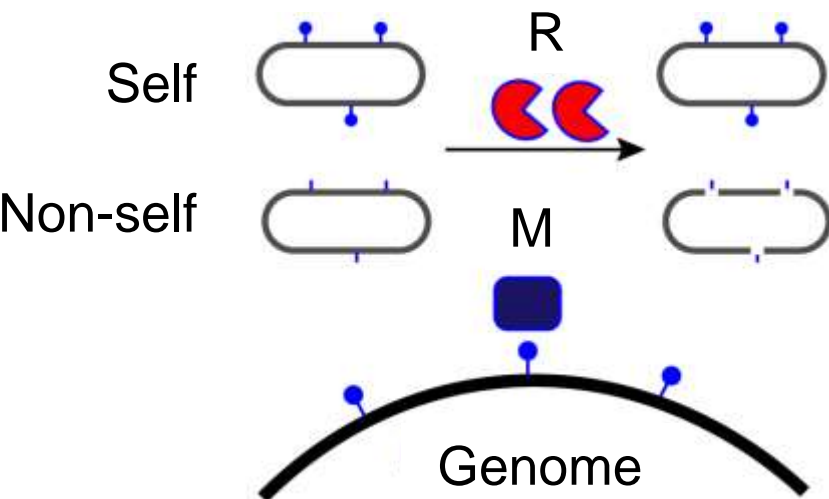
- Key Parameters**
- Ability to engineer
  - Microbe productivity
  - Microbe abundance

# Microbial Engineering Pipeline



# Need to quickly and efficiently overcome barriers to foreign DNA

## Restriction-Modification defense systems



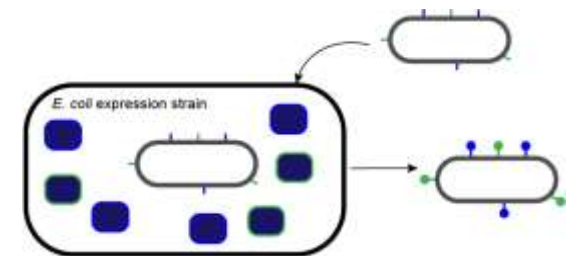
- Multiple MTases per bacterium
- MTases vary within species

## Standard approaches too slow and limited

Purified M proteins

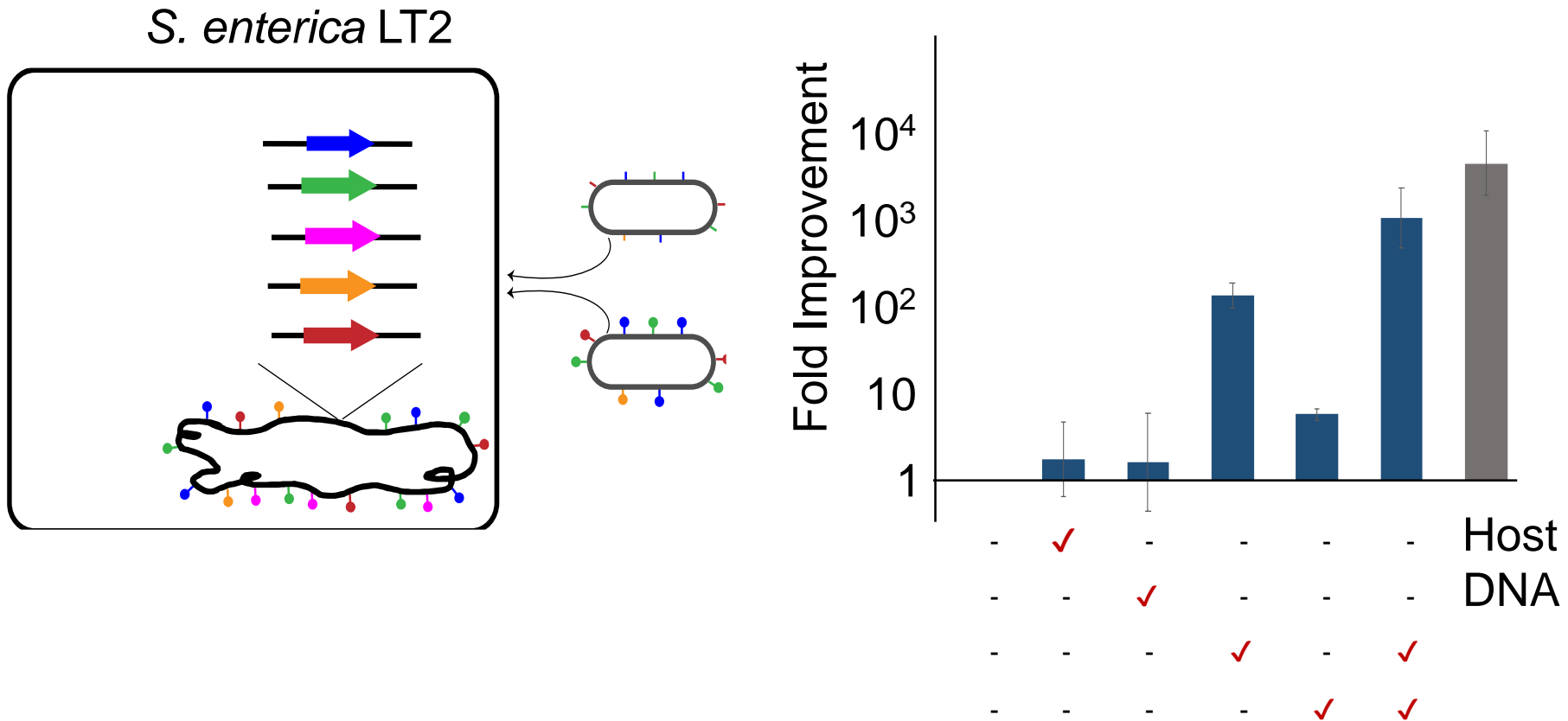


*E. coli* expressing M proteins

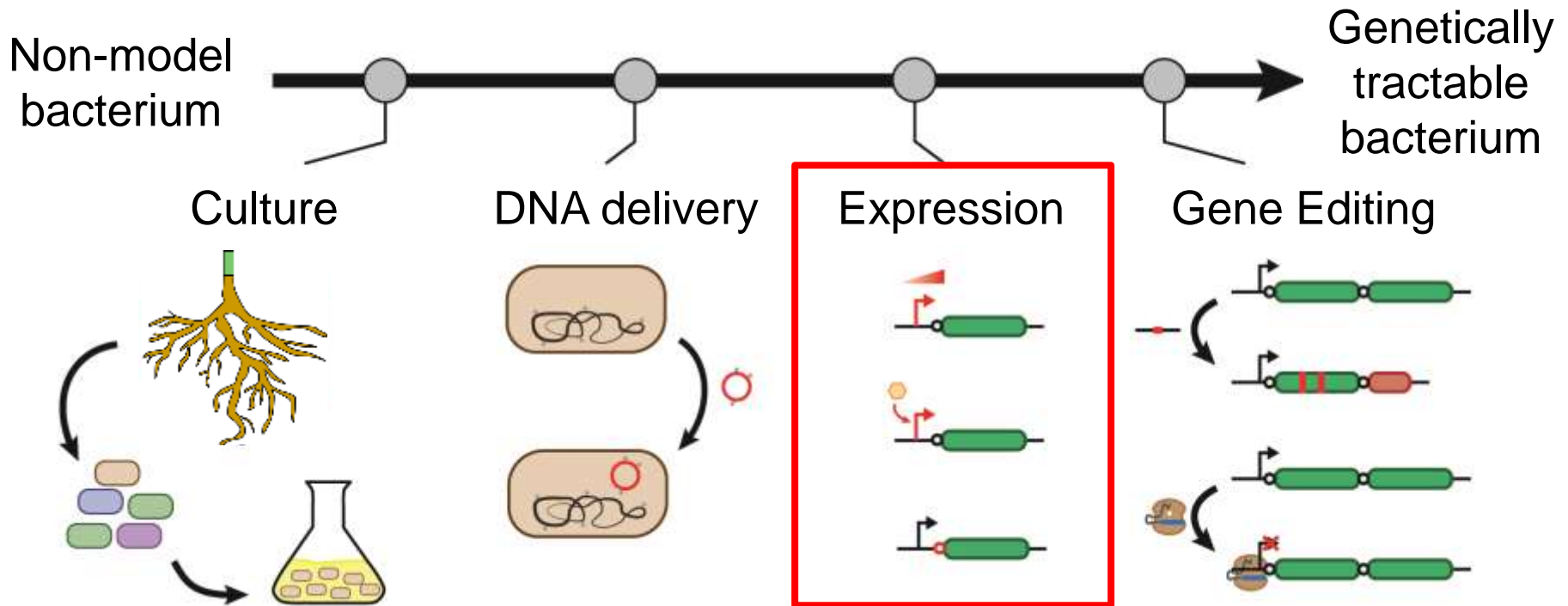


- M expression can be toxic
- Limited to 1 – 2 simple M proteins

# Enhanced plasmid transformation in *S. enterica*

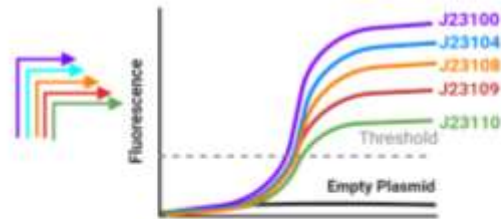


# Microbial Engineering Pipeline

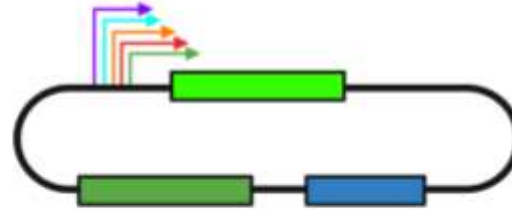


# Need for species-specific promoters

Anderson Promoter Set  
J23100-J23110



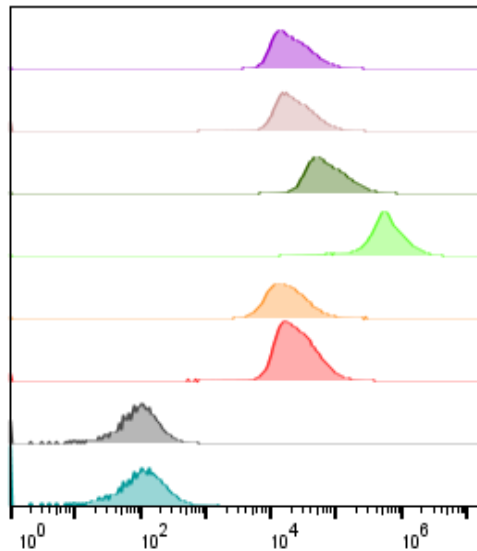
pMSBR-sfGFP-p\*\*\*



*E. coli* DH5 $\alpha$

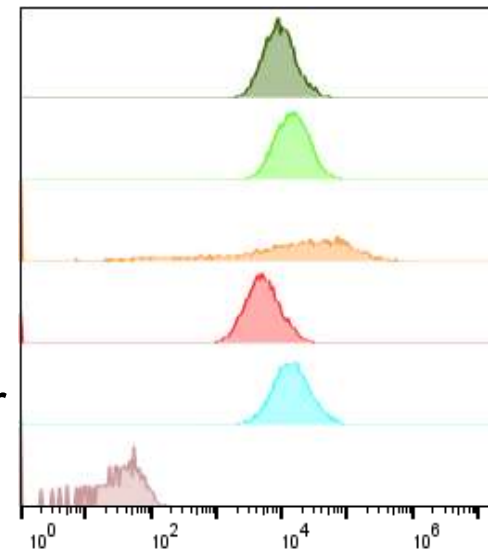
*E. cloacae*

p10  
p09  
p08  
p04  
p00



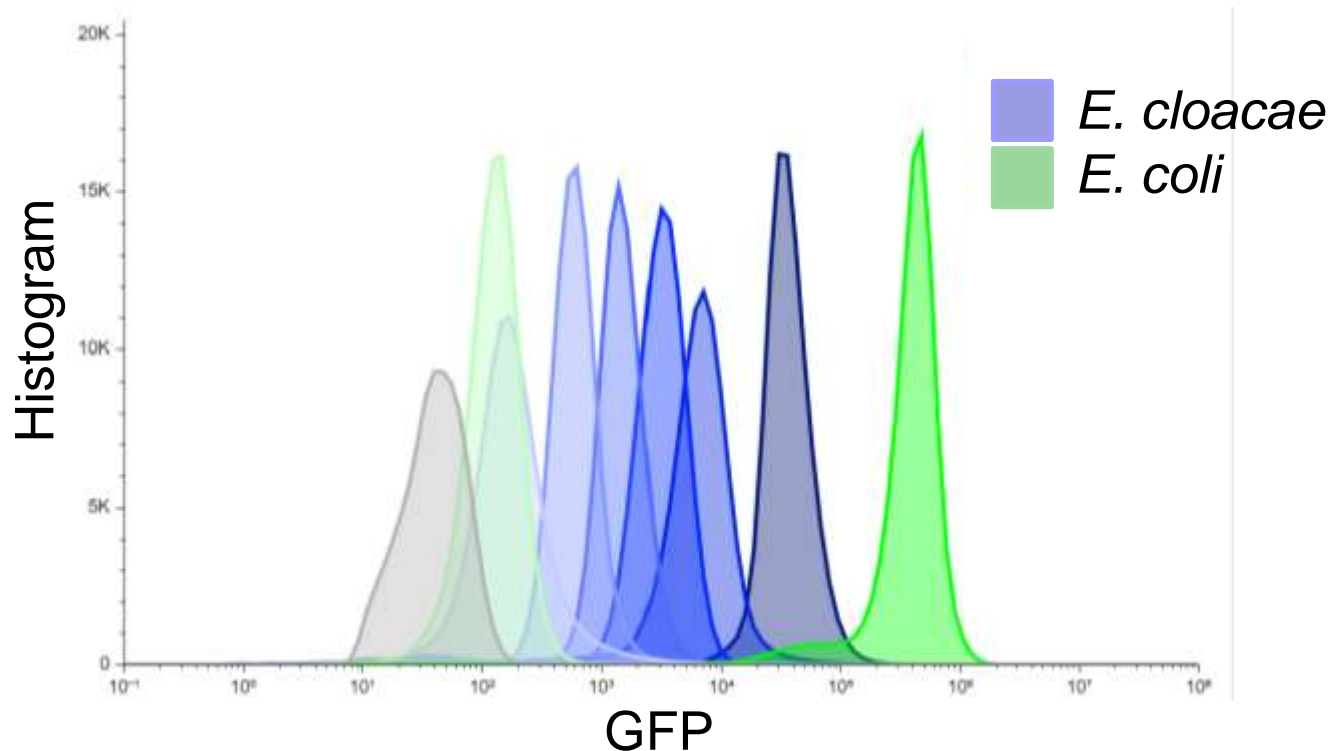
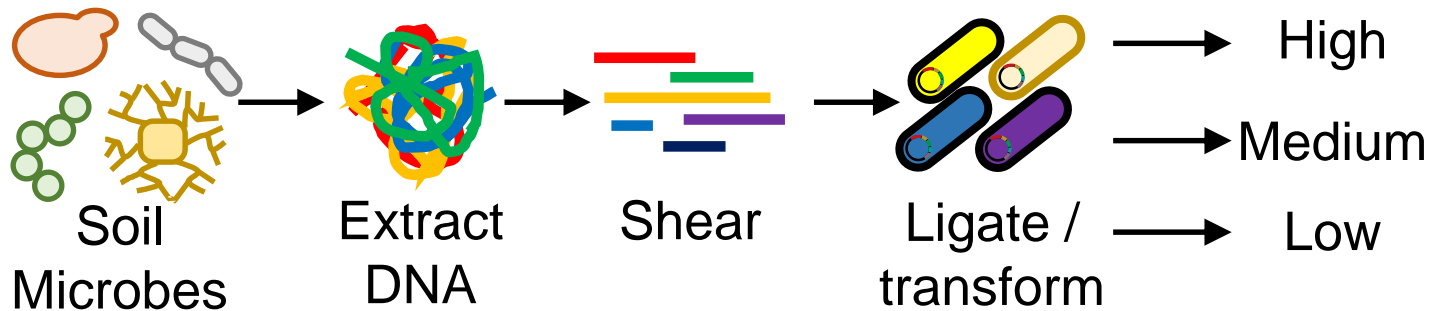
GFP

p10  
p08  
p04  
p00



GFP

# Mining the genome for promoters

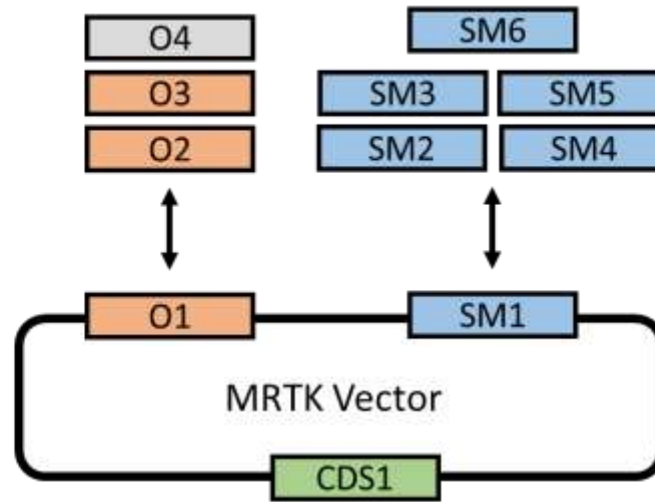




# A modular engineering toolkit

## Origin of Replication

BHR1*	O1
ColE1*	O2
SC101**	O3
R6Kγ***	O4

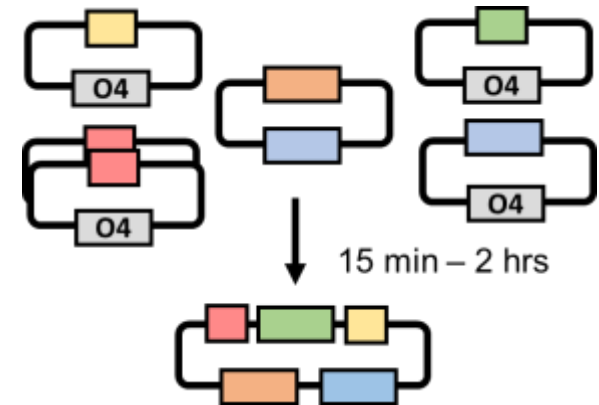
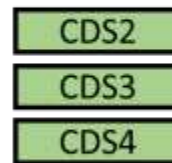


## Selection Markers

CmR	SM1
KanR	SM2
AmpR	SM3
SpecR	SM4
TetR	SM5
EryR	SM6

## Coding Sequence

sfGFP	CDS1
mCherry	CDS2
BFP	CDS3
YFP	CDS4

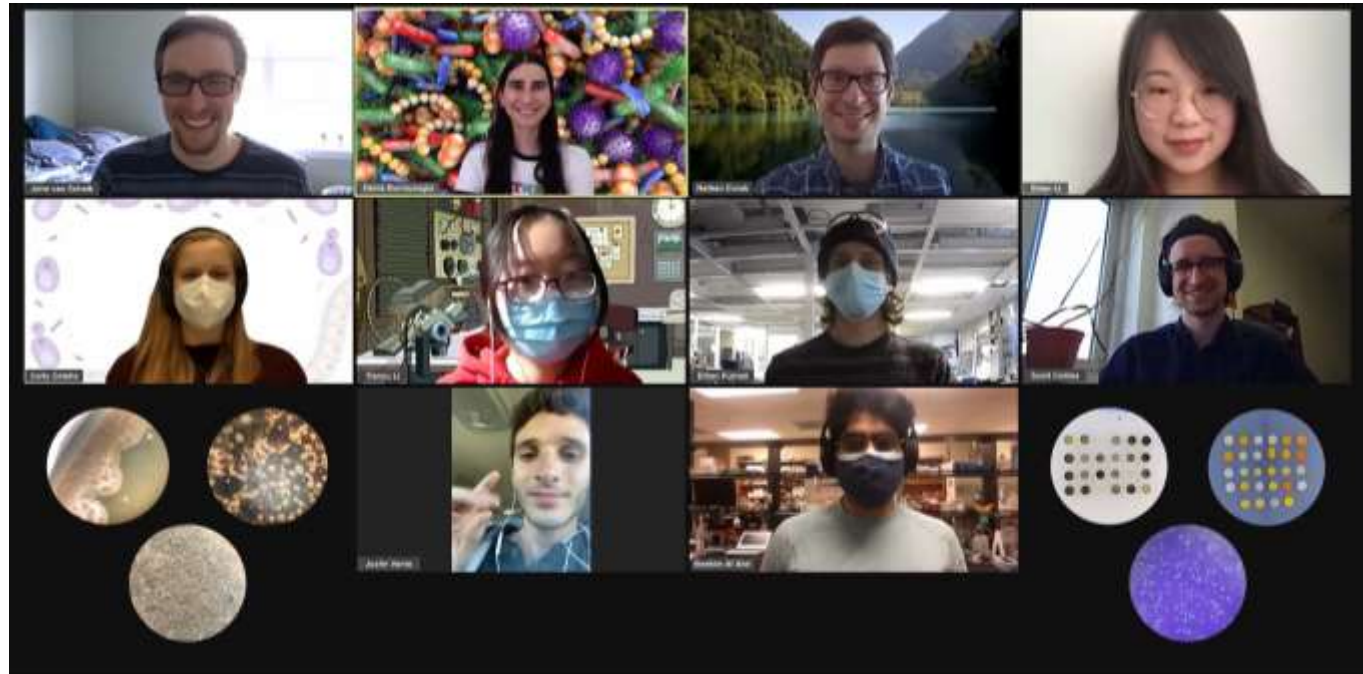


# Partnership opportunities

- Apply technologies to your favorite bacterium:
  - 6 mo – 1 yr projects, \$50k-\$100k
  - Improve transformation efficiency
  - Identify promoters at defined strength/conditions
  - Established IP process with NCSU ORC
  
- Longer-term collaborative interests
  - Bioprospect for colonization-enhancing genes
  - Directed evolution of catabolic or biosynthetic pathways

# Thank you!

Deniz Durmusoglu  
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John Van Schaik  
Zidan Li  
Tianyu Li



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