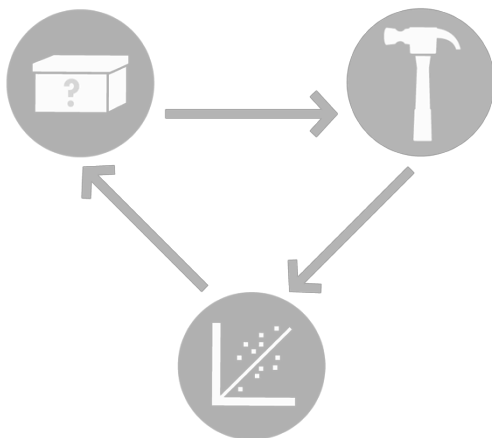


# BioBuilder

*Synthetic Biology for Students*

## Golden Bread

Explores the science, engineering and bioethics of a yeast that's genetically modified to make a vitamin-enriched food. Lab activities include PCR, yeast transformation, codon shuffling and quantitative analysis of data



# BioBuilder

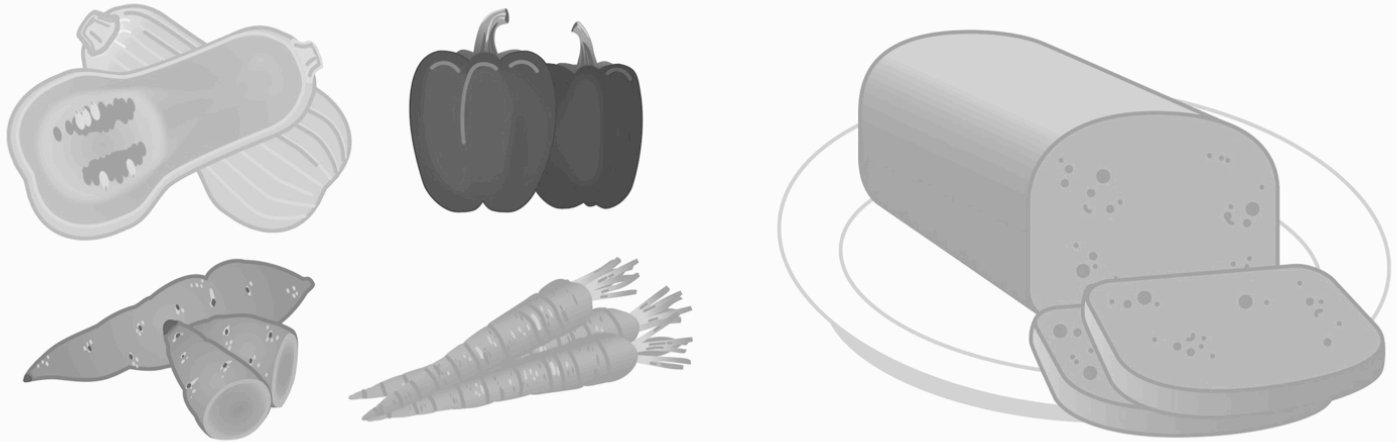
## Golden Bread

### Table of Contents

<i>About Golden Bread</i>	<i>03</i>
<i>Engineering Golden Yeast</i>	<i>04</i>
<i>Engineering Challenge</i>	<i>05</i>
<i>Engineering Toolbox</i>	<i>06</i>
<i>Re-Engineering Golden Bread</i>	<i>07-08</i>
<i>Pre-Lab Questions</i>	<i>09</i>
<i>Golden Bread poster</i>	<i>10</i>
<i>Golden Bread Protocol</i>	<i>11-12</i>
<i>Post-Lab Questions</i>	<i>13</i>

# ABOUT GOLDEN BREAD

This lab focuses on a strain of baker's yeast that has been modified to produce  $\beta$ -carotene, a nutrient we naturally obtain from eating foods such as carrots, sweet potatoes, and broccoli. In the body,  $\beta$ -carotene is converted to vitamin A, which is crucial for vision, the immune system, and other biological functions.



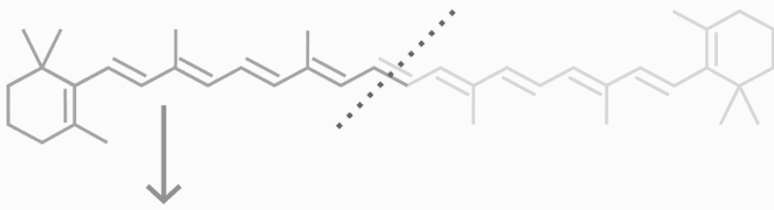
In some developing countries that struggle with malnutrition, vitamin A deficiency is a critical public-health issue. Researchers hope that an engineered strain of baker's yeast designed to generate  $\beta$ -carotene, like the one in this activity, could be used in bread to treat vitamin A deficiency. Such bread might appear a golden color from the added vitamin, hence the name, "Golden Bread."

The Golden Yeast was developed as part of an iGEM Project called "VitaYeast." The iGEM team wanted this yeast to substitute for standard baker's yeast, making it possible to bake vitamin A-enriched loaves of bread. The iGEM team worked with an engineered version of baker's yeast, extending some work published in 2007 by researchers who genetically manipulated the strain known as *Saccharomyces cerevisiae*. The modified yeast could express all their usual genes plus three  $\beta$ -carotene biosynthesis genes isolated from another fungus.

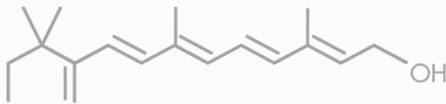


# ENGINEERING GOLDEN YEAST

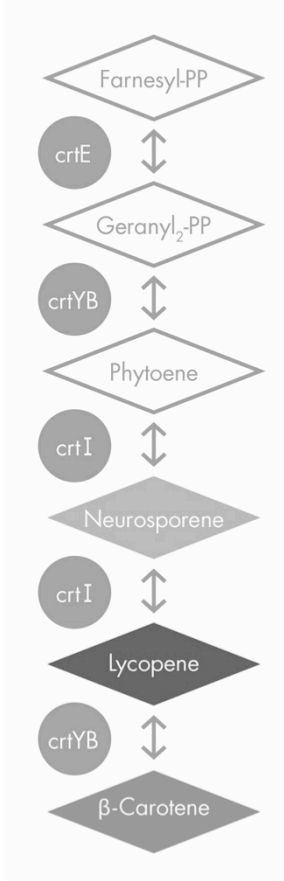
Beta carotene



Retinol (vitamin A)



The metabolic pathway for making vitamin A consists of three enzymes that convert farnesyl phosphate to  $\beta$ -carotene, which then spontaneously breaks in half to become vitamin A.



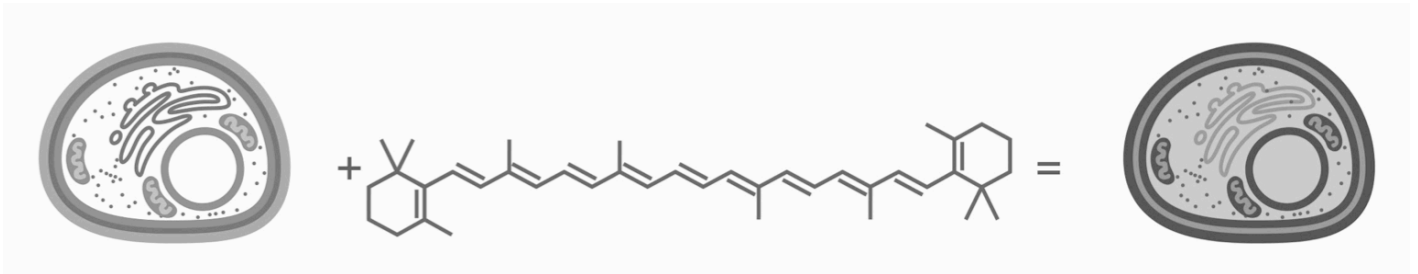
The baker's yeast strain, *S. cerevisiae*, naturally produces farnesyl diphosphate. The strain also expresses an enzyme encoded by the *BTS1* gene that converts the farnesyl diphosphate to geranylgeranyl diphosphate. Other yeast use a similar gene called *crtE* for this process. Converting geranylgeranyl diphosphate into  $\beta$ -carotene requires the action of two more genes, *crtYB* and *crtI*, which are not naturally found in *S. cerevisiae* and so they were engineered into baker's yeast from a different, red-colored yeast called *X. dendrorhous*.

Interestingly, each of these enzymes serves double duty when making  $\beta$ -carotene. The *crtYB*-encoded enzyme plays a role early in the synthesis, converting geranylgeranyl diphosphate into phytoene and then comes back into play for the last step of the synthesis, converting lycopene into  $\beta$ -carotene. Between the *crtYB* enzyme-catalyzed steps are two reactions that require the activity of the *crtI* enzyme, which was also imported into the baker's yeast strain from the red-colored yeast. The enzyme converts the phytoene first to neurosporene and then to lycopene.

Nature has provided a simple way to detect pigments produced by this pathway, resulting in a convenient visual test for the design. The first three compounds in this pathway are colorless, but the last three are colored yellow, red, and orange, respectively. Unengineered yeast appear white, whereas yeast successfully making  $\beta$ -carotene turn bright orange. Yeast making mostly lycopene turn red like a tomato, which has a naturally high lycopene concentration, and yeast making mostly neurosporene appear yellow. Engineered cells that appear white may have lost one or more of the *crt* genes.

# ENGINEERING CHALLENGE

Researchers knew they had successfully added the three  $\beta$ -carotene biosynthesis genes into baker's yeast, *S. cerevisiae*, because they saw that the normally white-colored cells were grown as orange colonies.



Much to the researcher's disappointment, though, the strain was not orange 100 percent of the time. When streaked out on a petri dish, the engineered yeast strain grew as orange colonies most of the time, but they could also see red, yellow, and white colonies, indicating that some of the steps in the pathway were not working.

Undaunted, they took two approaches to improving the reliability of the strain's  $\beta$ -carotene production.

First, they stopped using the easy-to-work-with plasmids and instead moved the *crtYB* and *crtI* genes into the chromosome of the baker's yeast they were building. These integrated copies of the genes were less likely to be lost from the yeast, and so the strains were expected to be more reliably orange-colored.

Second, they tried to improve the production of  $\beta$ -carotene by adding a second copy of the *crtE* gene and a second copy of the *crtI* gene. They hoped these extra copies would make more of the needed enzymes. The second copy would also provide a backup in case the first copy failed. This concept of redundancy is explored further in the engineering toolbox.

Much to their disappointment, however, the strain was still unstable, giving rise to orange, red, yellow and white colonies.



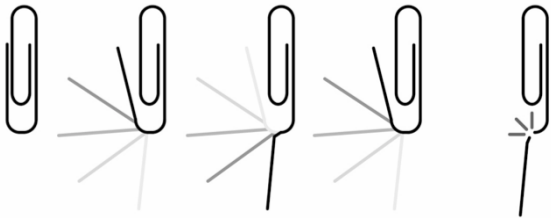
Your engineering challenge is to investigate this instability and to improve the strain's performance.

# ENGINEERING TOOLBOX

For any new food or drug to become widely available for use, manufacturers must show that they can reliably produce the material. In fact, reliability is crucial for nearly all engineering endeavors. How do engineers think about and then build-in reliability?

## Concept 1: Mean time to failure

The mean time to failure (MTF) helps designers predict when a system will break. It also guides the designer on when and how to intervene through regular maintenance of the system. Engineers include MTF calculations in their design process so that they can recommend when parts should be serviced and how to use them for greatest longevity.



A paperclip's MTF: Bending a paperclip back and forth, as shown here, will eventually cause it to break. The number of bends before breaking can be used to calculate MTF.

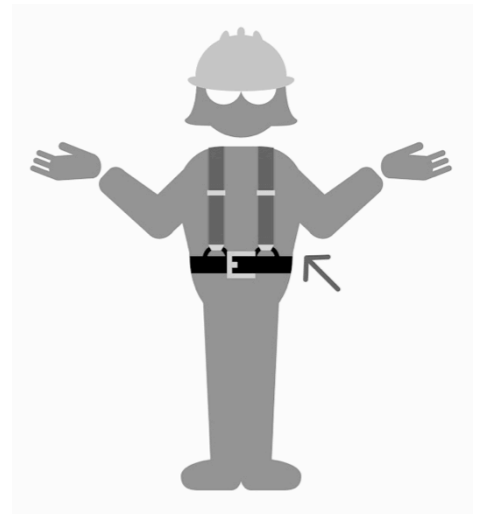


*In the context of this experiment you can ask: how easily does this system “break,” i.e. fail to produce  $\beta$ -carotene? Also, can we lengthen the MTF to improve the performance of the system?*

## Concept 2: Redundancy

Building redundancy into an engineered system is another technique used in many fields to ensure more reliable performance. For example, engineers working on automotive safety have chosen to deploy both seat belts and airbags to protect passengers in a car crash. Having two safety systems increases the likelihood that passengers will be unharmed in a crash, and so they are worth the added cost.

Redundancy in living cells is also important for their survival. DNA can be damaged by mutagens in the cell's environment, inducing changes in the DNA sequence and making some of the genetic instructions essentially unreadable. Having two copies of the genome is a natural form of genetic redundancy and provides the cell with some insurance.



**By adding suspenders to a belt, you can be extra confident that there will be no wardrobe malfunctions.**



*You will apply these engineering tools to investigate the effect of a second copy of *crtYB* for restoring  $\beta$ -carotene production in isolated white colonies.*

# RE-ENGINEERING GOLDEN BREAD

To fix the genetic instability seen in the Golden Yeast strain you will rely on the engineering strategy of redundancy.

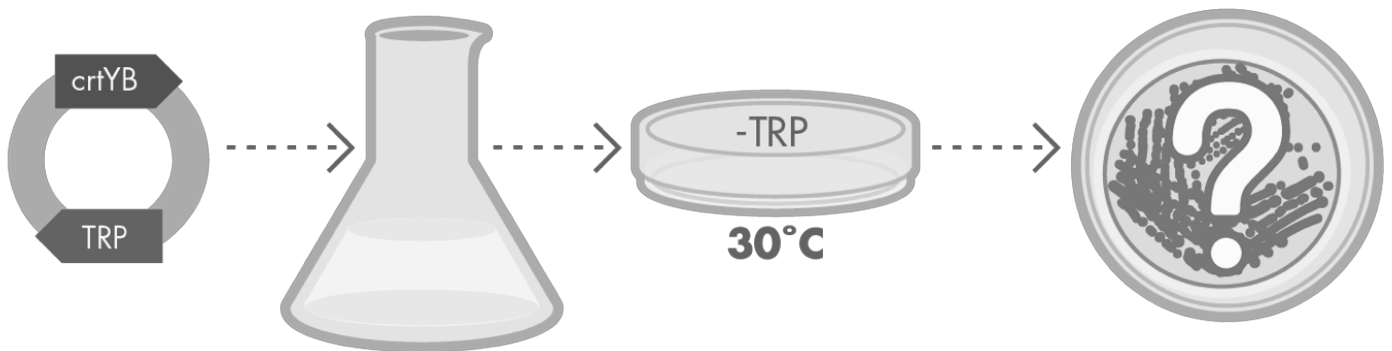
You will focus on the *crtYB* gene because it is the only gene of the system that does not have any redundancy already engineered into the strain. The second copy of the gene will be called *crtYB\** to distinguish it from the copy that is already in the strain. To make *crtYB\**, the DNA sequence of *crtYB* was “**codon shuffled**” and then ordered from a DNA synthesis company. Codon shuffling substitutes synonymous codons for the ones in the natural gene, for example writing TTC in place of TTT in the gene since both DNA sequences encode the same amino acid, phenylalanine. The *crtYB\** gene sequence was inserted into a plasmid to more easily insert the gene in the engineered yeast.

Your experiment will also focus on the white colonies of engineered yeast, asking if the second copy of *crtYB* can restore their orange color. This kind of experiment is called a “**genetic complementation**” test or sometimes a “*cis-trans* test.” If a component of a pathway can be added back to restore a function, then you have good evidence that the component was the broken part in the first place.

## Experimental Methods

### TRANSFORMATION

The process of introducing new DNA into cell is called transformation. The Golden Yeast strain is not naturally ready to take new DNA in from the environment. To prepare the yeast for this experiment, they must be washed with water and then mixed with a salt solution that makes the strain porous. New DNA can be introduced into the strain when the cells are in this “competent” state.



In order to distinguish cells that have taken in the *crtYB\** DNA from those that haven't, the experiment depends, again, on genetic complementation. Because of a defect in their genomic DNA, the Golden Yeast are not able to make their own tryptophan, a necessary amino acid for growth. The cells will grow on “complete” or “rich” media called YPD because the media provides tryptophan for the cells to use. If asked to grow on media that does not have tryptophan, the cells will die unless they have taken in the a plasmid that encodes the tryptophan synthesis enzyme. For this experiment, the *crtYB\** plasmid also carries the gene for tryptophan synthesis.

# RE-ENGINEERING GOLDEN BREAD Continued

## Samples

### NEGATIVE CONTROL

No DNA. This reaction is done to confirm that there will be no growth on media lacking tryptophan if there is no external DNA added to the cell.

### POSITIVE CONTROL

DNA encoding the TRP gene but no *crtYB\** sequence. This reaction is done to confirm that the process of transformation is working. It also confirms that any changes seen with the experimental sample are due to the *crtYB\** gene and not to the process of transformation itself.

### EXPERIMENTAL SAMPLE

DNA encoding the TRP gene and the *crtYB\** gene. If the white colonies have are defective in the *crtYB* enzyme but encode *crtI* and *crtE*, then the DNA will complete the  **$\beta$ -carotene** pathway and the cells should appear orange.

## Predictions

What are the expected outcomes for the three transformations if the media shown below has no tryptophan? It may not be possible to know how many colonies will grow on some of the plates, but if the white colonies have a defect in both the TRP gene and the *crtYB* gene, then you can predict the color of the colonies on the templates below



Negative Control (no DNA)



Positive Control DNA (+TRP gene)



Experimental (+TRP +*crtYB\**)



# PRE-LAB QUESTIONS

The genus and species name for commonplace baker's yeast is \_\_\_\_\_.

By engineering baker's yeast with three genes from another yeast species, the baker's yeast should express \_\_\_\_\_, and so should appear \_\_\_\_\_.

An engineered yeast that appears red instead of orange might have a defective \_\_\_\_\_ gene.

The addition of an airbag into a car that has seatbelts is an example of \_\_\_\_\_ which is expected to increase reliable passenger safety in the event of a car crash.

To increase the reliable production of  **$\beta$ -carotene** by the engineered yeast, the researchers made two modifications to the strain. Name them.

(1)

(2)

Why was the *crtYB* gene chosen for the transformation experiment performed here?

The gene design technique that replaces the triplet codons in a gene with a synonymous codons is called \_\_\_\_\_. This technique maintains the amino acid sequence in a protein but adjusts the DNA sequence encoding it.

If the pathway for  **$\beta$ -carotene** synthesis is broken so the cells grow as white colonies rather than orange, which genes might be defective?

If transformation of the *crtYB\** gene into the white cells turns them orange, what do you know about the genetic defect in the white ones?

Prior to transformation the engineered strain is expected to grow on "complete media" but not media that lacks \_\_\_\_\_.

You are expecting the negative control for transformation \_\_\_\_\_.

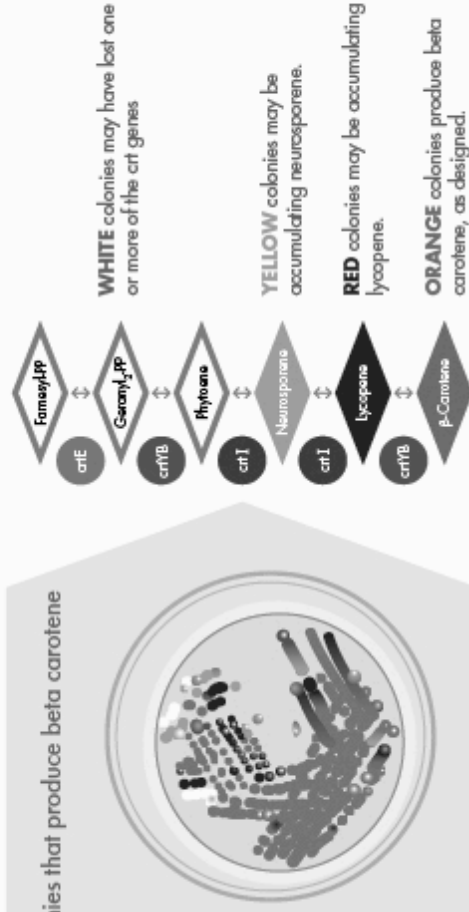
You are expecting the positive control for transformation \_\_\_\_\_.

## THE SYSTEM

Genes from *X. dendrorhous* can be added to *S. cerevisiae*, resulting in colonies that produce beta carotene

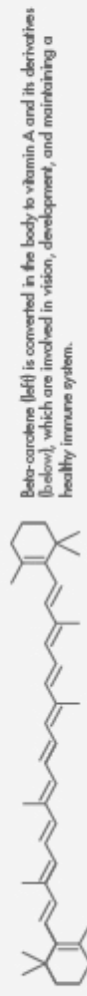


## A NEW HYPOTHESIS



## THE BIG PICTURE

Baking with beta-carotene producing yeast creates bread "biofortified" with vitamin A

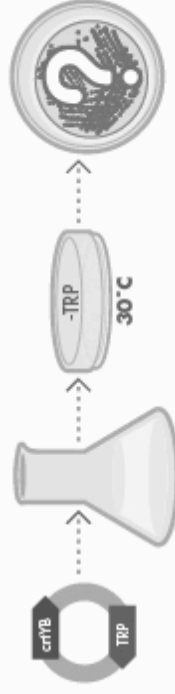


## ENGINEERING WITH REDUNDANCY

GENES in Vita YEAST ENZYMES in Vita YEAST

- Bifunctional enzyme: Phytoene Synthase/Lycopene B-Cyclase
- Geranylgeranyl Diphosphate Synthase
- Phytoene Desaturase

The Vita Yeast strain has two copies of every enzyme in the pathway except for crtYB. Will adding an "extra" copy of this gene increase the strain's robustness, i.e. eliminate non-orange colonies?





ATCG...

QUICK GUIDE: GOLDEN BREAD



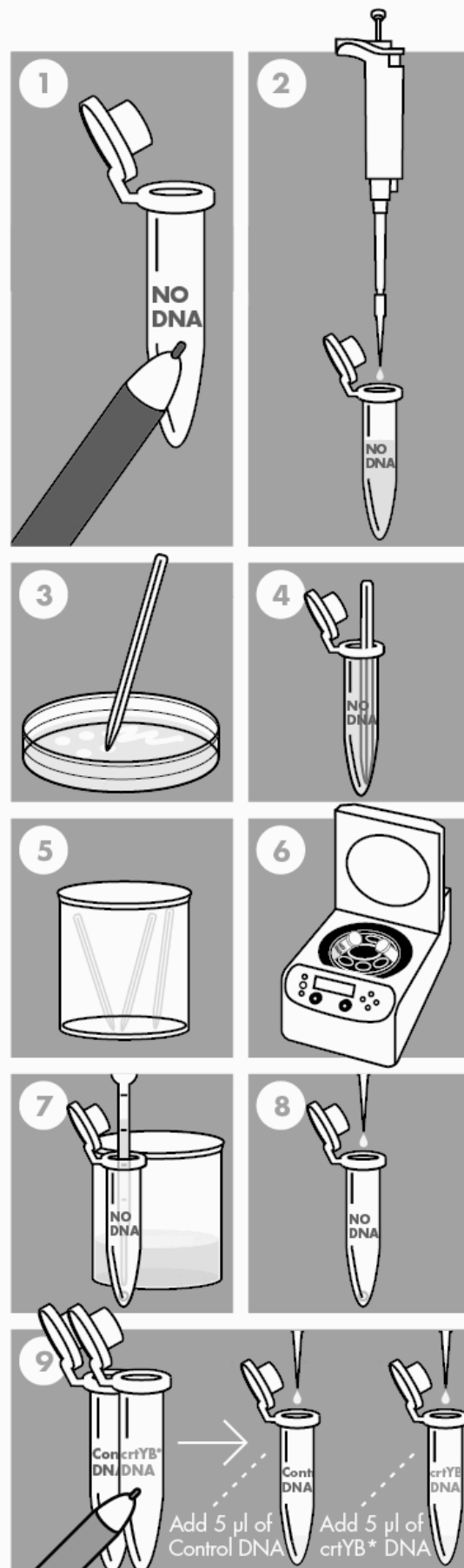
# IN ADVANCE

Melt YPD in microwave and pour plates.

Restreak "Golden Yeast" onto YPD to isolate single white colonies\*\*

# DAY OF LAB

1. Label 1 microfuge tubes: **NO DNA**.
2. Add 500  $\mu$ l of **EZ - Solution I** to the **NO DNA** tube.
3. Use a sterile pipet tip, toothpick or inoculating loop, scrape an isolated white colony of yeast off the petri dish.
4. Swirl the colony into the **EZ - Solution I** in the **NO DNA** tube.
5. Discard the pipet tip, toothpick or inoculating loop into a waste receptacle to be decontaminated.
6. Microfuge the tube you have prepared, coordinating with another group or setting up a "blank" microfuge tube with 500  $\mu$ l of water. Spin the tubes for 30 seconds at full speed.
7. After microfuging the tubes, the cells will have collected as a white pellet in the bottom of the tube. Remove as much of the supernatant as you can, using a pipet and discarding the liquid into a waste receptacle to be decontaminated.
8. Resuspend the pellet in 150  $\mu$ l of **EZ - Solution II**, pipetting up and down to make a homogeneous solution.
9. Label two more microfuge tubes: **Control DNA**, **crtYB<sup>+</sup> DNA**. Add 5  $\mu$ l of the appropriate DNA to the tubes, changing tips between aliquots. Skip this step if the DNA has already been aliquoted for you.



10. Add 50  $\mu$ l of cells from the **NO DNA** tube to the **Control DNA** tube. Pipet up and down to mix.

11. Add 50  $\mu$ l of cells from the **NO DNA** tube to the **crtYB\* DNA** tube. Pipet up and down to mix.

12. Add 500  $\mu$ l of **EZ - Solution III** to each microfuge tube. Solution III will be goopy, but the amount you pipet does not need to be precise. Pipet up and down to mix, changing tips between tubes.

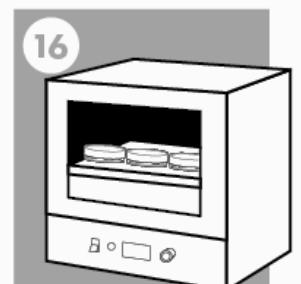
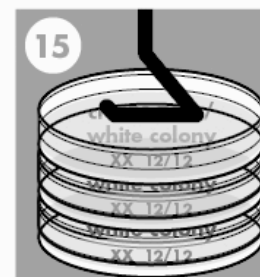
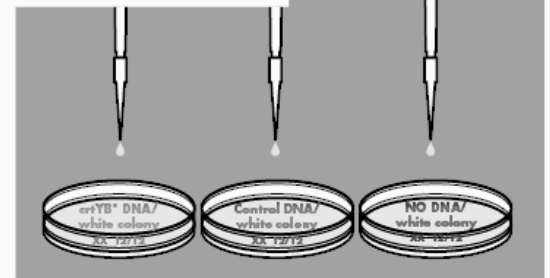
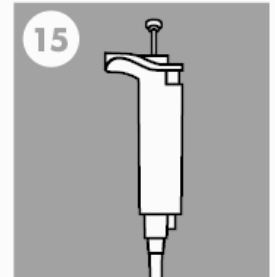
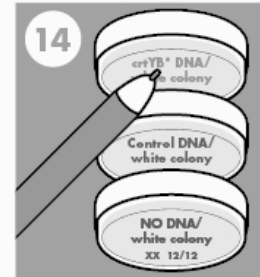
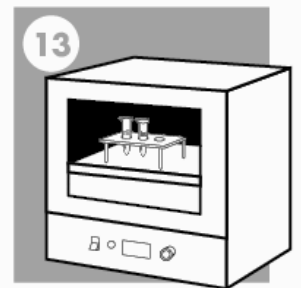
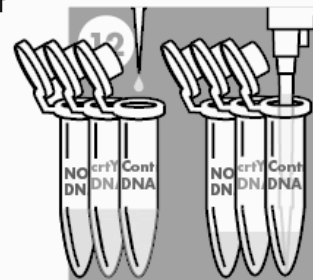
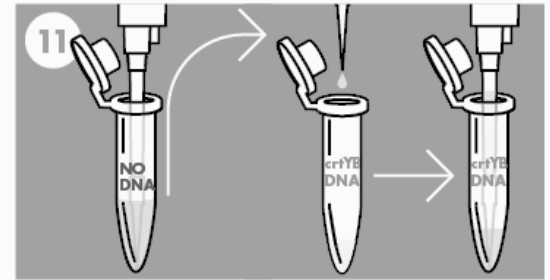
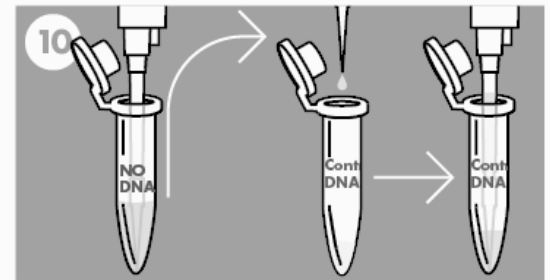
13. Incubate the tubes at 30° Celcius (C) for one hour. Periodically flick the tubes to mix during the incubation.

14. Label the media side of three SC-trp petri dishes as **NO DNA/white colony**, **Control DNA/white colony**, and **crtYB\* DNA/white colony**. Add your initials and today's date to each.

15. Pipet 200  $\mu$ l of each sample onto the media of the appropriate petri dish. Spread evenly across the dish with a sterile spreader. \*\* Discard spreader and microfuge tubes into the waste receptacle to be decontaminated.

16. Incubate petri dishes, media side up, for 2 days at 30° C.

After the petri dishes have incubated for 2 days, count the colonies of each color in every dish.



\*\* VIDEO OF PROCEDURE AVAILABLE ONLINE

# POST-LAB QUESTIONS

The negative control (no DNA) you ran was expected to have no colonies because the yeast cannot make their own tryptophan and the cells were grown on media that \_\_\_\_\_.

If you DO see colonies of yeast growing on the negative control -trp plate, you might think that \_\_\_\_\_.

If you accidentally plated the negative control (no DNA) on YPD, what would you expect to see?

If you see white colonies growing on the positive control, then you know that \_\_\_\_\_.

If you see orange colonies growing on the experimental petri dish, then you know that \_\_\_\_\_.

If you see orange and red and yellow and white colonies growing on the experimental petri dish, what can you conclude?

Can you think of other ways to improve the reliable performance of the system?

If you wanted to build a business around the health benefits of Golden Yeast, what would you need to do?

Would you eat bread made with Golden Yeast? Why or why not?



*Ideally, the interpretation of these results should encourage more experimentation, provide ideas for improved designs, and build excitement to explore and do more.*

STUDENT  
MANUAL

# BioBuilder

Do you have an idea for improving and extending the units? Please email us: [info@biobuilder.org](mailto:info@biobuilder.org)