

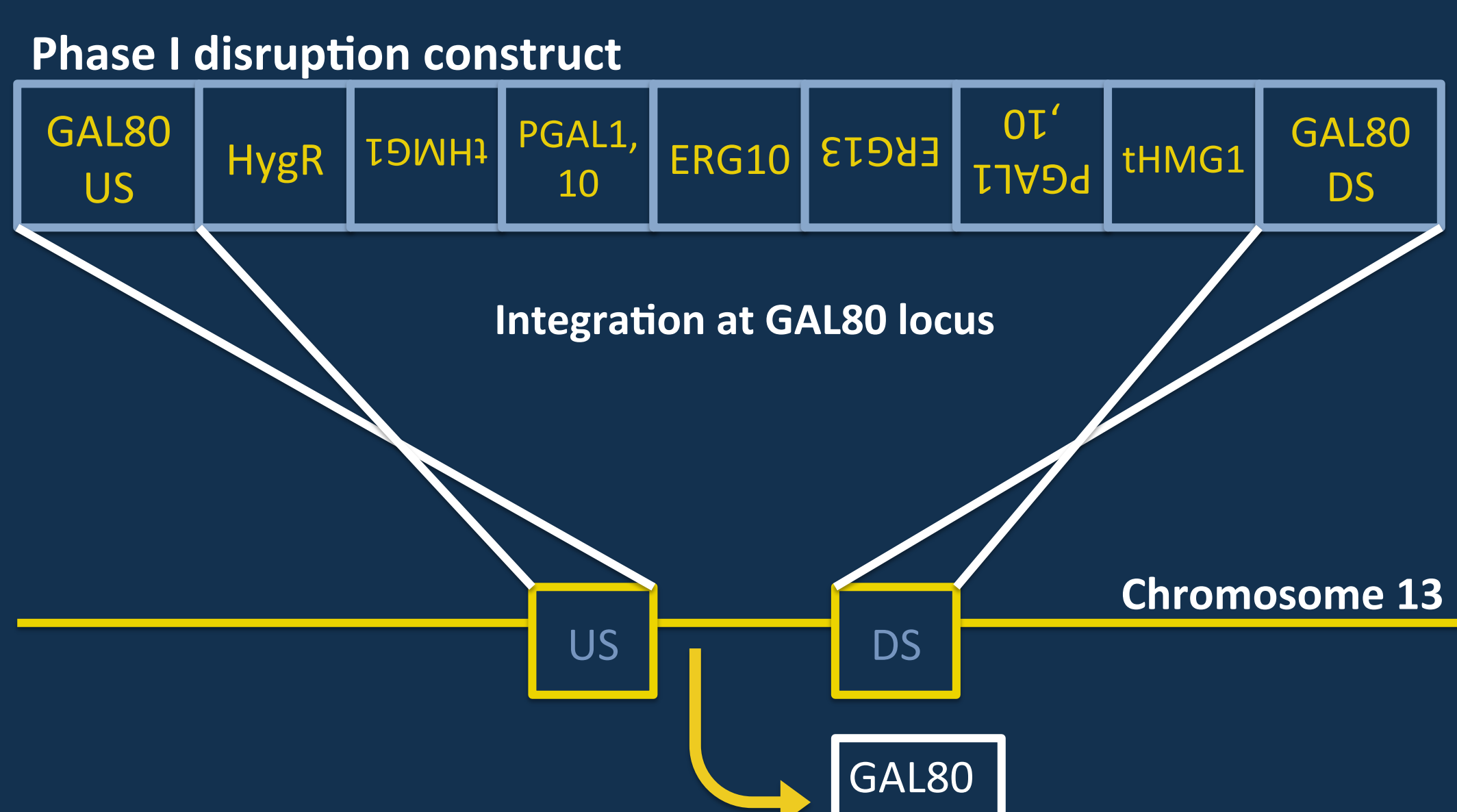
as a biojet fuel production in *Saccharomyces cerevisiae*

Metabolic engineering and synthetic biology

OBJECTIVES

- Learn how to modify *Saccharomyces cerevisiae* metabolic pathway and which strain is the best for the invention.
- Deal with molecular techniques and their applications.
- Learn how to design an expression plasmid.
- Evaluate new biofuel sources.

Disruption construct example



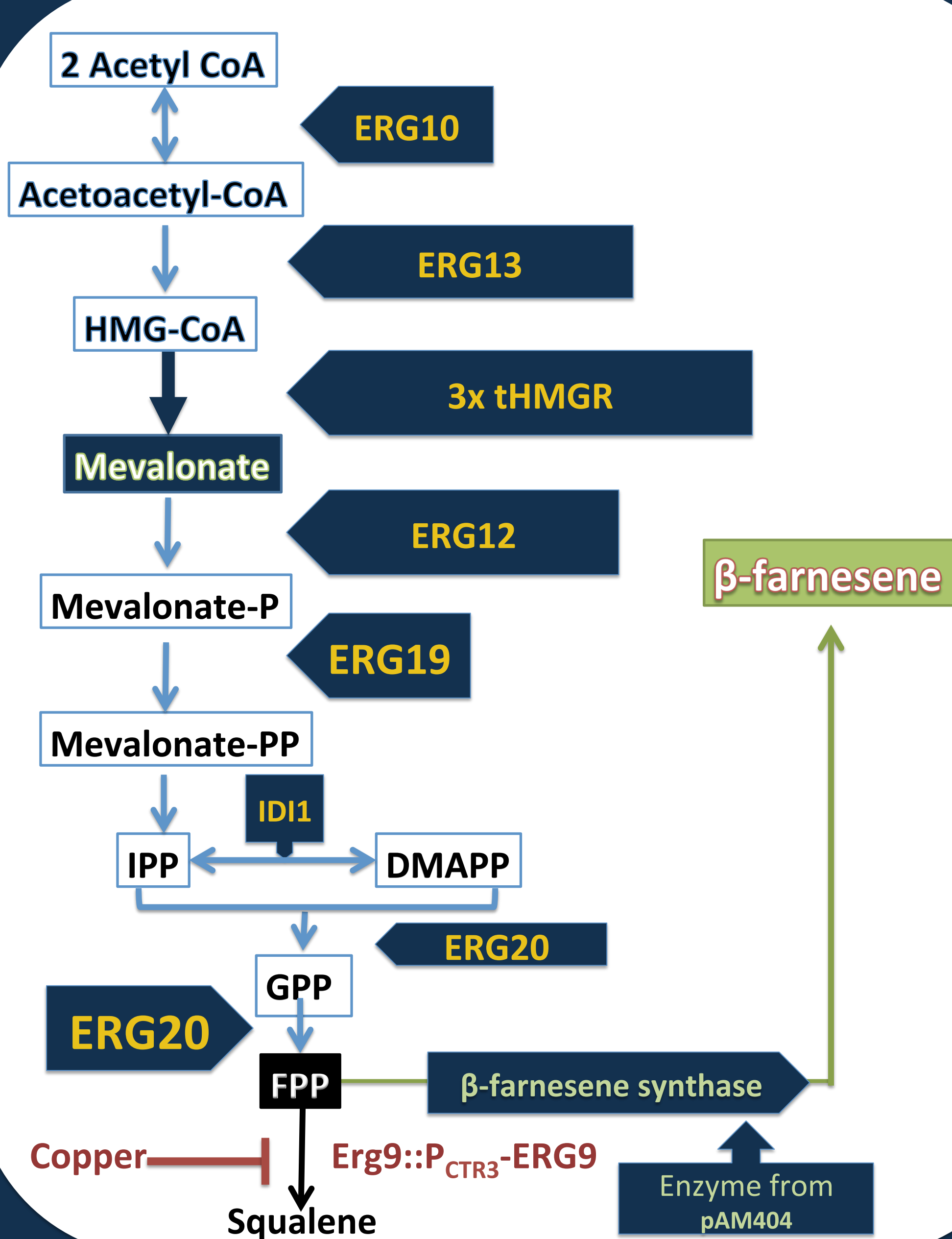
▲ **Figure 1.** Disruption construct cassette with promoters and homologous recombination in GAL80 locus example

Disruption cassette triple function:

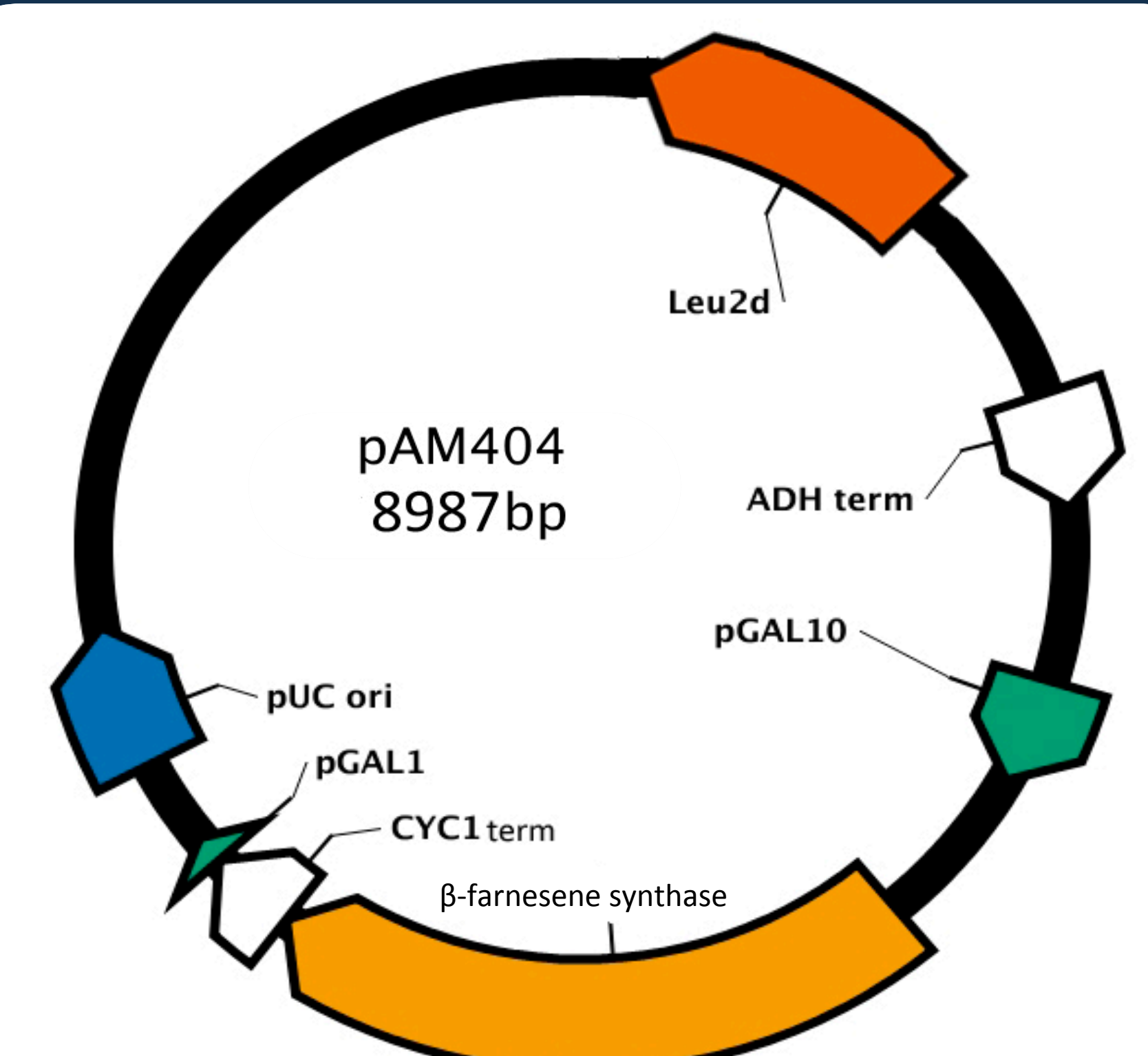
- 1. **Overexpressed mevalonate genes** by homologous recombination.
- ② Selection genes used: **hygA**, **natR**, **kanA (resistance)** and **URA3 (without it, no growth)**.
- ② **Promoters: pGAL1 ↔ pGAL10, pGAL4 (!), pGAL7.**
- 2. **Knock-out genes** by homologous recombination.
- ② **STE5** and **IME1**.
- 3. **Promoter substitution** for expression repression by homologous recombination.
- ② **ERG9 promoter** is substituted by a CTR3 promoter under Copper control.

Carried out techniques

Modified mevalonate pathway



▲ **Figure 2.** Metabolic pathway showing which genes have been modified to synthesize β -farnesene



▲ **Figure 3.** Overview of pAM404 and genes that makes up of it

Transformed plasmid

Inactivated genes by homologous recombination.

- STE5.** Mating – pheromone response gene. Without it, the possibility of sexual reproduction is 6 times less possible than with it (WT).

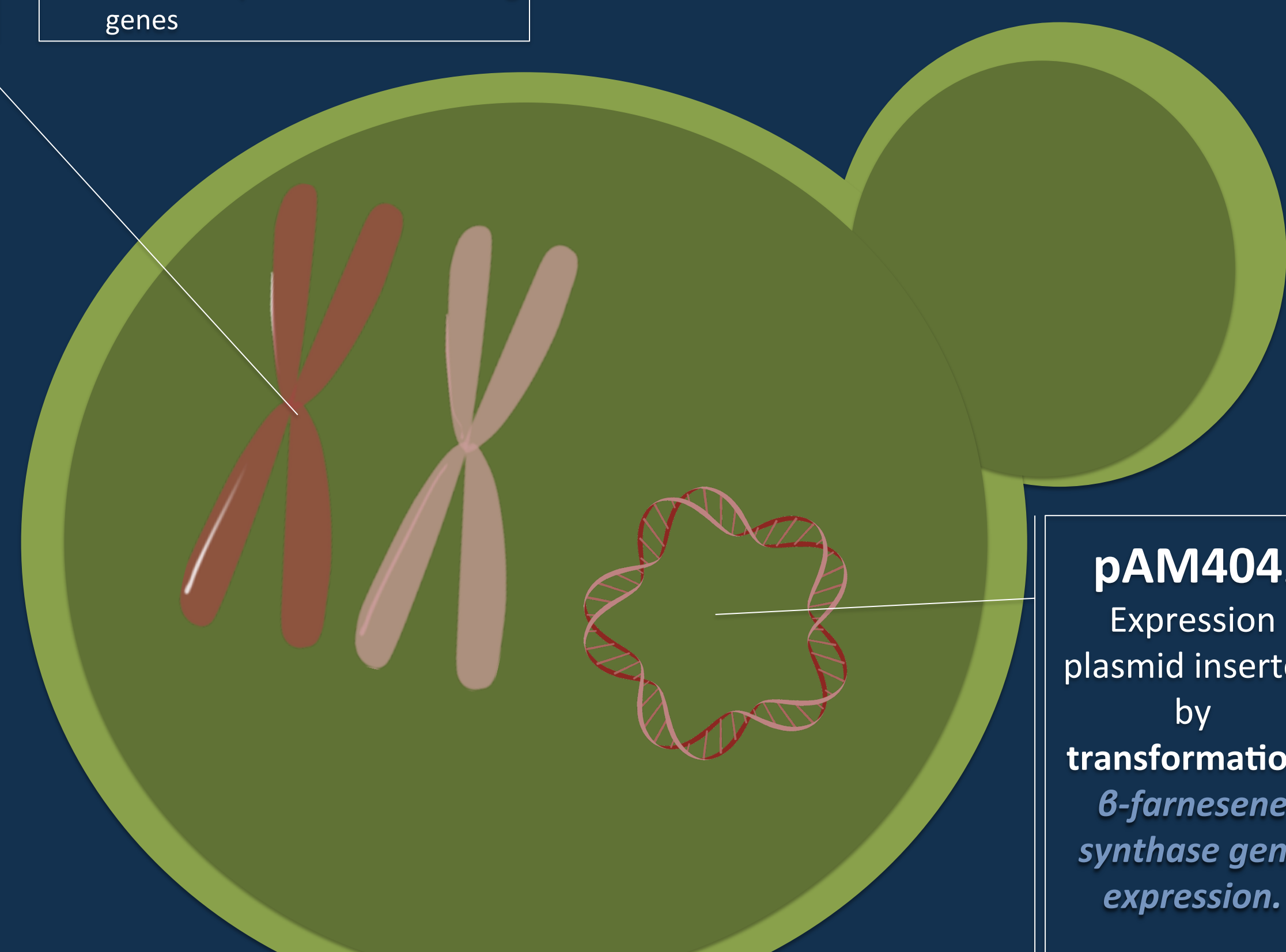
IME1. Meiosis transcriptional regulator

Diploid and an immutable strain.

Modified yeast used | Y1979

Modified genomic DNA.

- Modified mevalonate pathway
- Modified sporulation and mating genes



▲ **Figure 4.** Schematic figure of a *Saccharomyces cerevisiae* strain and what includes on it

pAM404.
Expression
plasmid inserted
by
transformation.
β-farnesene
synthase gene
expression.

Codon optimized for
Saccharomyces cerevisiae.

Under a very strong **pGAL10** promoter

Synthetically generated β -farnesene synthase gene from *Artemisia annua*.

Inserted into a pAM178,
yielding **pAM404**, using
restriction enzymes.

Leu2 gene as a selectable marker

Techniques used



Company and application

Conclusions

- ⊙ Part of the information taken in this project comes from patents. That is why there is a lack of information in different points.
- ⊙ There are different strategies for modifying *Saccharomyces cerevisiae*. The use of disruption cassettes involving more than one gene is innovative and different from what other research group have tested.
- ⊙ Farnesane production shows that yeasts, and in this case *Saccharomyces cerevisiae*, are a good platform for sesquiterpenoids production.
- ⊙ Farnesane can be one of the future biojet fuels available against the imminent lack of oil and petrol.

- [1] Mcphee, D. J., & Toomer, P. E. D., Stephen Renninger, N. (2010). Oakland, California, (US). Patent. Fuel compositions comprising farnesane and farnesane derivatives and method of making and using same. Patent No.: US 7846222 B2.
- [2] Ubersax, J.A., Emeryville, (2010), California, (US). Methods for generating a genetically modified microbe. Patent. Pub. No.: US 2010/0304490.
- [3] Nielsen J, Larsson C, van Maris A, Pronk J. Metabolic engineering of yeast for production of fuels and chemicals. Curr Opin Biotechnol [Internet]. Elsevier Ltd; 2013;24(3):398–404.
- [4] Asadollahi MA, Maury J, Schalk M, Clark A, Nielsen J. 2010. Enhancement of farnesyl diphosphate pool as direct precursor of sesquiterpenes through metabolic engineering of the mevalonate pathway in *Saccharomyces cerevisiae*. *Biotechnol Bioeng* 106:86-96.

Main references