

# Getting started

Week 1

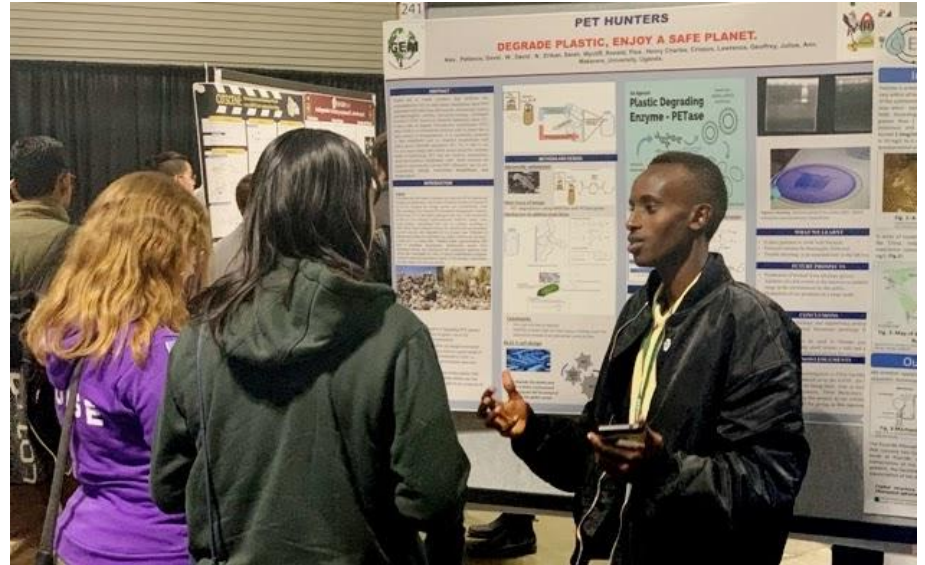
iGEM 2020 Summer webinars

Alexis Casas

# Where do you start ? Where do you end ?



Team Marburg 2018, finalist presentation



Team Makere 2018, poster presentation



Team Pasteur Paris 2018



Team Valencia UPV 2016

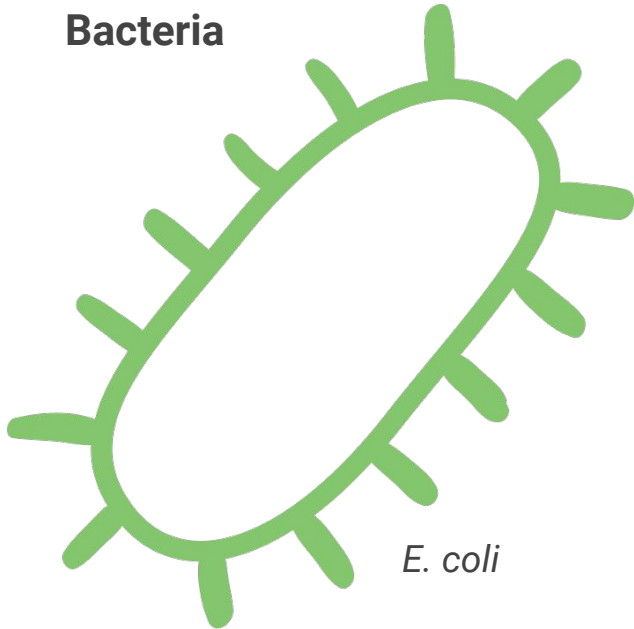
# Where do you start ?

- What are you solving ? What are you engineering ?
- What should you design *in-silico* (on your computer) ?
- What are you going to be doing in the lab and why ?
- Choice of organism ?
- How does Human Practices influence your work in the lab ?

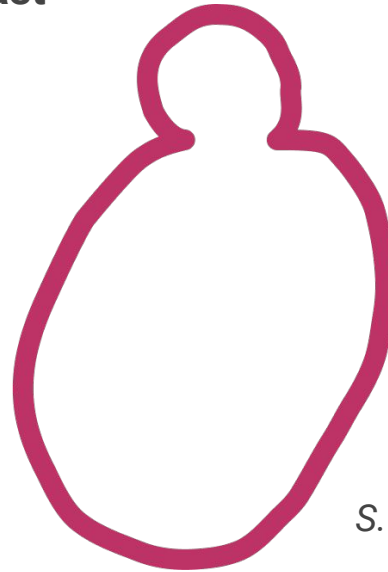
# Where do you start ?

- Choice of organism ?

**Bacteria**



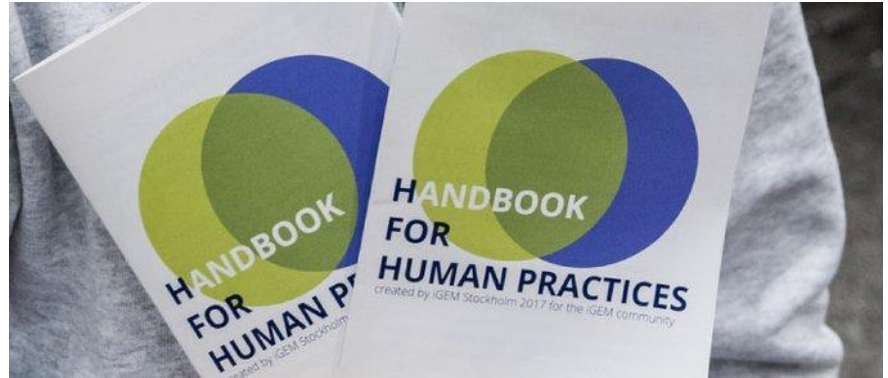
**Yeast**



**Other ?**

# Where do you start ?

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  - → Integrated Human Practices

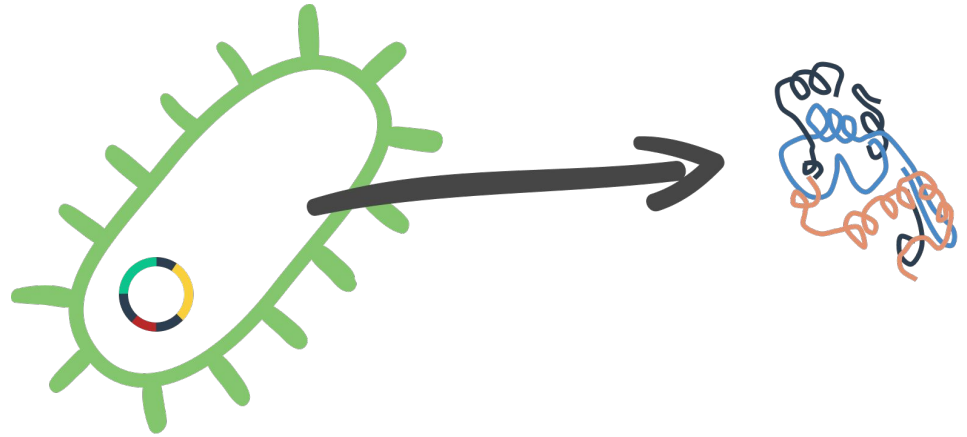


# What are you engineering ?

- DNA circuit for protein expression / production
- DNA circuit as a sensor
- Biotransformation
- Data storage in DNA

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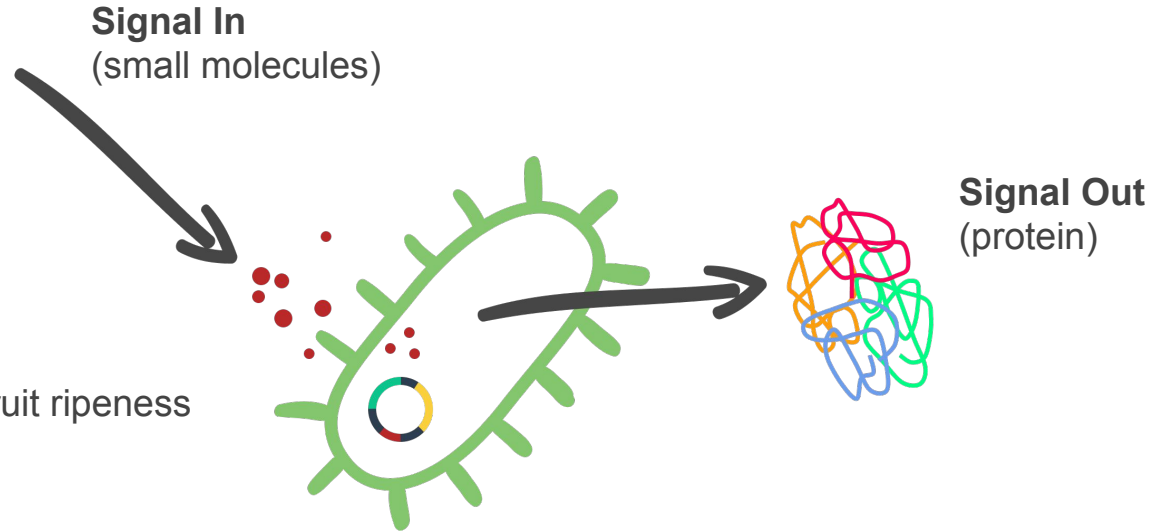


Example: Spider silk protein production  
Team GreatBay SZ 2019



# What are you engineering ?

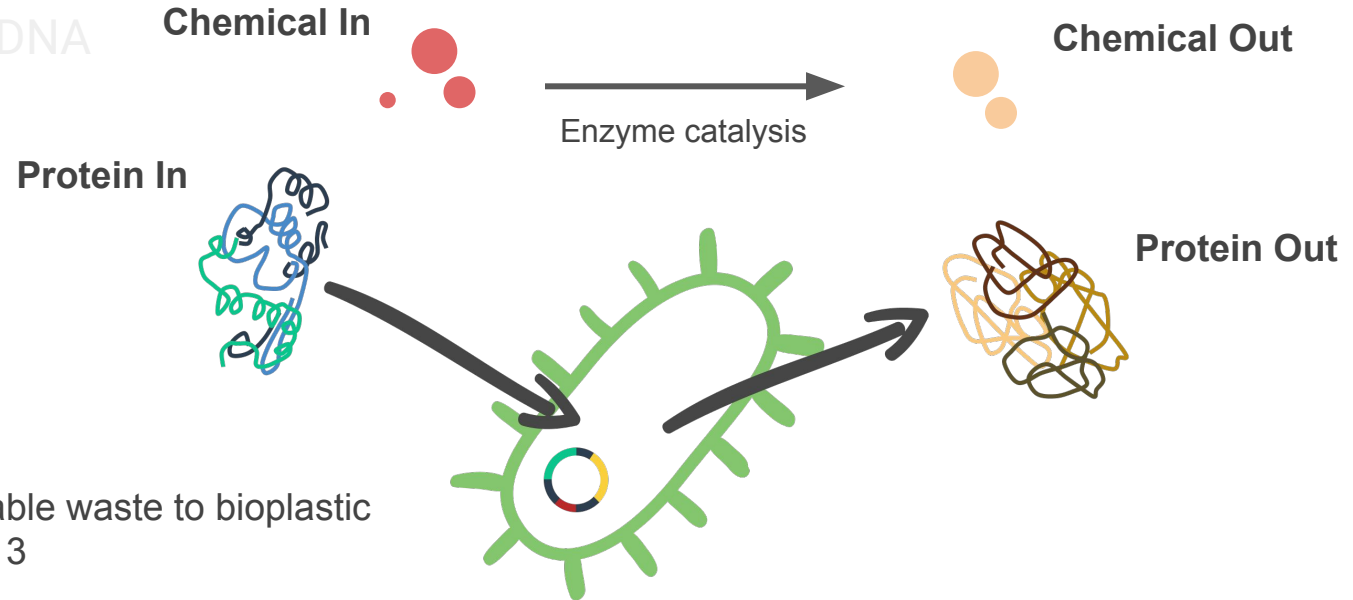
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Example: Biosensor to detect fruit ripeness  
Team Sydney 2016

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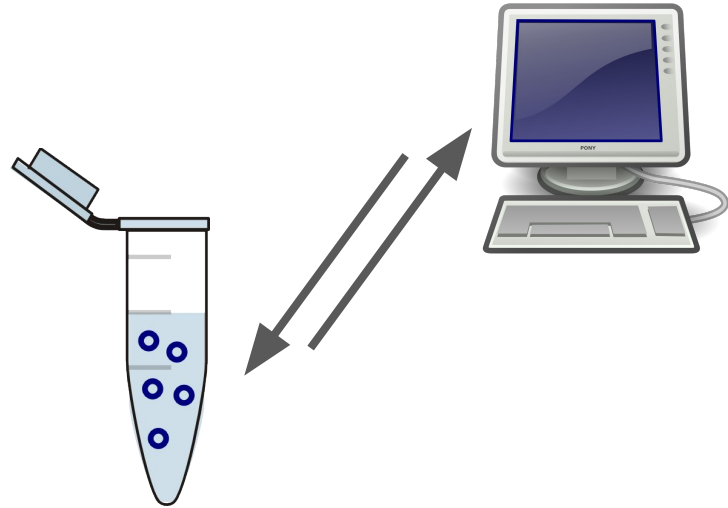
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Example: Turn non-recyclable waste to bioplastic  
Team Imperial College 2013

# What are you engineering ?

- DNA circuit for protein expression / production
- DNA circuit as a sensor
- Biotransformation
- Data storage in DNA



Example: DNA storage and encryption systems  
Team Edinburgh 2016, Team Groningen 2016

# Resources from iGEM 2020 Opening Week-end

- *Planning and Designing an iGEM Project*

- <https://youtube.com/watch?v=39nLyxYun38>



- *Cloning and Assembly Plans*

- <https://www.youtube.com/watch?v=f0Q1xeX2xzA>



- *Modelling for Synthetic Biology*

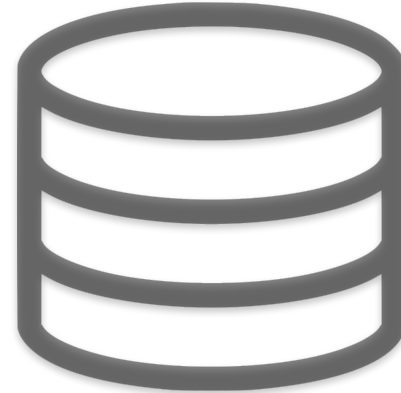
- <https://www.youtube.com/watch?v=z7isDdt0IS0>



# Tools to design your circuit



DNA design tools



Databases

# Tools to design your circuit

## DNA design tools

geneious

 Benchling

 python™

 SnapGene®

SBOL

  
biopython

# Tools to design your circuit

- Databases
  - Find proteins
  - Find genes
  - Find other parts (promoters, RBS etc..)



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Next talk: **How to utilize databases effectively** - Ian Schlander



# Designing your circuit

- Getting your DNA sequences
  - Where to search ?
  - How to use and do lookups in different databases ?
- Getting your Genes
- Designing your Transcriptional Unit (TU)
  - Choosing the promoter
  - Choosing the RBS
  - Putting the CDS together
  - Choosing the terminator
- Designing your Device (or Circuit)
  - Backbone (=Vector)
  - Resistance

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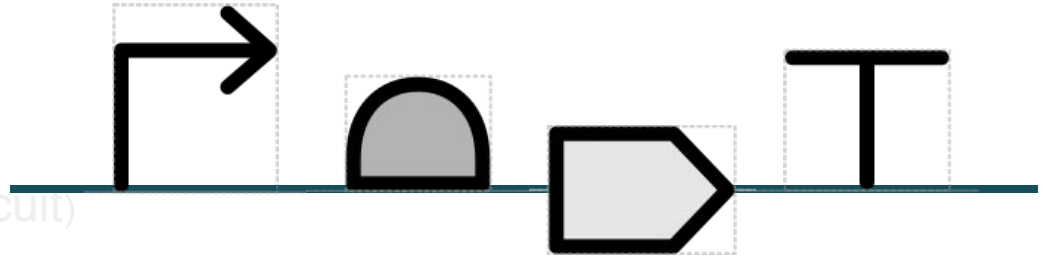
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Next talk: **Cloning Strategy** - Sonja Billerbeck

# Building your circuit

- Physically getting the parts and the genes
- Steps to get your DNA from existing plasmid
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**Week 2 - DNA parts and Basic Molecular Biology**

- Cloning to get many copies of your gene
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**Cloning Strategy** - Sonja Billerbeck



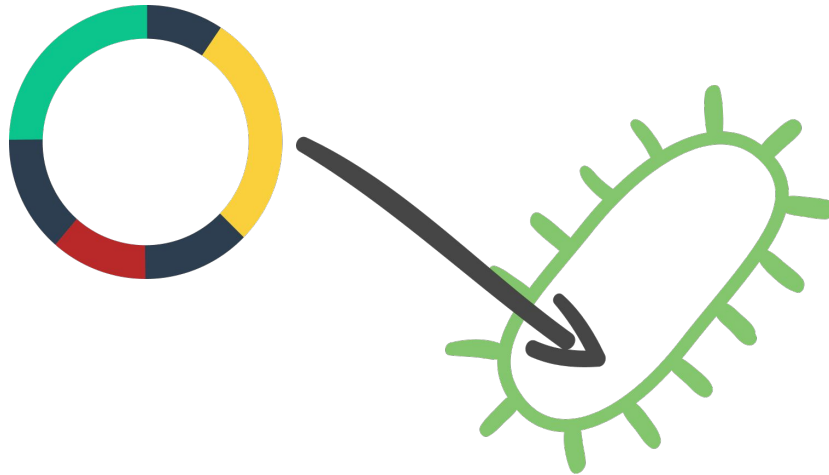
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**Week 3 - DNA assembly techniques**

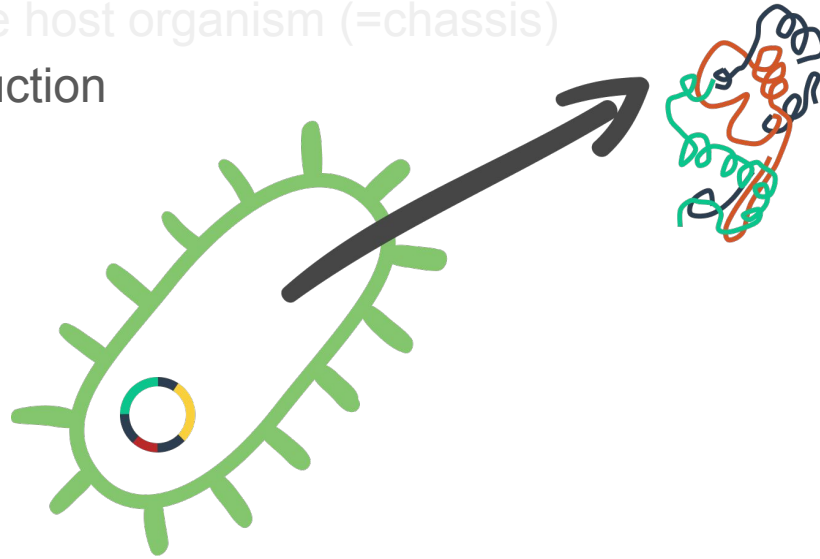
# Expressing / Transformation

- Expression vector
- Cloning in the host organism (=chassis)



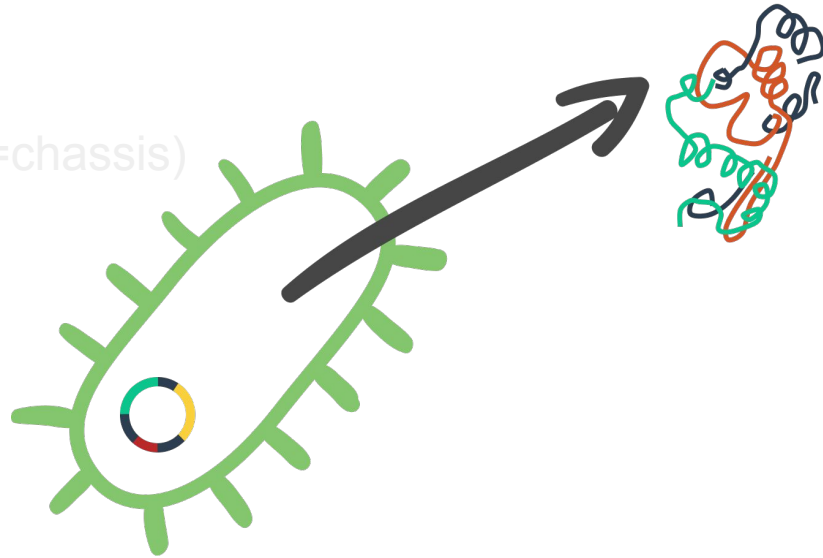
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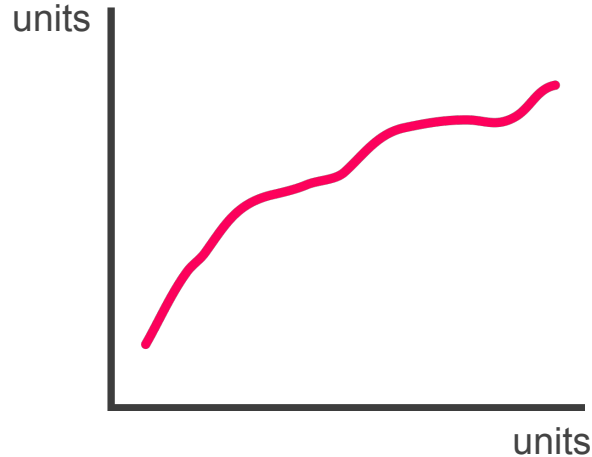
Week 2 - **DNA parts and Basic Molecular Biology**

Week 6 - **Transforming and Sequencing**

Week 8 - **Protein characterization**

# Measuring

- What do you measure ?
- Designing your measurement assays



Week 5 - **Quantifying fluorescence and cell count with plate readers**

# Troubleshooting

- Verify at each step
  - Gels electrophoresis
  - Sequencing
  - Demos (simulate troubleshooting steps) of software
  - Troubleshooting guides in products (ex: miniprep guide, vendor websites etc..)
  - Other assays
- Discuss with your mentor / PI
- iGEM Measurement Committee Office Hours
  - Biweekly office hours to answer questions
  - Every other Tuesday at 3am EDT and 1pm EDT starting June 9.

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Next Office Hours: **Tuesday 23rd of June 2020**

# Future Webinars & Events

Thank you



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The next two sessions in this webinar:

**Getting started** - Alexis Casas

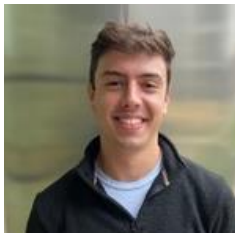
**How to utilize databases effectively** - Ian Schlander

**Cloning Strategy** - Sonja Billerbeck



# How to Use Biological Databases and Tools for Beginners

Measurement Committee  
Ian Schlander (NREL)



# Biological Databases Covered Today

1. **BioCyc** – Sequenced genomes with predicted metabolic pathways
2. **BRENDA** – Enzyme database
3. **UniProt** – Protein sequences and functional information
4. **PDB** – Protein structural data
5. **NCBI** – Biomedical and genomic information
6. **GenBank** – DNA sequences and their protein translations

# Tools Demonstrated Today

1. **NCBI: BLAST** – Aligning homologous sequences for finding a protein of interest
2. **GenBank: RefSeq** – How to find a DNA sequence of your protein in the genome of an organism
3. **Clustal Omega** – Multiple sequence alignments and constructing phylogenetic trees
4. **ExPasy** – Bioinformatics resource portal

# By the end you will be able to...

1. Understand what databases are available and how to navigate the database
2. Use database information to decide on sequences of proteins from different organisms that can be useful for their project
3. Use bioinformatic tools in conjunction with the database to analyze and visualize these proteins and their pathways to eventually decide which one to use.



# Time for you to practice!

- “Database & Bioinformatic Tools for Beginners Practicum”
  - Download the document available on our webpage:  
<https://2020.igem.org/Measurement/Webinars>
- Contact [measurement@igem.org](mailto:measurement@igem.org) and request **Ian Schlander** for further questions 😊
- Next is a presentation by Sonja Billerbeck to discuss Cloning Strategies!

# Cloning Strategy

**Sonja Billerbeck**

Assistant Professor  
University of Groningen  
Netherlands



@billerbeck\_s



[www.BillerbeckLab.com](http://www.BillerbeckLab.com)

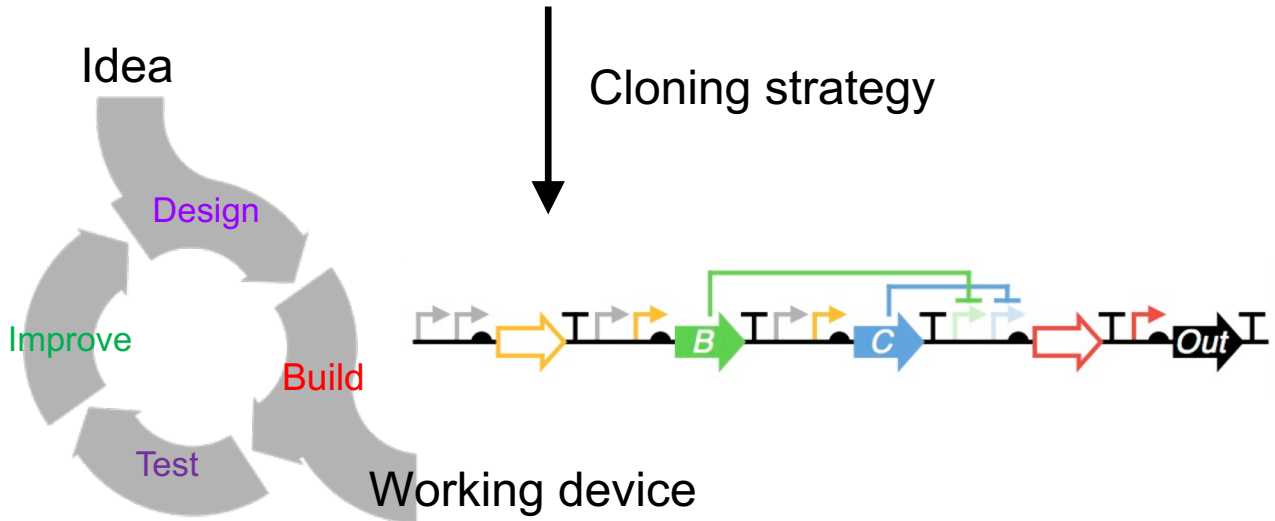


# Cloning Strategy



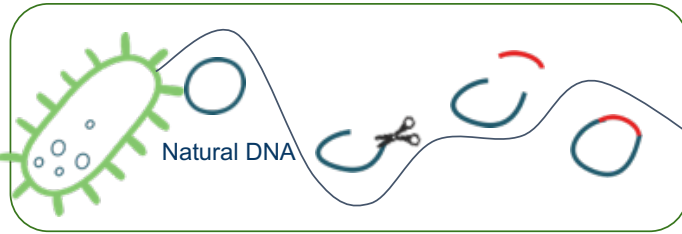
Genetic information

Cloning strategy

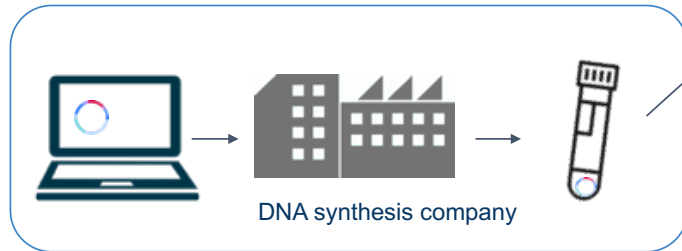
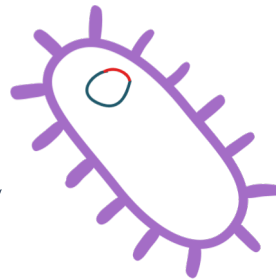


# A cloning strategy needs three major considerations

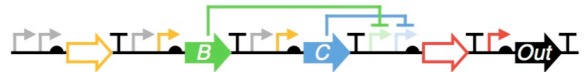
## Cloning



1. Host organism: Cloning *Backbone/vector*



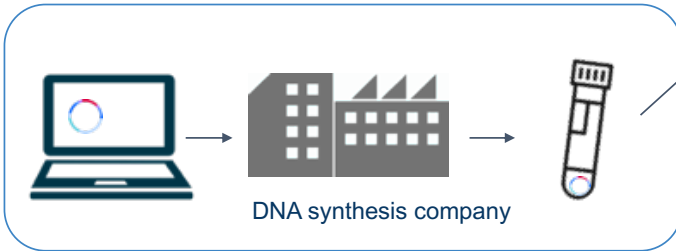
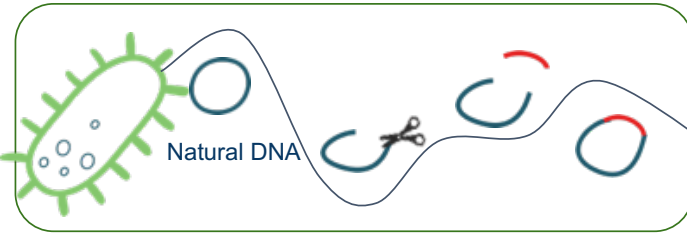
2. Assemble genes to circuits: *Assembly method*



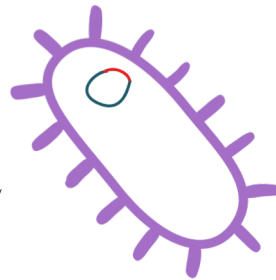
3. Troubleshoot the circuit performance:  
Flexible and *modular design and assembly*  
strategy of the backbone and the circuits

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## Cloning

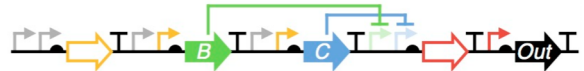


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## Cloning Strategy

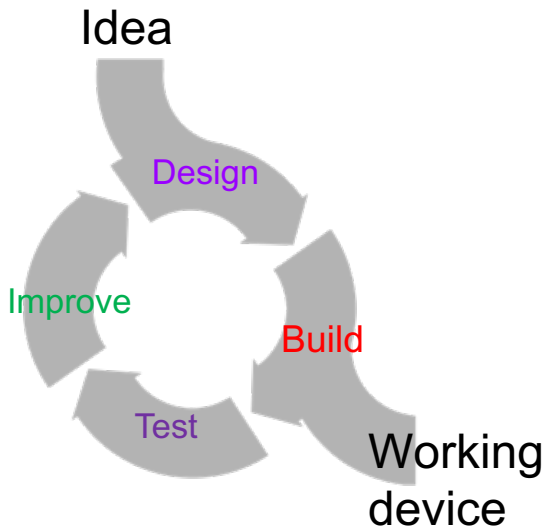
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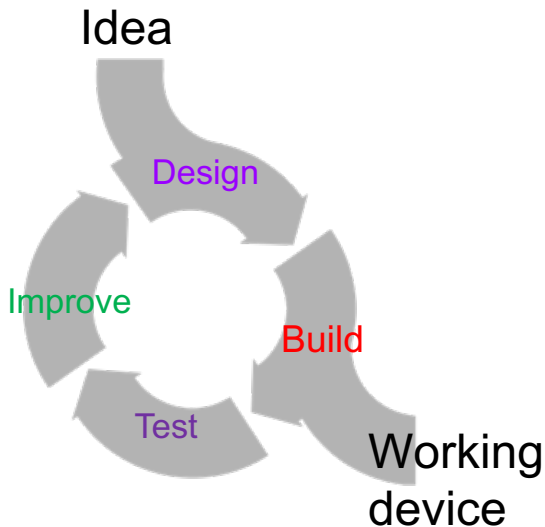
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- Cloning is an essential component of the **built** phase
- A cloning strategy enables the **improvement** phase
- The cloning strategy depends on the **design** and **test** phase



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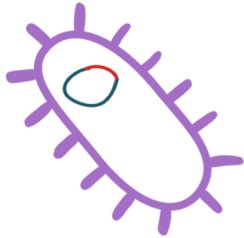


Points for consideration

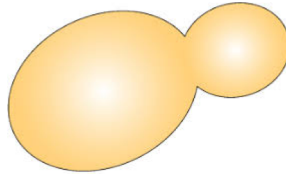


# Cloning strategy question 1: Host organism

**Bacteria** (*E. coli*)



**Yeast** (*S. cerevisiae*)

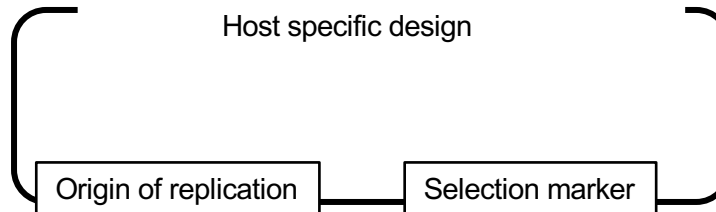


- Mammalian cells
- Plants
- Non-conventional yeast and bacteria

Circuit



**Backbone/  
Vector**



Host organism: Backbone/vector collections

# Bacteria

**pSEVA** collection

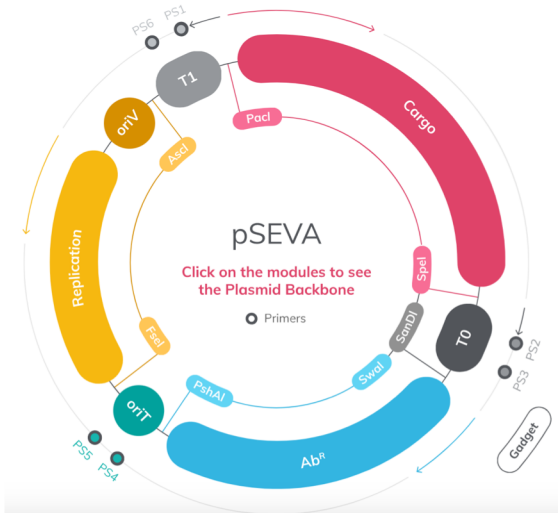


## Interchangeable

## Selection markers

## Origin of replication

## Multiple cloning site for cargo



## Yeast

- pRS series of plasmids

## Interchangeable

## Selection markers

## Origin of replication

## Multiple cloning site for cargo

Addgene #11258 pRS410

Addgene #11256 pRS418

Addgene #11256 pRS420

- Yeast MoClo Toolkit (Dueber Lab)
- Make pRS series via golden gate assembly

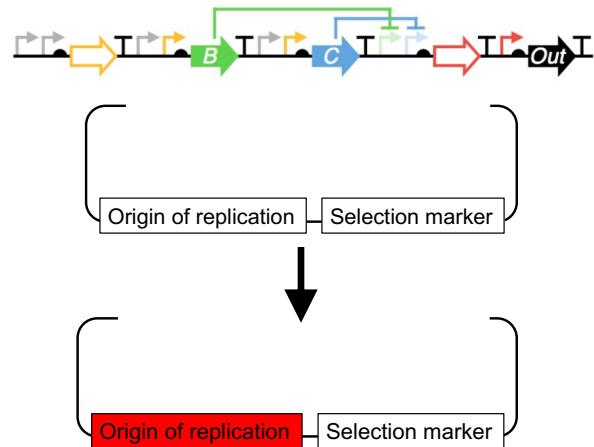
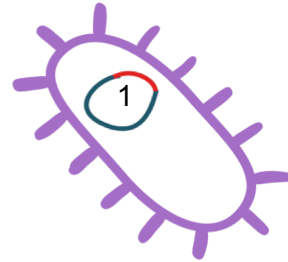
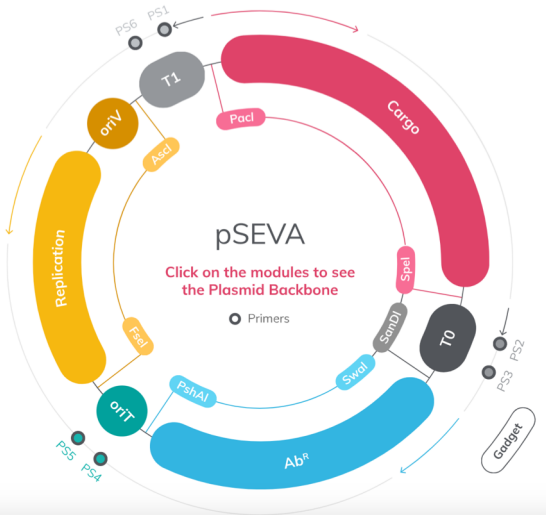
Addgene Kit #1000000061



Backbone collections: Change copy number of plasmid

## Interchangeable

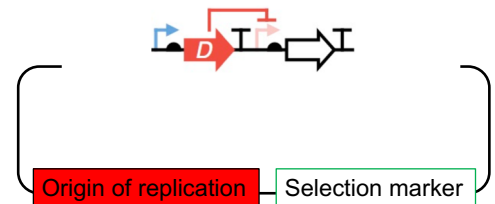
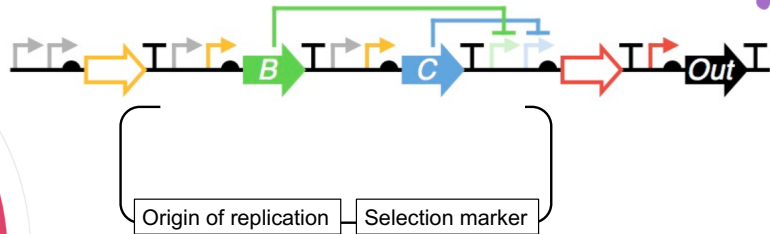
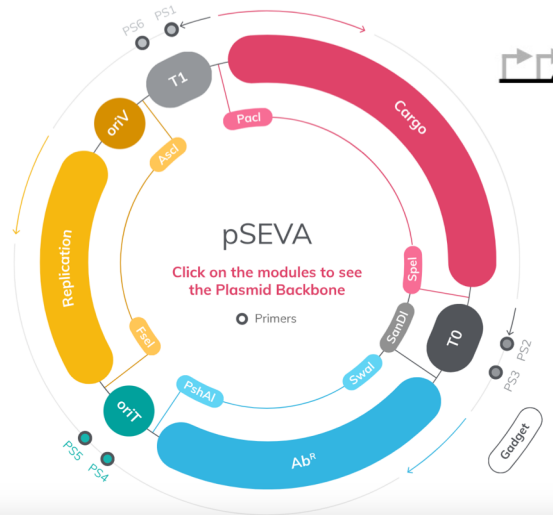
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## Backbone collections : [Add a circuit](#)

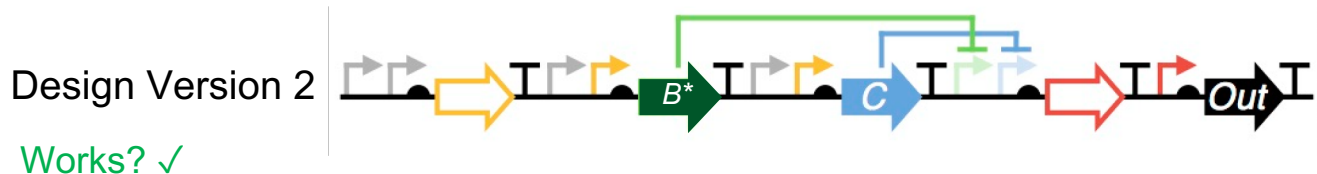
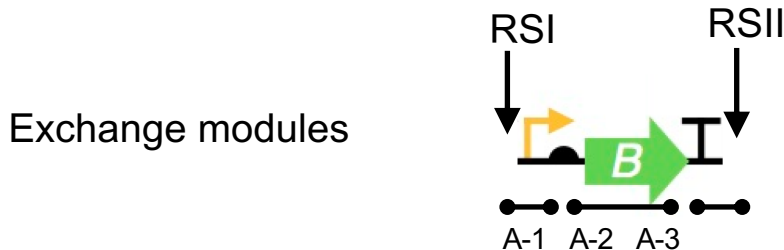
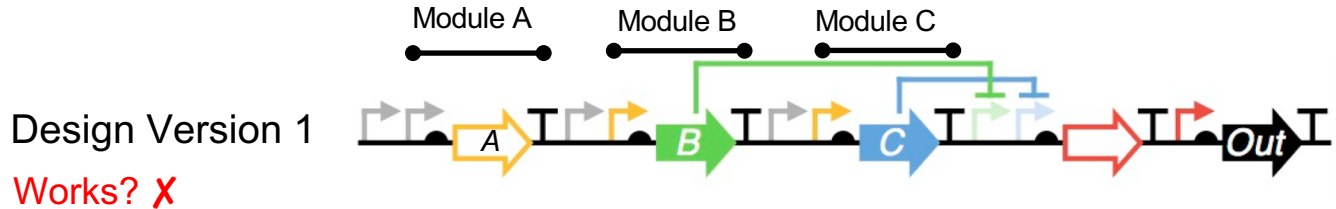
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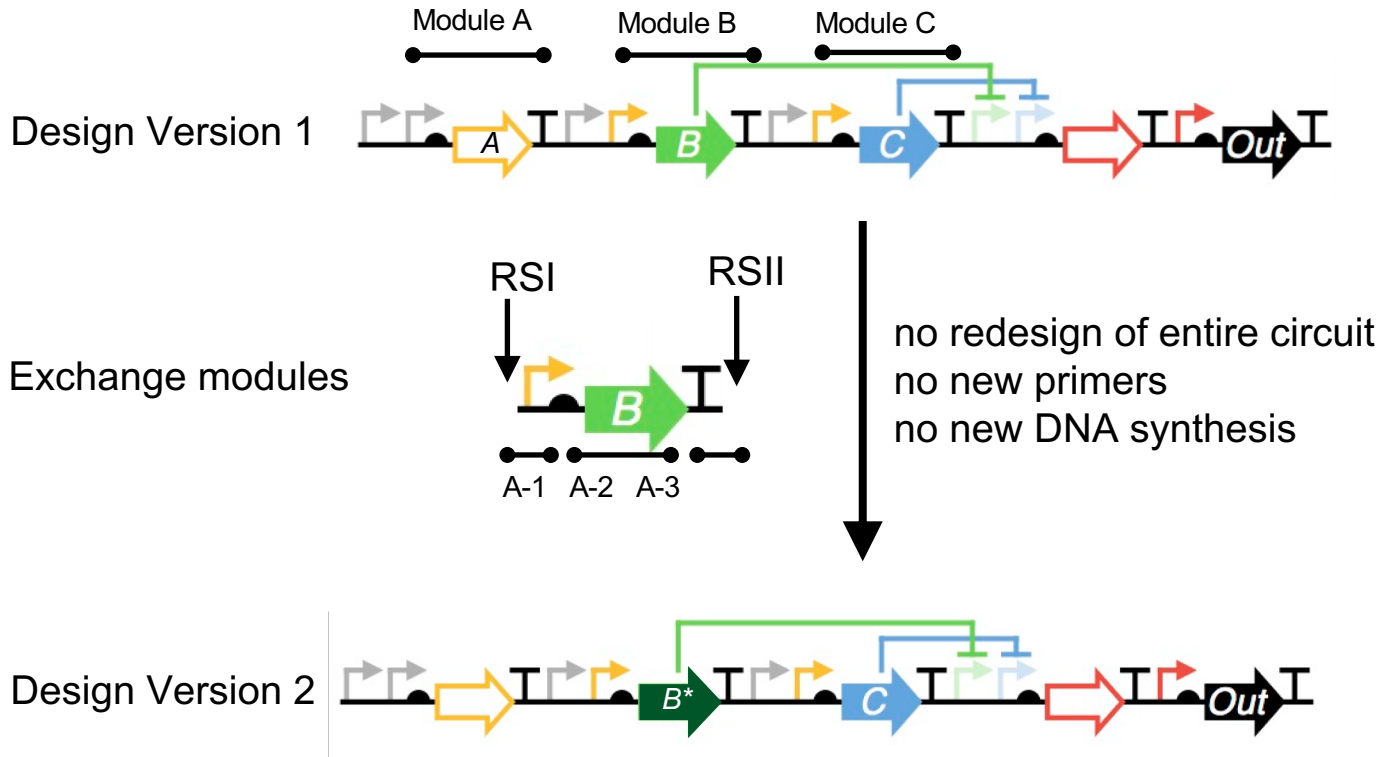


# Cloning strategy question 2: Assembly of circuits:

## Modularity and troubleshooting



# Modularity allows for **quick and cheap** troubleshooting

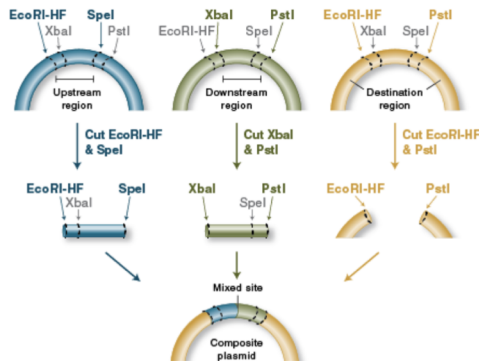


# Assembly of circuits: **Assembly method**

## Webinars

### BioBrick Assembly

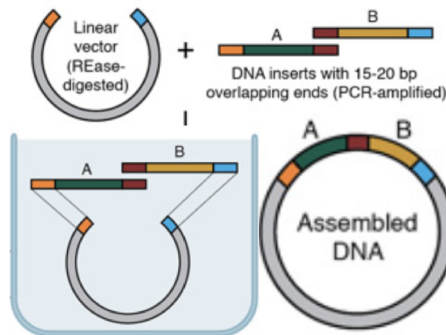
- Restriction sites flank every gene fragment, allowing “parts” to be interchanged, but introduces scar sequences



Modular ✓

### Gibson Assembly

- Exonuclease creates large overhangs for annealing fragments, allowing for more accurate assemblies

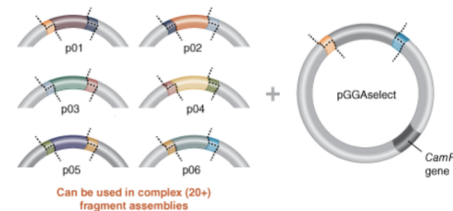


Modular ✓

Scar-less ✓

### Golden Gate Assembly

- Endonuclease creates fragment-specific overhangs allowing 20+ fragments to be assembled at once in a relatively short time



Modular ✓

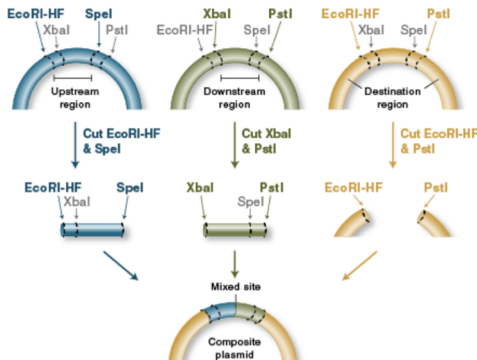
Scar-less ✓

Scalable ✓

## Assembly of circuits: Assembly method

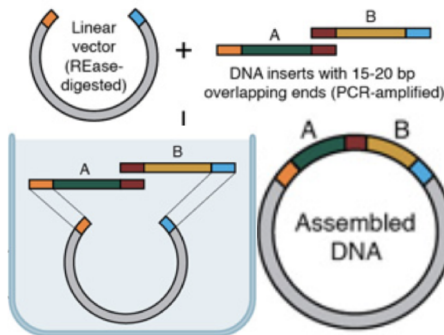
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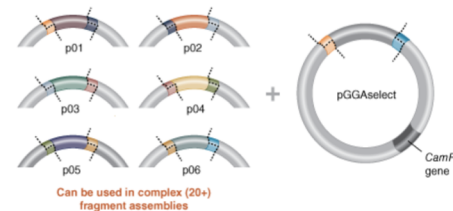
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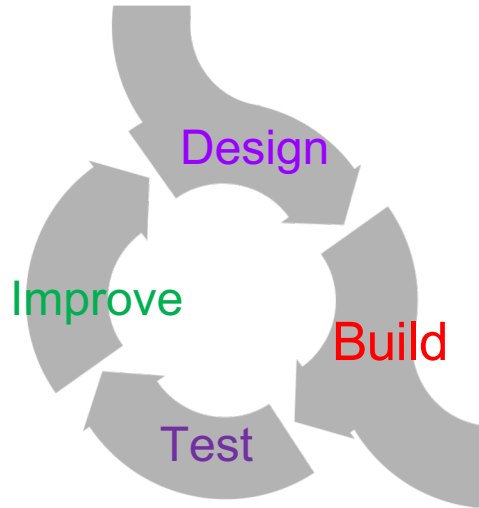
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Good choice for most cloning strategies

A well-defined cloning strategy is...



...essential to run the full cycle and improve a design until it works.

**Specialized webinars on:  
Cloning Apps and Assembly methods**