# Disclosing the Causes Underlying Sterility in *Zygosaccharomyces*Allodiploid Yeast by NGS-assisted Approaches

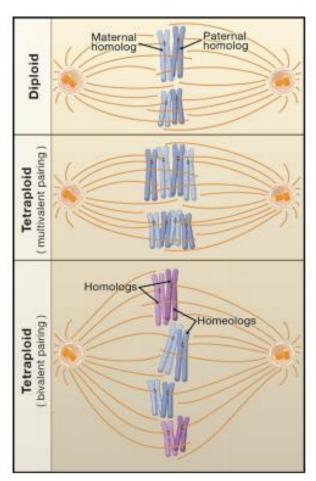
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# The Evolutionary Impact of Ploidy Variation in Eukaryotic Lineages



Adapted from Otto P. Sarah, 2007.

#### **Autopolyploidization** displays:

➤ Positive short-term and long-term effects:



Enrich organism with novel functions and beneficial properties

Deleterious effects:



Pre- and post-zygotic reproductive isolation of alloploid genomes

### Allodiploid offspring can have:

- Advantages since their increased genome size and complexity enhance heterosis and adaptive flexibility;
- Strong selective <u>disadvantages</u> due to their sterility.

# **Zygosaccharomyces rouxii** Clade: Multipurpose Salt-Tolerant Food Yeasts

### Microbial Cell-Factories

Candidate yeasts for producing glutaminase, chiral compounds, and heterologous proteins.

## Food-Starter and Spoilager

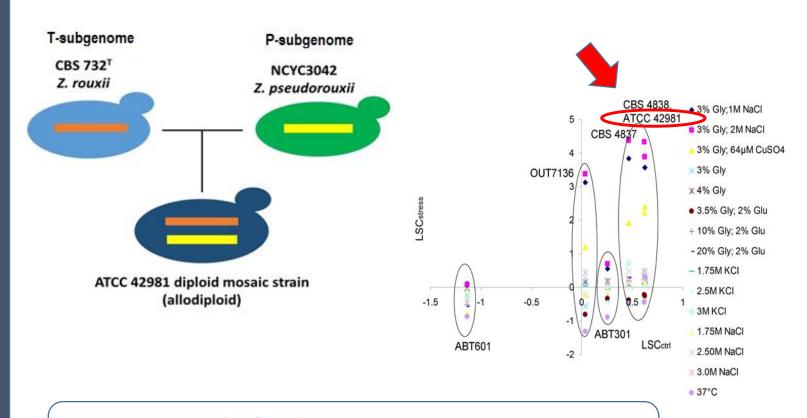
Starter for producing non-alcholic fruit-based beverages/condiments and vinegars.

Main responsible of food spoilage.

### **Genetic Engineering**

Source of genes for engineering the osmo/halo-tolerance in other organisms.

## Mosaic ATCC 42981 Strain: a Model for Gene-network Interplay and Adaptative Flexibility in Allodiploids



**Chimeric Mating System** 



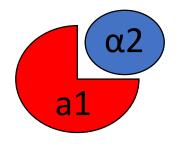
Two phylogenetically divergent
 MATa/MATα expression loci

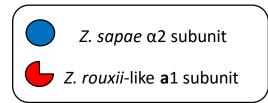
Despite its diploid DNA content ATCC 42981 is unable to undergo meiosis

Two copies of HO endonuclease gene

## Is the $a1/\alpha 2$ Heterodimer Functional?

a/α allodiploid ATCC 42981



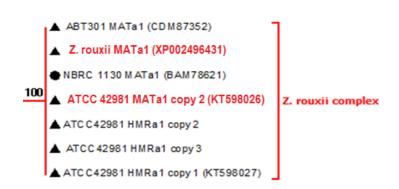




Master mating-type transcriptional factor.



Generation of negative epistasis contributing to allodiploid sterility and dysregulation of cell-type identity.

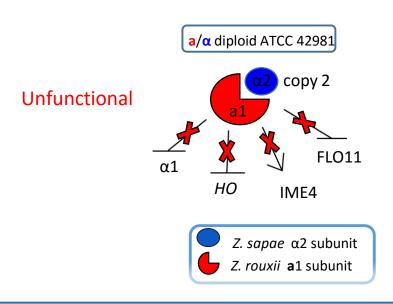


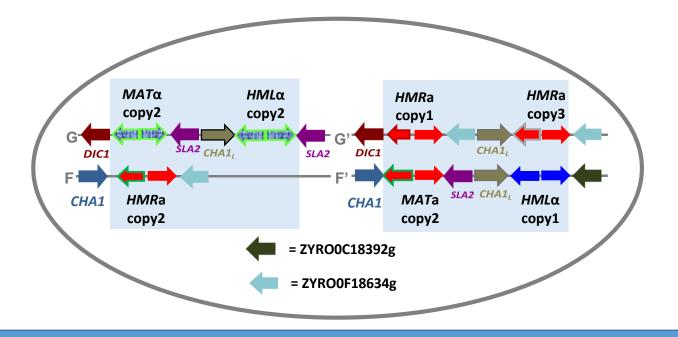
# Validation of our Model by Restoring a Functional a $1/\alpha$ 2 Dimer in ATCC 42981

<u>Hypothesis</u>: the chimeric a1-α2 heterodimer contributes to hamper the switching from mitosis to meiosis in our sterile allodiploid model.



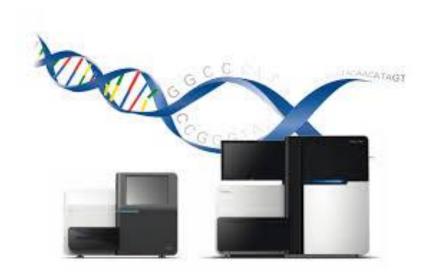
Verification: restoring a putatively functional heterodimer by genetic engeneering of ATCC 42981 cells.





# Three NGS approaches to unveil how introgressed parental genomes interplay at different levels

- 1. De-novo genome assembling to establish genome-wide patterns of introgression;
- 2. Transcriptome analysis of salt stress response;
- 3. Chip-Seq focused on a1- $\alpha$ 2 heterodimer target genes.



### ...and if you want to know more...





RESEARCH ARTICLE

Chimeric Sex-Determining Chromosomal Regions and Dysregulation of Cell-Type Identity in a Sterile *Zygosaccharomyces* Allodiploid Yeast

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