

## How do specific genes in yeast contribute to flavor profiles?

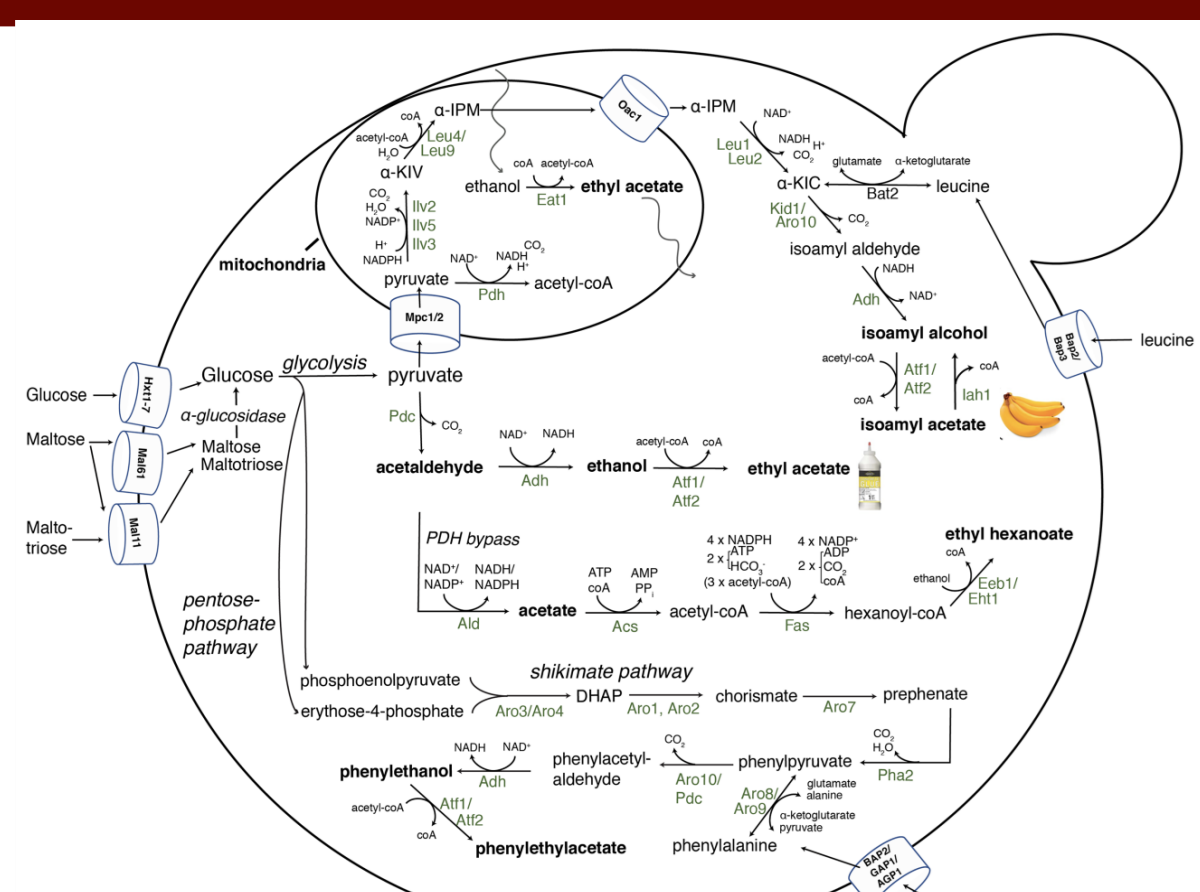


Figure 1: Genetic Pathways of Brewing Process

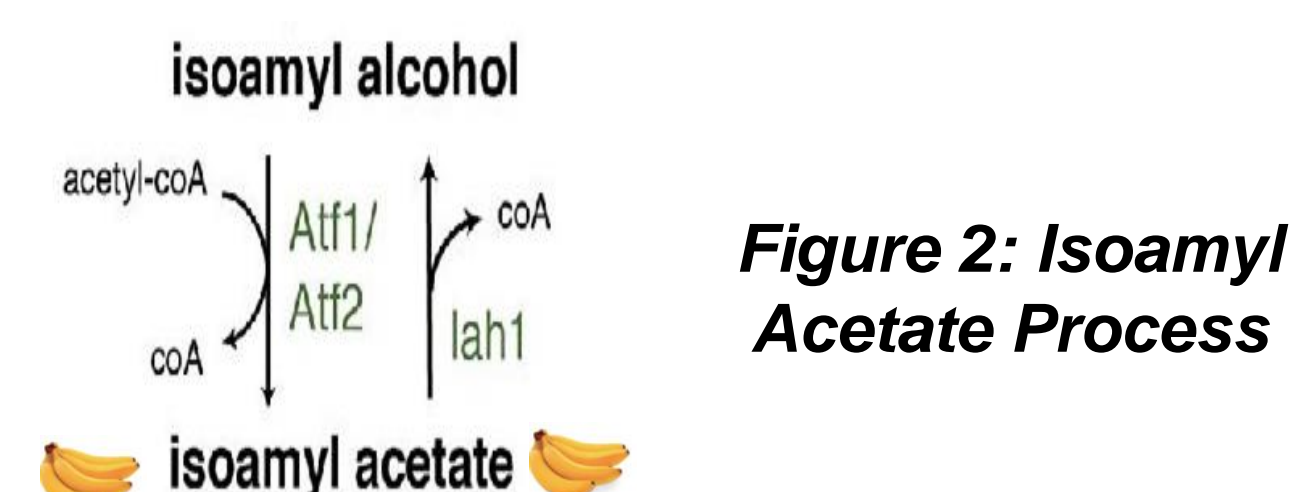


Figure 2: Isoamyl Acetate Process

- ❖ Took samples over the course of 4 days for 4 types of beer: Hefeweizen, Belgian Saison, California Ale, and San Diego Super to understand gene expression.
- ❖ Ran mRNA extractions to determine gene expression and an overhead GC to determine the isoamyl acetate levels.

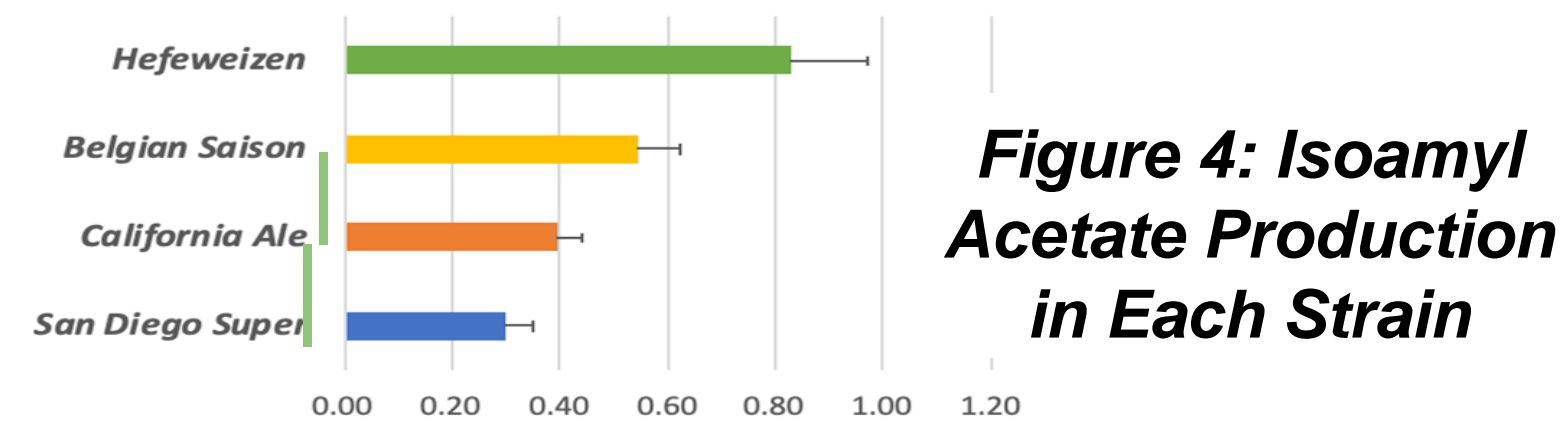


Figure 3: Beer Strains Flavor Profile

- ❖ The Hefeweizen strain has a significantly higher isoamyl acetate to isoamyl alcohol ratio
- ❖ mRNA expression fold calculated using  $\Delta\Delta Cq$  method.
- ❖ ACT1 was used as a reference gene compared to California Ale as the control strain.
- ❖ Figure 5: There was significant difference between Belgian Saison and every other type of beer.
- ❖ Figure 6: There was significant difference between beer types.
- ❖ Figure 7: There was significant difference between Belgian Saison and every other type of beer.

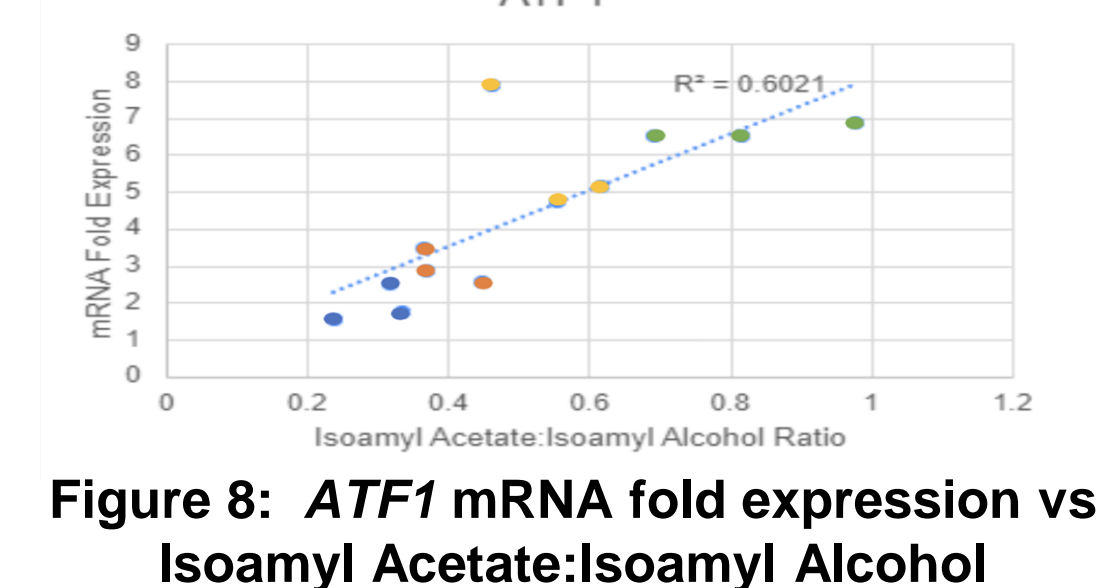


Figure 5: Relative ATF1 Gene expression

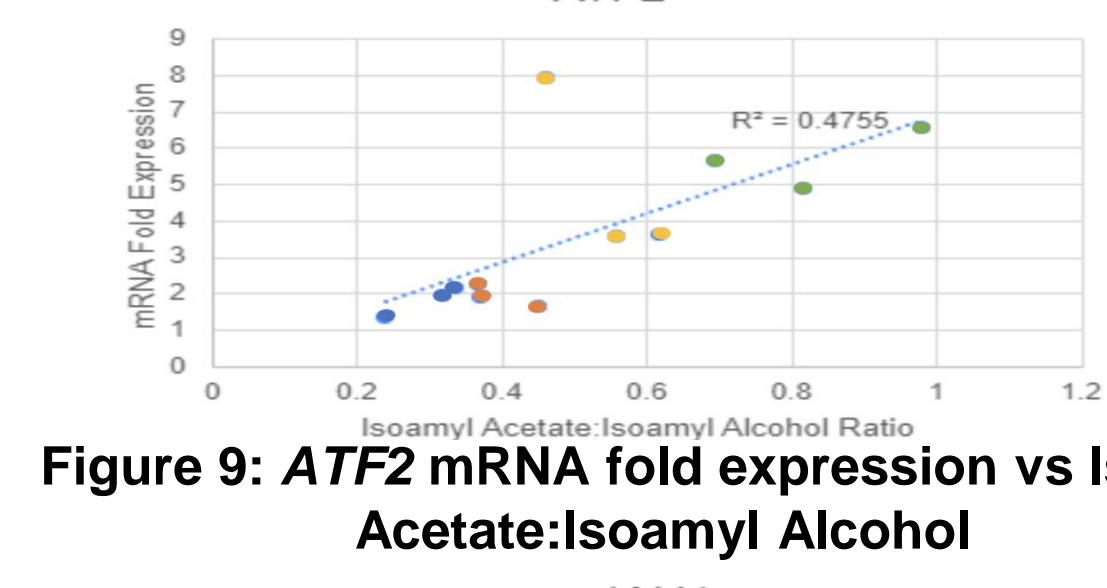


Figure 6: Relative ATF2 Gene expression

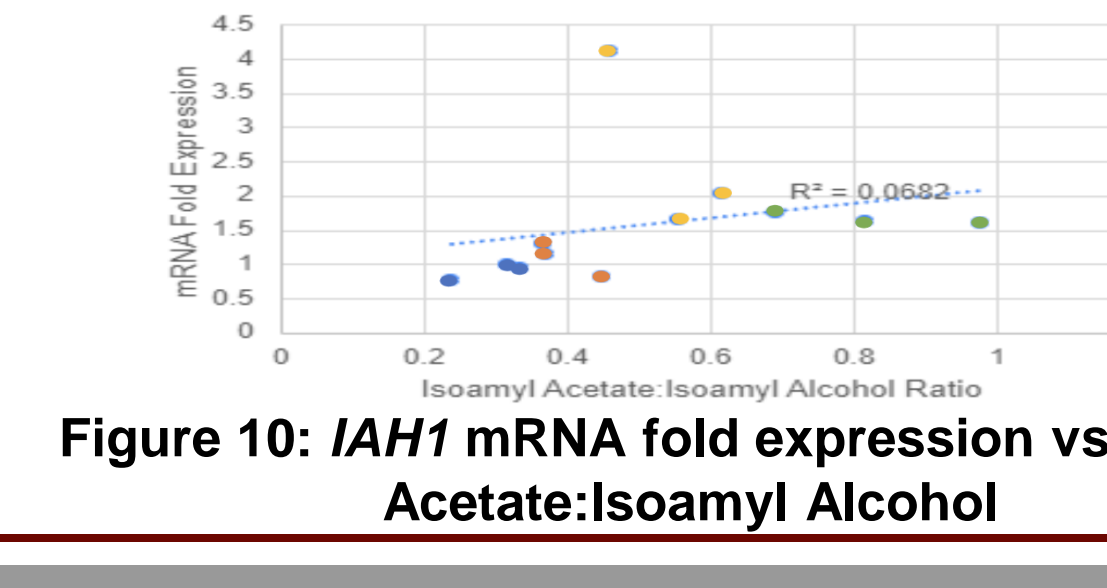


Figure 7: Relative IAH1 Gene expression



Figure 8: ATF1 mRNA fold expression vs Isoamyl Acetate:Isoamyl Alcohol

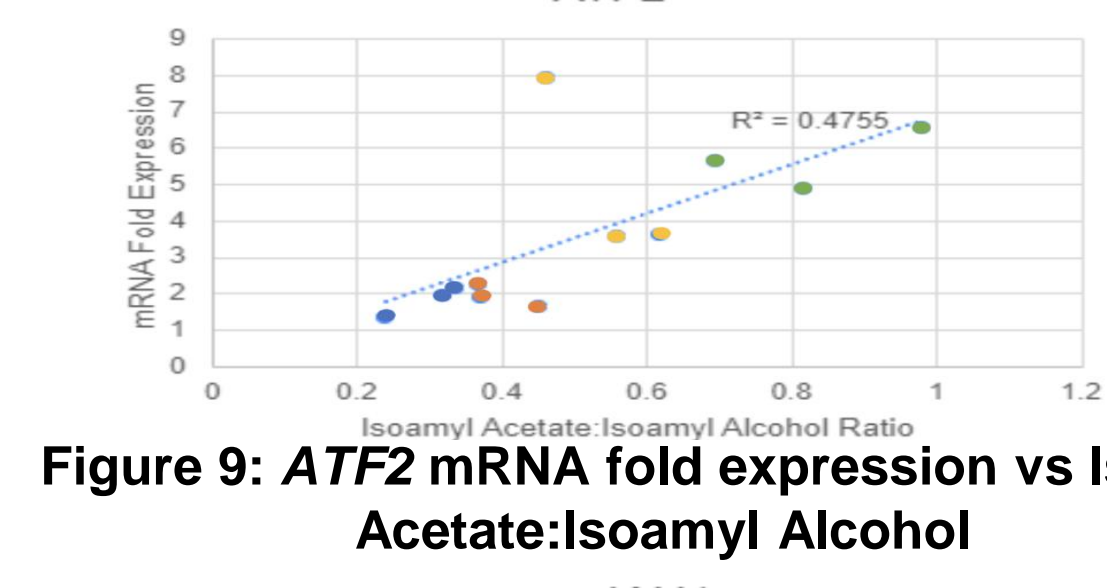


Figure 9: ATF2 mRNA fold expression vs Isoamyl Acetate:Isoamyl Alcohol

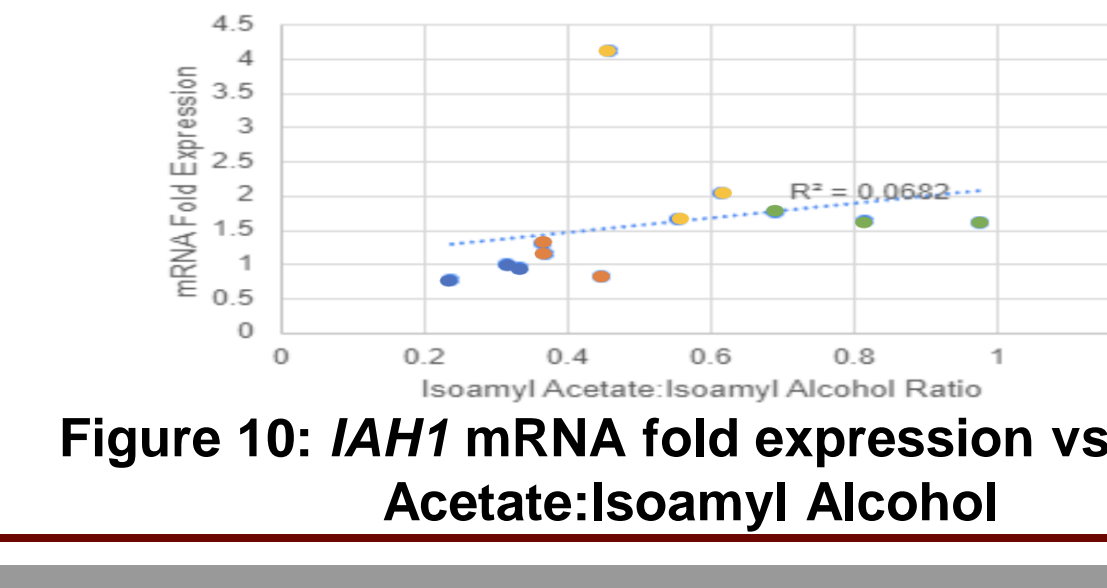


Figure 10: IAH1 mRNA fold expression vs Isoamyl Acetate:Isoamyl Alcohol

- ❖ The figures 8, 9, and 10 represent XY scatter plots comparing mRNA fold expression to isoamyl acetate:isoamyl alcohol ratio.
- ❖ There is not a strong linear correlation when plotting any individual gene's expression vs. isoamyl acetate:isoamyl alcohol ratio.
  - ATF1 does have the strongest correlation for any individual gene.

## How do ATF1, ATF2, and IAH1 work together?

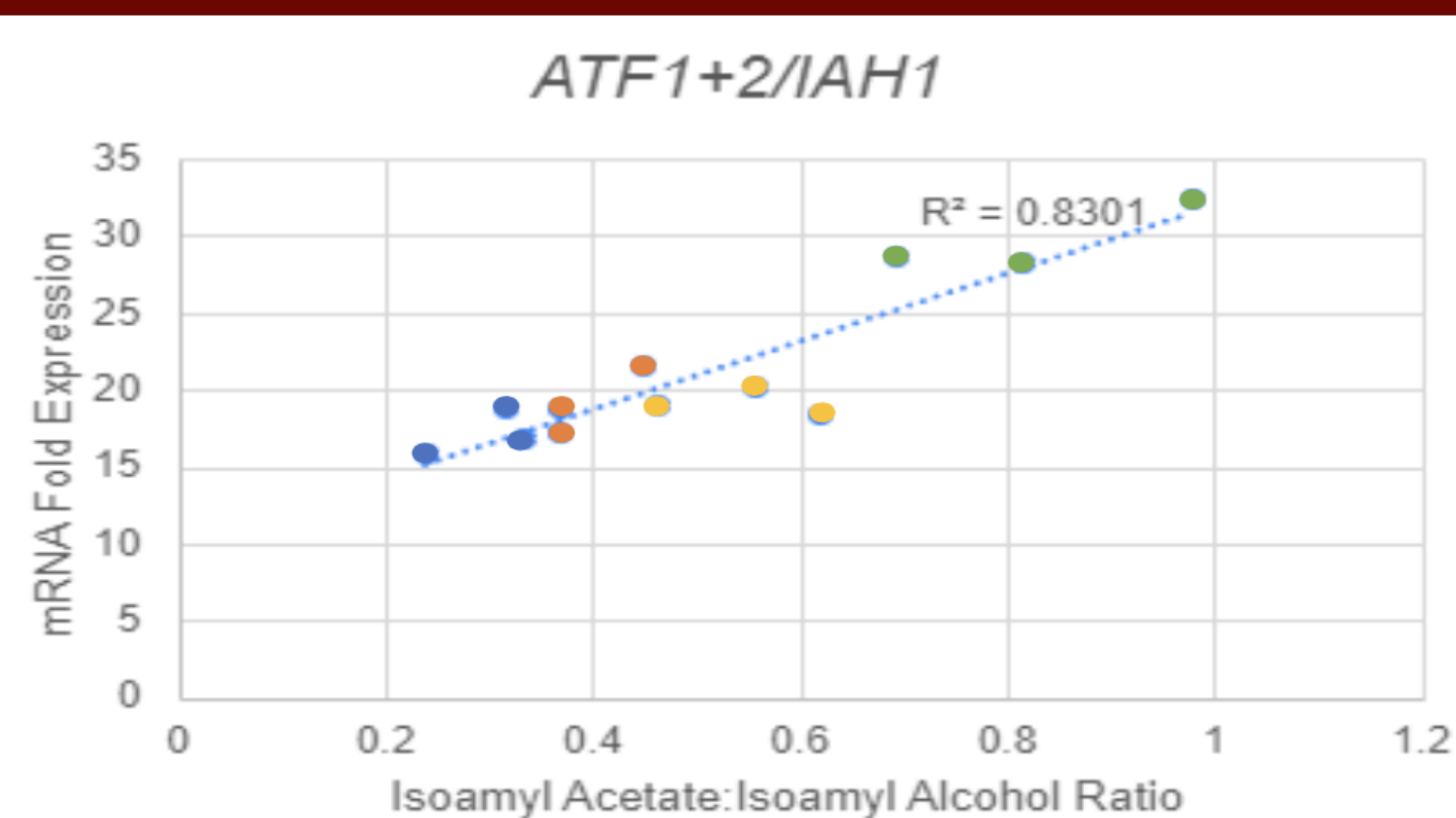


Figure 11: (ATF1+ATF2)/IAH1 mRNA fold expression vs Isoamyl Acetate:Isoamyl Alcohol

- ❖ When factoring in the balance of ATF1 & ATF2 driving the forward reaction, and IAH1 driving the reverse reaction, there is a strong linear correlation ( $r = 0.8301$ ) when comparing (ATF1+2):IAH1 ratio vs. isoamyl acetate:isoamyl alcohol ratio.
  - ATF1 has a relative good correlation of  $r=0.6021$ , but not as good as the gene fold expression vs isoamyl acetate:isoamyl alcohol ratio.

### This means.....

- 1) The expression of any one gene is not indicative of isoamyl acetate production.
- 2) The relative expression of all three genes factored together is indicative of isoamyl acetate production. There is a strong correlation between ATF1&ATF2:IAH1 expression and Isoamyl acetate production.

## Hypothesis:

*By lowering (or completely removing) IAH1 production, there will be an increase in isoamyl acetate.*

## Modifying the Genes

- ❖ Now knowing this gene ratio, we aim to modify isoamyl acetate production by genetically modifying IAH1.
- ❖ Goal: Increase the amount of isoamyl acetate in beer by knocking out IAH1 via the use of CRISPR-Cas9.

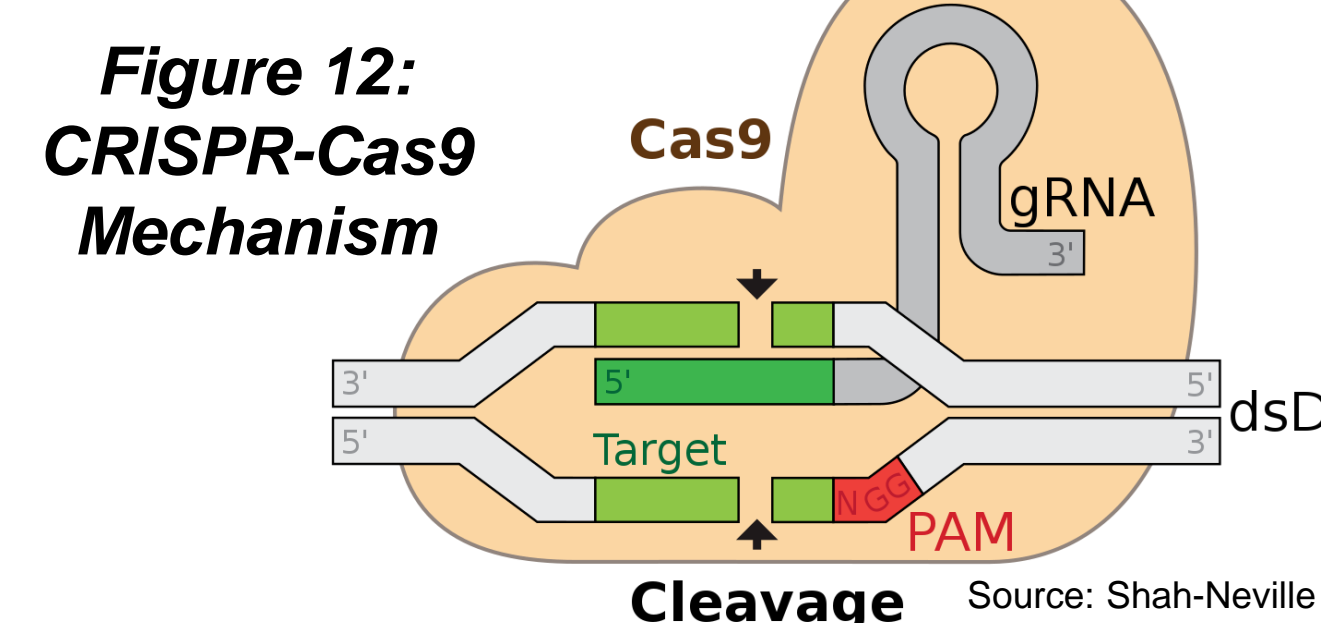
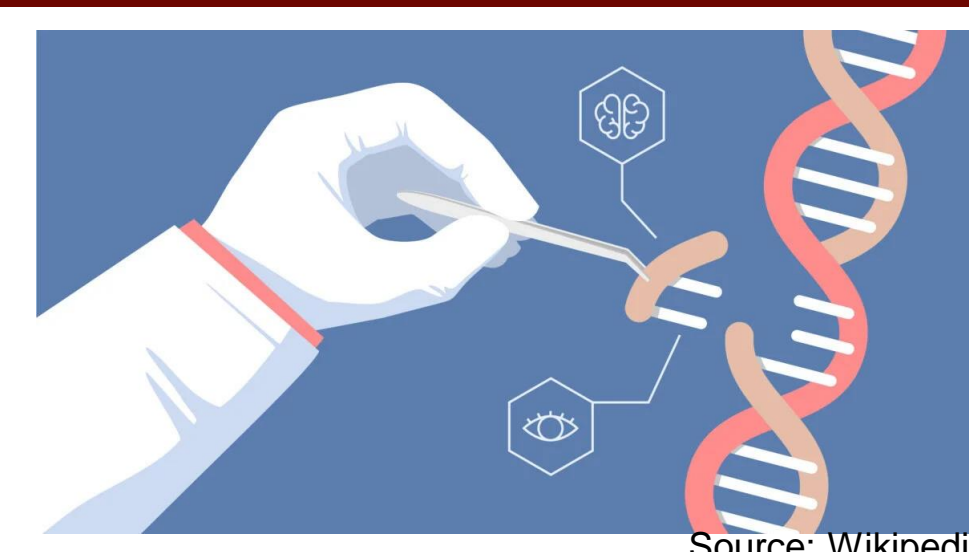


Figure 12: CRISPR-Cas9 Mechanism

- ❖ CRISPR-Cas9 is a novel engineering technique that is able to change genes on a base precision level.
- ❖ It is independent of copy number variations.
- ❖ It will insert a stop codon and an mRuby gene into the IAH1 gene preventing the translation.
- ❖ This will prevent the transformation of isoamyl acetate to isoamyl alcohol.

## Plasmid Design

- ❖ Step 1: Designing sgRNA site
  - Using E-Crisp, design the site that would be inserted into Cas9 in order to show the target site.
  - Using the web protocol, the sgRNA was designed to avoid the mutation found within the IAH1 sequence within our genome.
  - New primer was designed in order to work around the mutations found in:
    - 23T (ATT>ACT) missense mutation
    - A81A (GCC>GCT) silent mutation
    - K163N (AAA>AAT) missense mutation
  - Identified mRuby gene in one of the plasmids ordered in order to add to our sgRNA.
  - The following oligos were created:
    - Forward oligo: oJM120 – 5'-GACTTTGGCAAAGATCAGTATGCTCT-3'
    - Reverse oligo: oJM121 – 5'-AACAGAGCATACTGATCTTTGCCAA -3'
    - Annealed oligos
      - 5'-GACTTTGGCAAAGATCAGTATGCTCT-3'
      - 3'-AACCGTTTCTAGTCATACGAGACAAA-5'
  - The resulting sgRNA plasmid was added to benchling for the next steps.
  - Run a virtual digest to confirm and compare to actual digest.

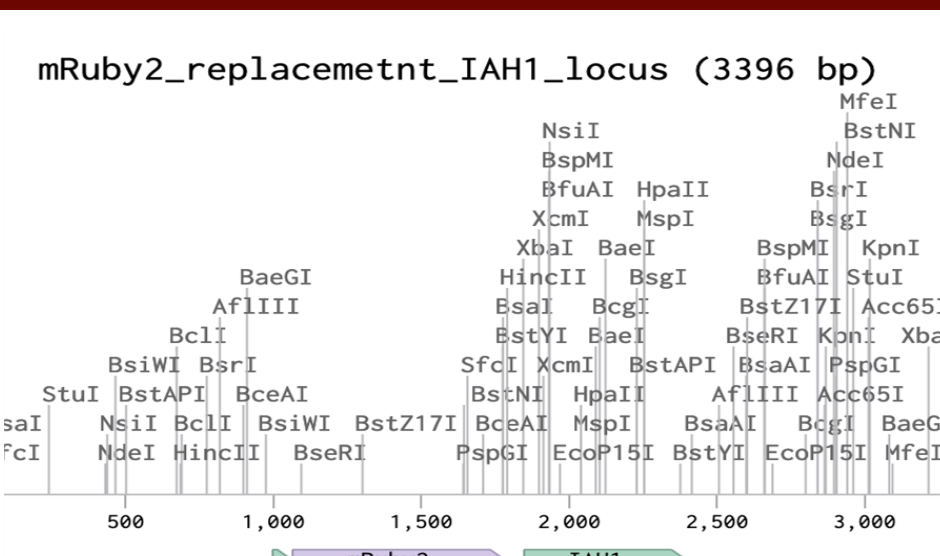


Figure 13: Plasmid for Insertion (linear)

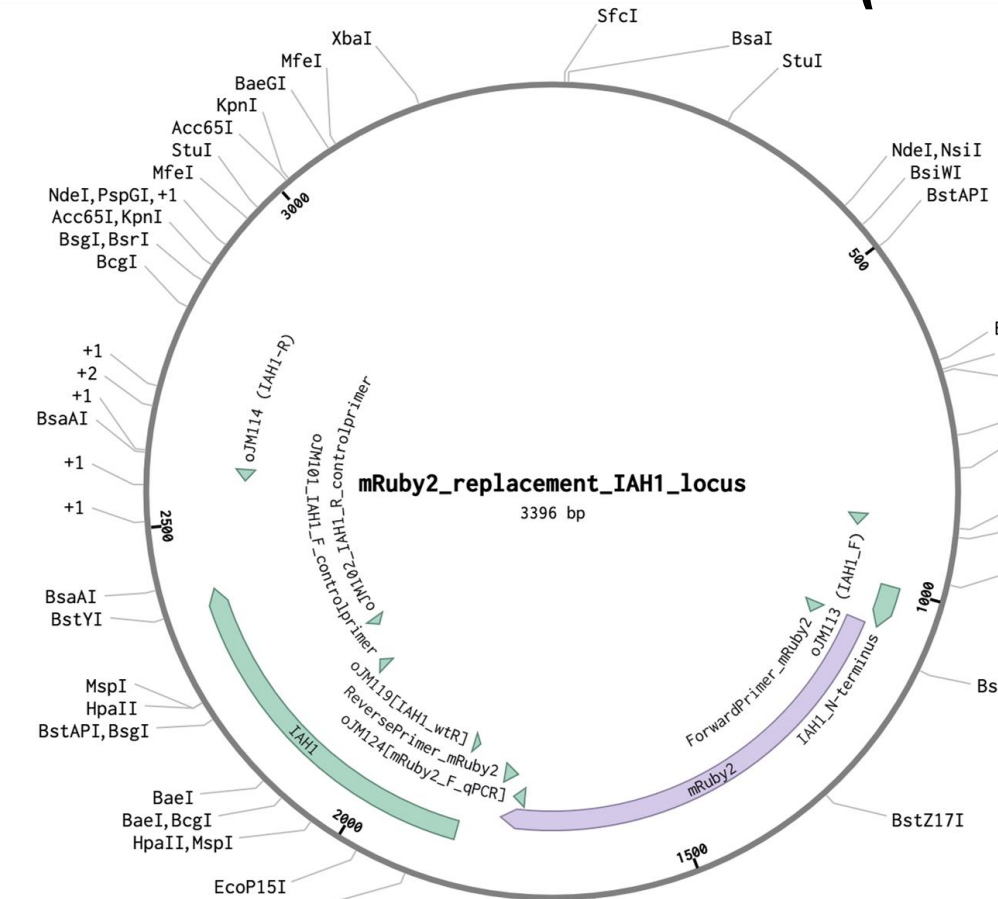


Figure 14: Plasmid for Insertion

## Creating Plasmid for Insertion

- ❖ Step 2: Prepping for CRISPR editing
  - Prepare plasmids:
    - pYTK050, pYTK003, pYTK068, pYTK095, pYTK-DN1, pYTK-DN2, pYTK-DN4, and pYTK-053
  - Prepare Plates and Media
    - Grow up E. Coli and
- ❖ Step 3: Anneal sgRNA Oligos
  - Prepare 100 uM stock of oligos in molecular grade ddH2O
  - Mix 10 uL of forward and 10 uL of reverse oligo in PCR tube.
  - Anneal the oligos using the thermal cycler under the following conditions
    - 95 C for 5 min
    - 55 C for 15 min
    - 25 C for 15 min
    - 4 C forever

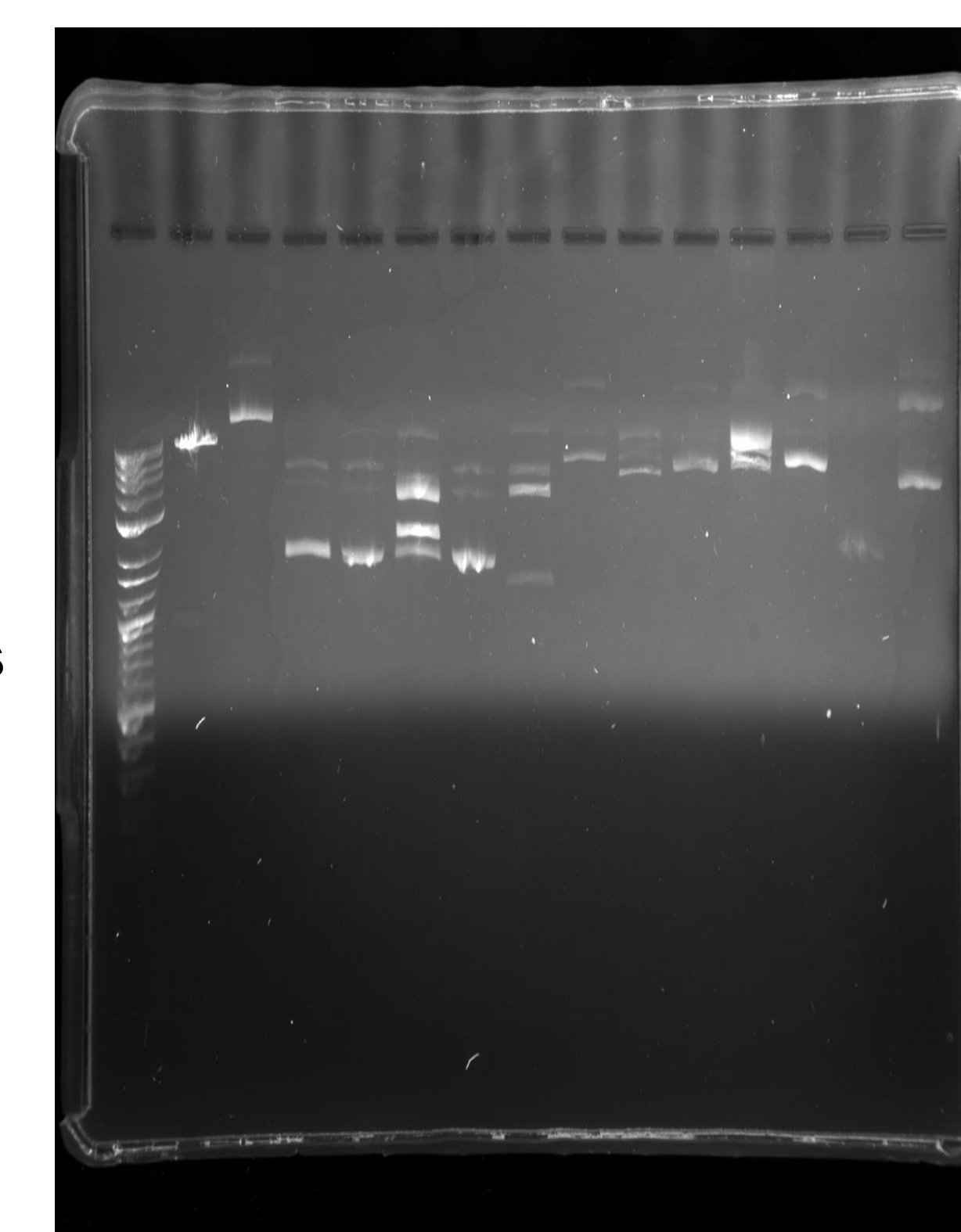


Figure 15: Confirmation of Plasmid on Gel

- ❖ Step 4: Run 1st golden gate reaction
  - Create 2 reaction mixes: a negative control and the golden gate.
    - Normal reaction has the sgRNA
  - In each one, add
    - 1 uL 10x T4 DNA ligase Buffer
    - 0.5 uL T7 DNA ligase
    - 0.5 uL BsmBI-V2
  - Ran in thermal cycler under the following conditions:
    - 42 C for 1 hr
    - 16 C for 1 hr \*(ligation step not included in NEB protocol for 1 insert, but can't hurt)
    - 60 C for 10 min
    - 80 C for 10 min
    - 4 C forever
  - Plate samples and preform extraction to see if the first editing work
  - Run a mini prep to confirm the knockout of GFP gene

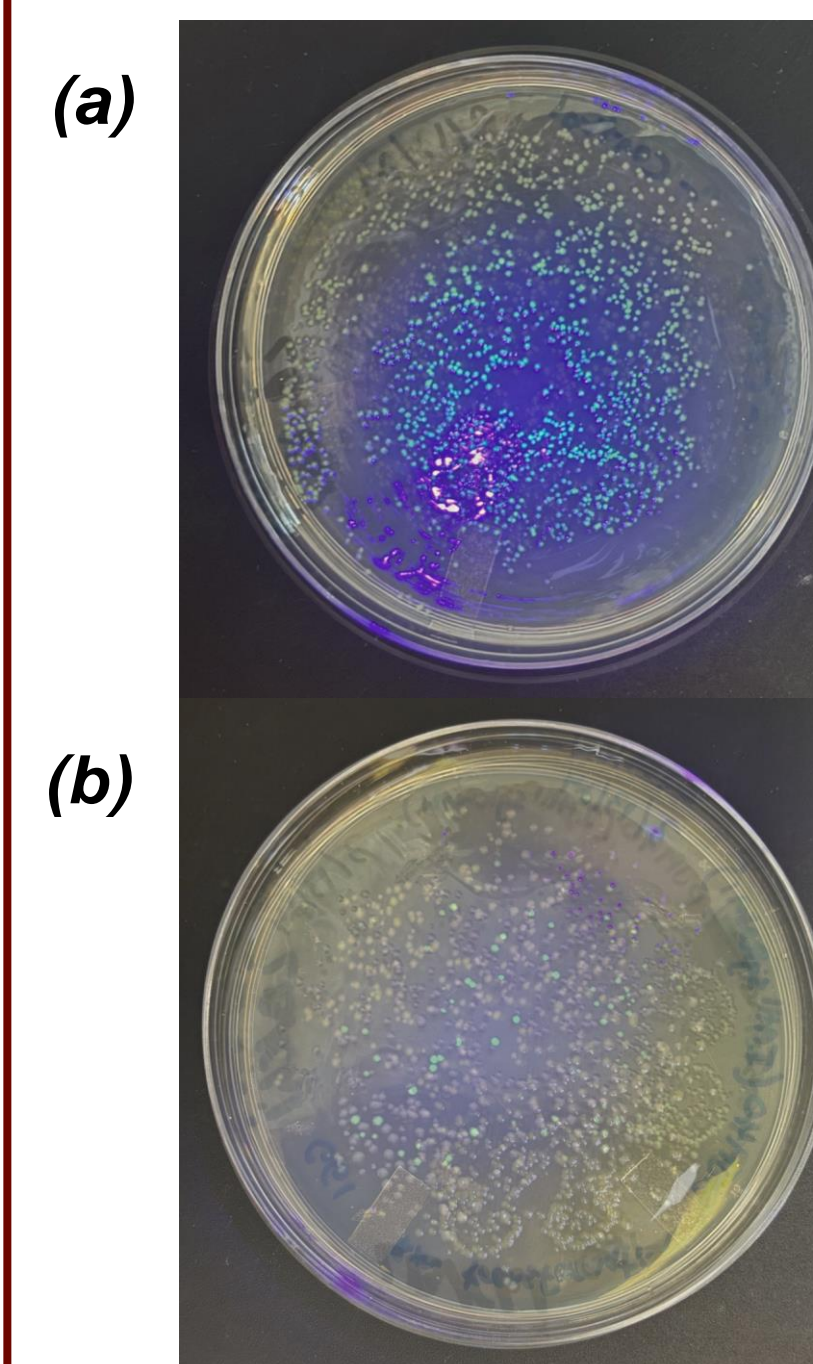


Figure 16: (a) negative control, (b) Golden Gate 1 Transformation product

## Future Steps

- ❖ Run second and third golden gate reactions to insert plasmid into the bacteria
- ❖ Transform the plasmid into the yeast after confirmation it works via plate
- ❖ Brew beer and test to see if the gene of interest was seen
- ❖ Test Isoamyl Acetate levels and gene expression

Figure 17: Overview of Transformation Steps

## Acknowledgements and Works Cited

Thank you to Jennifer Kennell of Vassar College for the help with the GC-MS and data collection. I would also like to thank Dr. Joost Monen and Dr. Ashley Stuart for their help with all of my research.

Thank you to The Ramapo College Foundation for funding the research



Scan above for references