

# Synthetic biology approaches to design and construct microbial cell factories for the production of fructooligosaccharides

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# CONTEXT

- New products that improve the quality of life and reduce the risk of suffering from some diseases;
- Functional food: are **foods** that have a potentially **positive effect** on health **beyond basic nutrition**;



# CONTEXT

## Prebiotics

International Scientific Association for Probiotics and Prebiotics → **“a substrate that is selectively utilized by host microorganisms conferring a health benefit”**

### **A rising market!**

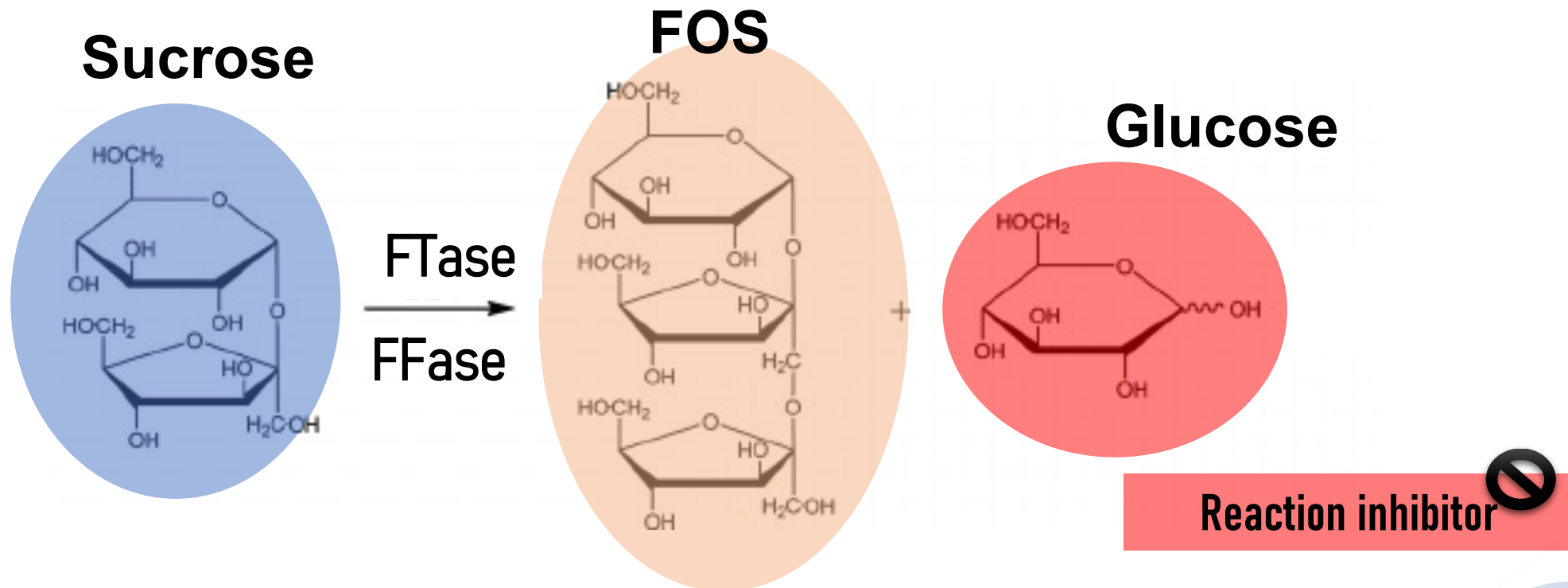
Expected to reach a value of  
**USD 7.37 billion by 2023**



# CONTEXT

## Fructooligosaccharides (FOS)

Non-digestible carbohydrates that are composed by fructose residues with a terminal glucose molecule residue linked by  $\beta(2\rightarrow1)$  glycosidic bonds



# CONTEXT

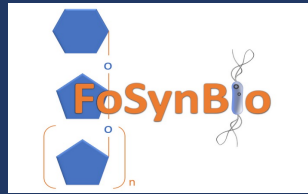
## Problems



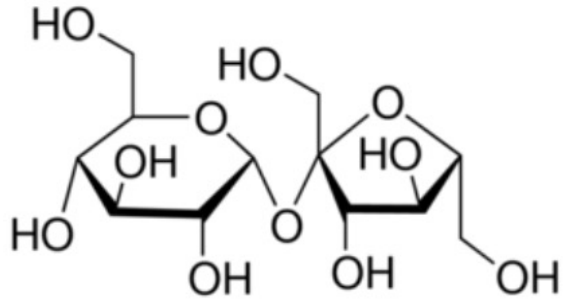
- Glucose inhibition;
- Presence of high concentrations of glucose, together with some sucrose and fructose, compromises the prebiotic effect;
- FOS purification steps → increase the production costs



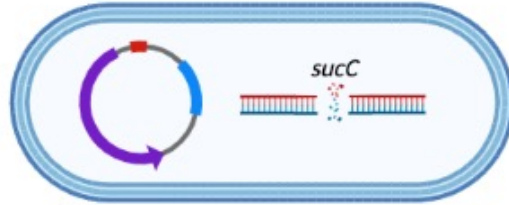
# INNOVATIVE SOLUTION



Sucrose



OR



*Zymomonas mobilis*



↑ FOS Purity

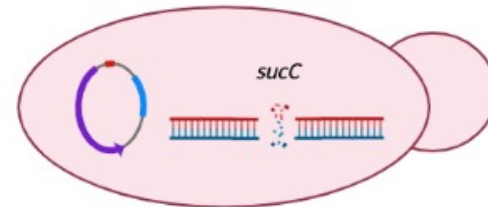
↓ Glucose + Fructose



**Final product with high prebiotic effect and commercial value**



Agri-food industry by-products



*Saccharomyces cerevisiae*



T1 - Design and construction of ZM and SC strains able to produce FOS



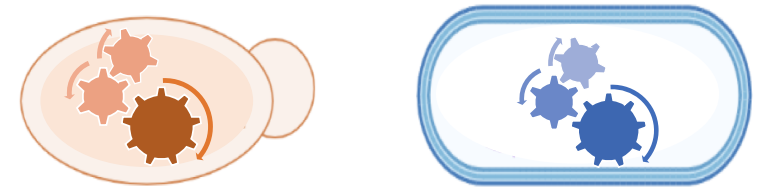
Mutant strains to prevent substrate consumption  
Overexpression of a Ftase/FFase from a recognized  
FOS producer

T3 - Design and optimization of the FOS production bioprocess



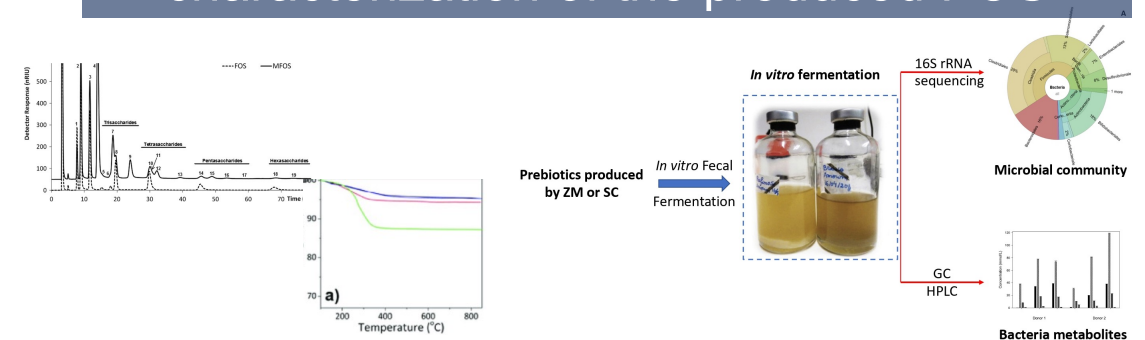
Agroindustrial by-products as alternative low-cost substrates  
Process scale-up

T2 - Genetic optimization of ZM and SC to increase the FOS yield and purity



Engineer the strains to consume the excess of glucose and fructose

T4 - Physicochemical and functional characterization of the produced FOS



Physicochemical and functional characterization  
Prebiotic potential → *In vitro* gut model

A scanning electron micrograph (SEM) showing a dense population of Zymomonas mobilis bacteria. The bacteria are rod-shaped, with some appearing as single cells and others as pairs or short chains. They have a textured, slightly irregular surface. The background is dark, making the lighter-colored bacteria stand out.

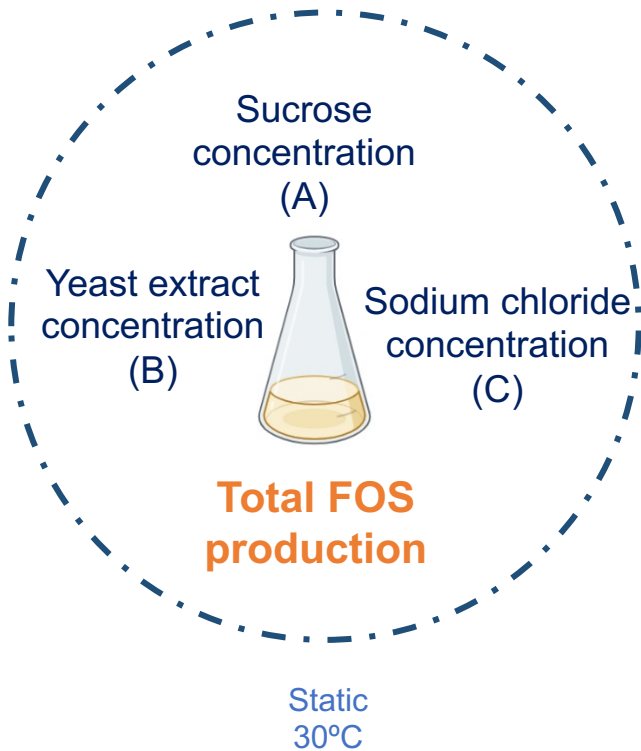
***In vivo* prebiotic production  
using *Zymomonas mobilis***



# In vivo prebiotic production using *Z. mobilis*

Box-Behnken  
(RSM)

15  
experiments

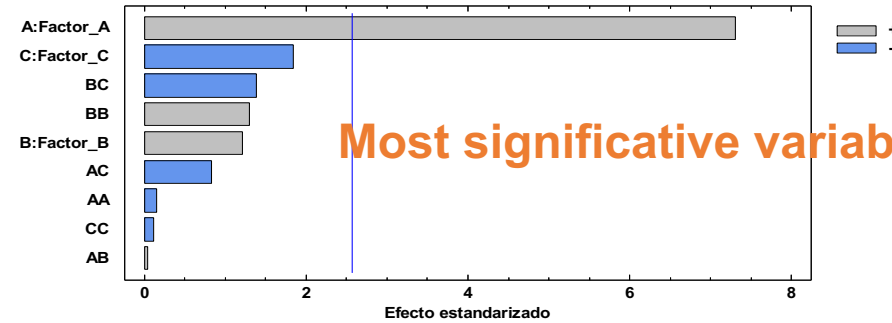


Total FOS  
51.6 g/L

Sucrose 350 g/L  
YE 20 g/L  
NaCl 2.5 g/L

ANOVA

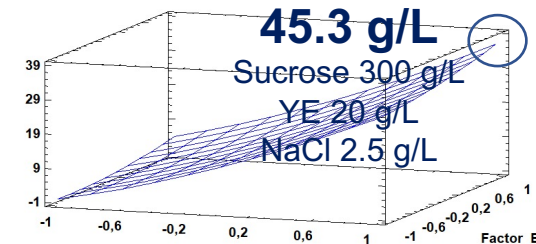
Diagrama de Pareto Estandarizada para Var\_1



Most significant variable- Sucrose

Model for predicting Total FOS production

Max. Total FOS

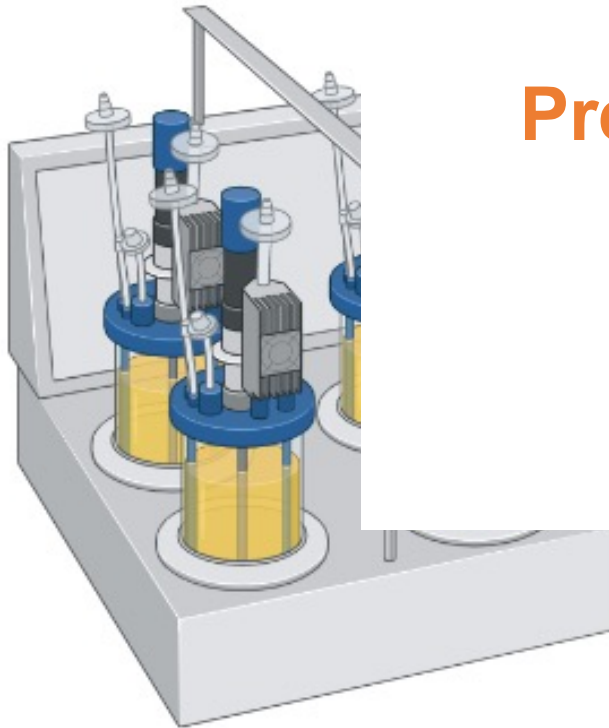


~ experimentally obtained

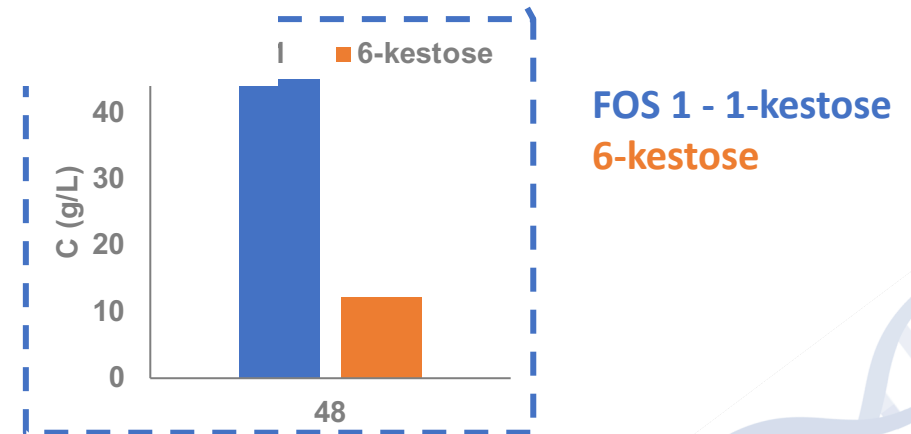
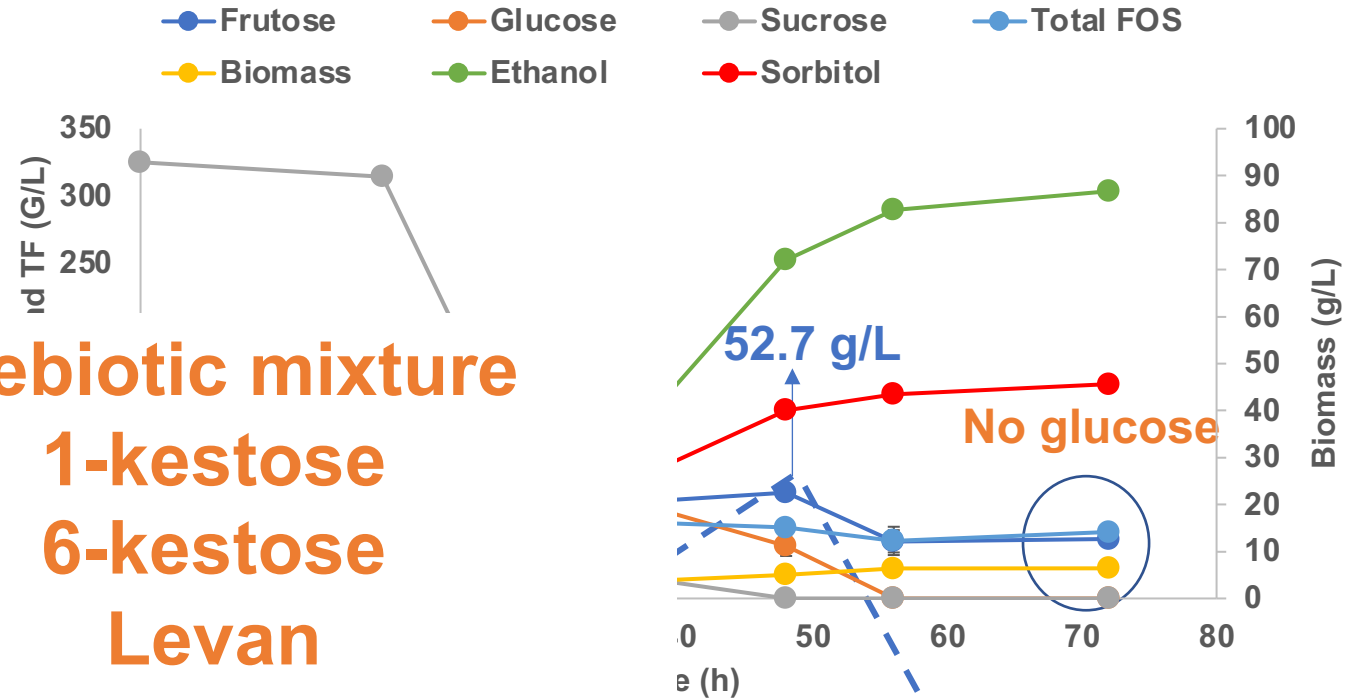
# In vivo prebiotic production using *Z. mobilis*

## Process scale up

Sucrose 350 g/L  
 YE 20 g/L  
 NaCl 2.5 g/L  
 Without air  
 Agitation 100 rpm



**Prebiotic mixture**  
 1-kestose  
 6-kestose  
 Levan  
 Sorbitol



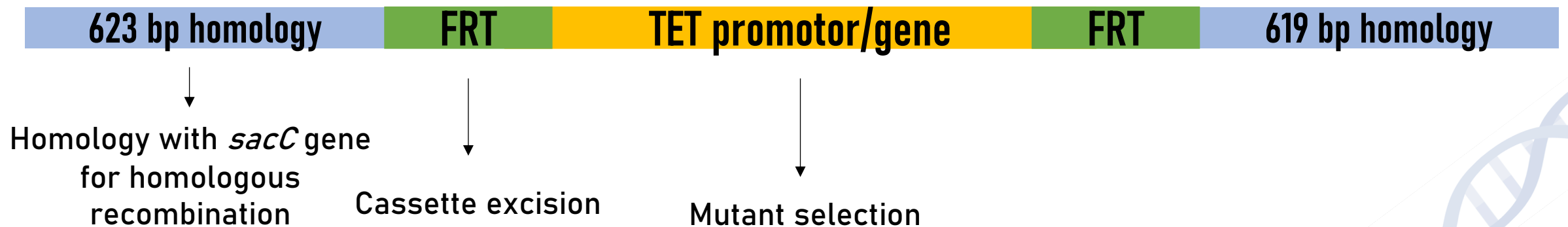
# *In vivo* prebiotic production using *Z. mobilis*

## Deletion of sucrase genes in *Z. mobilis*

**Purpose:** Prevent substrate consumption by the engineered *Z. mobilis*

**Objective:** Development of a sucrase deficient strain by *sacC* deletion

**How?** By Integration of a cassette carrying the Tet resistance gene and FRT sequence in *sacC* gene of *Z. mobilis* genome using homologous recombination



# *In vivo* prebiotic production using *Z. mobilis*

## Deletion of sucrase genes in *Z. mobilis*

### Cassette excision

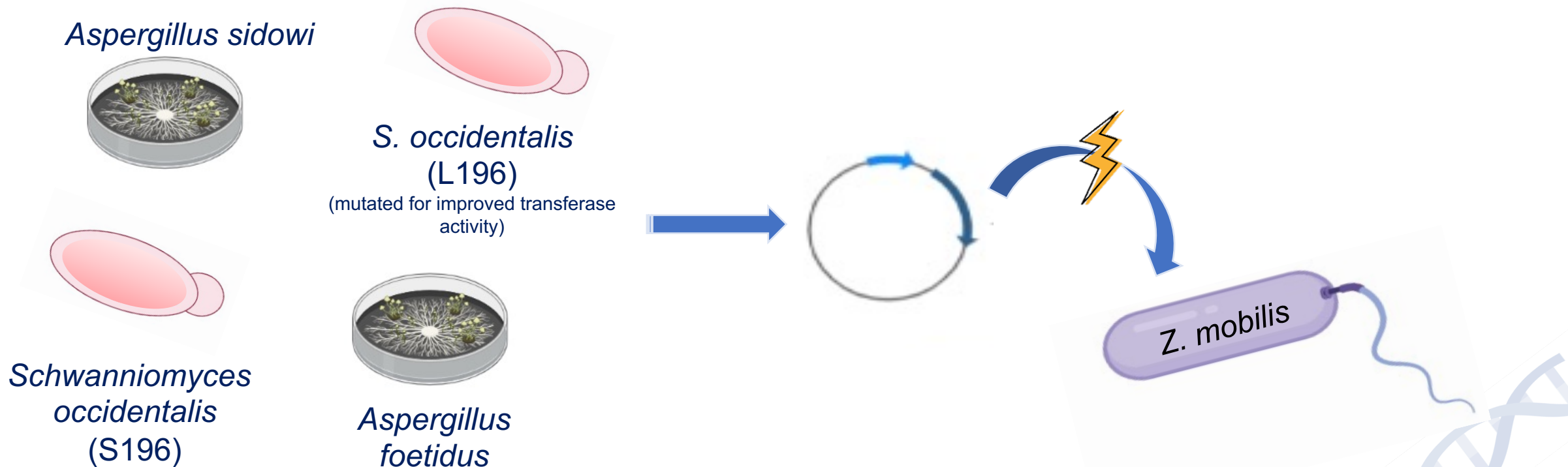


**GRAS status**



# *In vivo* prebiotic production using *Z. mobilis*

## Overexpression of a Ftase/FFase from a recognized FOS producer

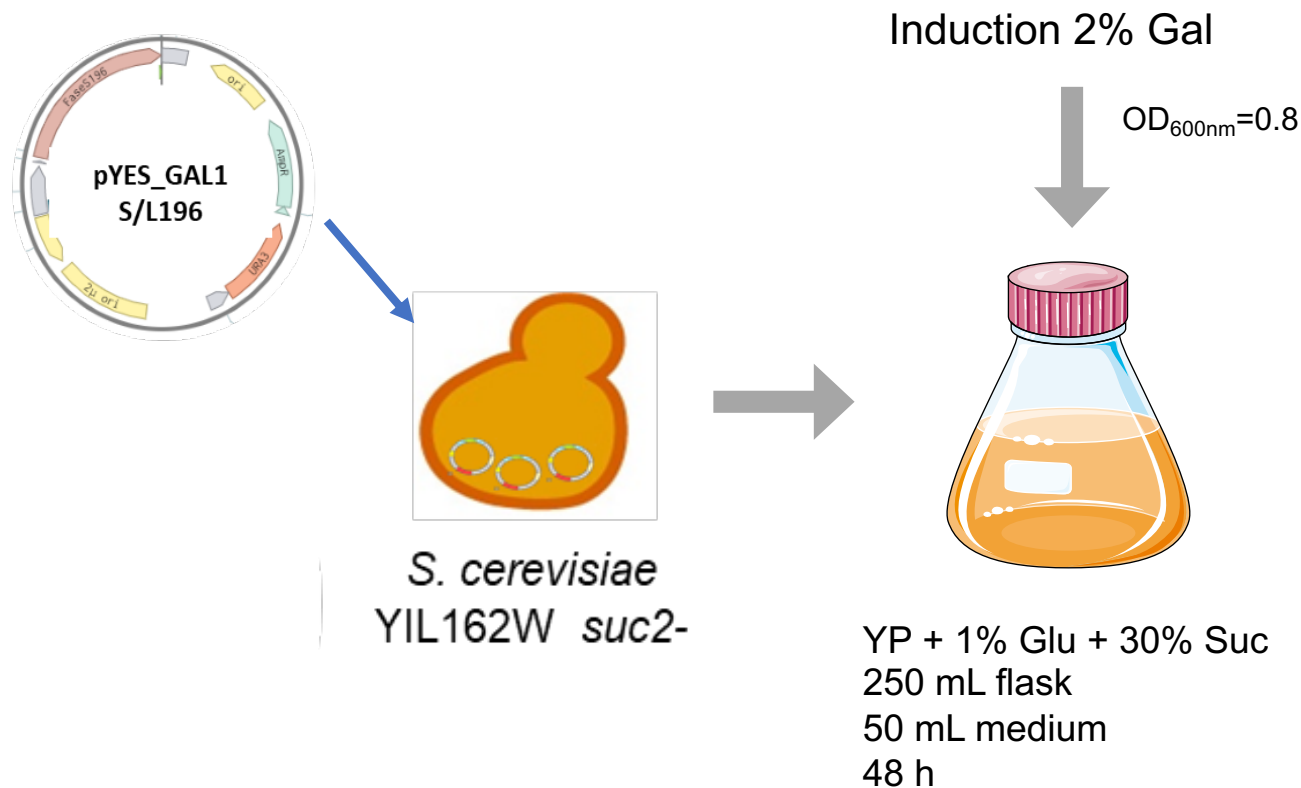


A microscopic view of numerous yeast cells, likely *Saccharomyces cerevisiae*, showing their characteristic oval shape and budding structures. The cells are densely packed and appear in various stages of growth and division.

***In vivo* FOS production using  
*Saccharomyces cerevisiae***

# *In vivo* FOS production using *S. cerevisiae*

Construction of *S. cerevisiae* Suc2<sup>-</sup> mutant harboring the pYES\_SoS196 or pYES\_SoSL196 (frutosyltransferase gene from *Schwanniomyces occidentalis* )



**S196: Higher sucrose hydrolysis**  
**L196: Higher FOS production**

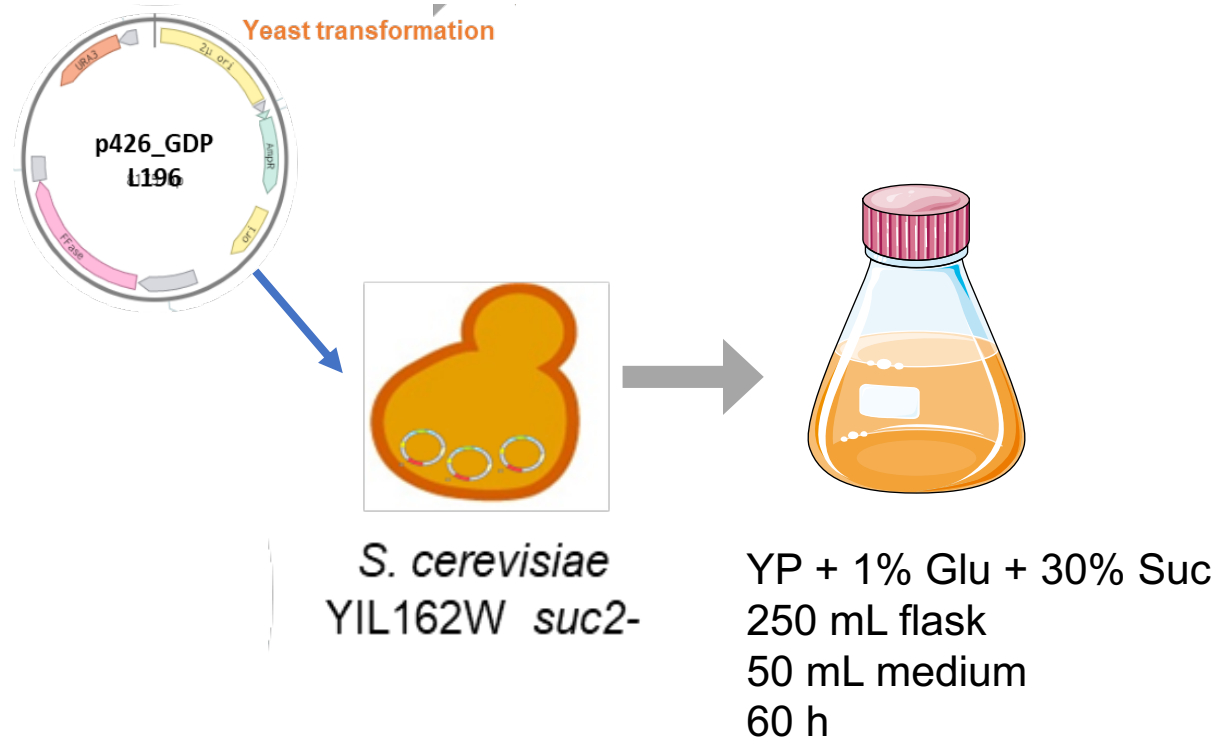
**Max. FOS production at 12 h after induction:**

**S196  $Y_{FOS/Sac} = 54 \pm 3$  mg/g**

**L196  $Y_{FOS/Sac} = 269 \pm 12$  mg/g**

# *In vivo* FOS production using *S. cerevisiae*

**GOAL:** Transform the SoL196 gene on p426\_GPD → Constitutive promotor (production cost reduction)

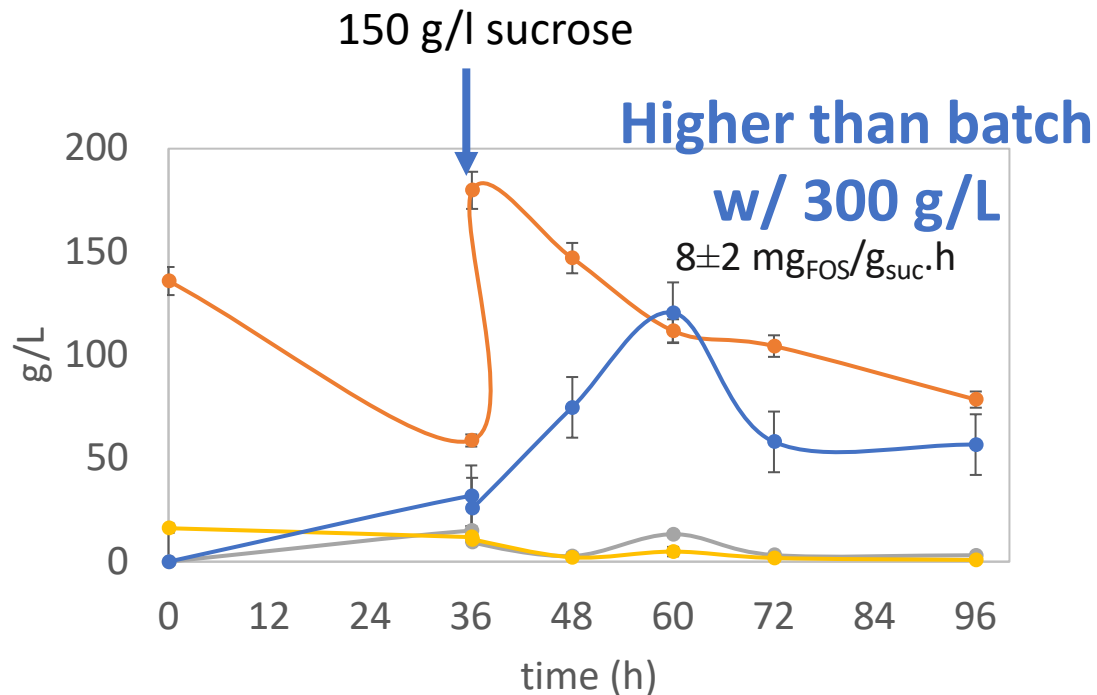


	GAL1 L196	GPD L196
FOS (g/L)	78 ±4	61±4
Maximum extracellular sucrolytic activity (U/mL)	6±0,5	8±1
Monosaccharides (g/L)	51±7	6±1
Ethanol (g/L)	17.5±1	13.0±0.9
Glycerol (g/L)	4±1	3.8±0.5
Productivity <sub>max</sub> (mg/g.h)	9.58±0.08	5.0±0.1



# In vivo FOS production using *S. cerevisiae*

## Effect of sucrose concentration



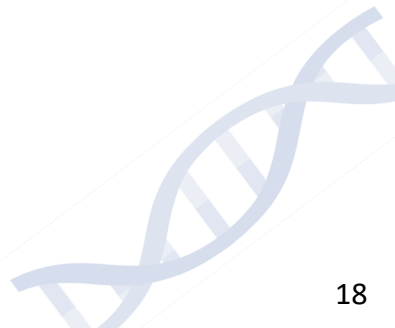
Maximum productivity and respective optimal time achieved during sucrose fermentation using *S. cerevisiae* GPD L196 clone.

Sucrose (g/L)	Productivity <sub>max</sub> (mg/g.h)	Optimal time (h)
120	4,6±0,4	36
210	5,1±0,2	48
300	5,9±0,3	48
390	5,8±0,3	60

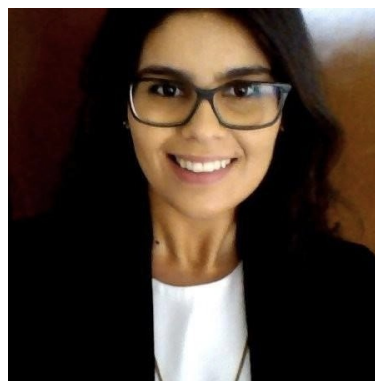
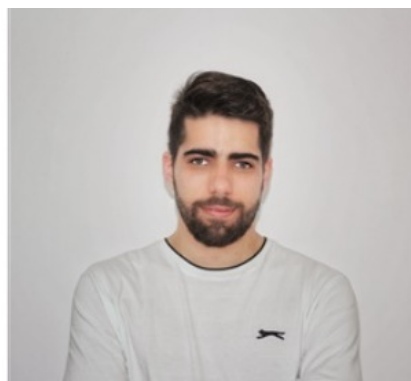
## Step-wise fed-batch

# Perspectives and Final Remarks

- New prebiotic “mix” → Prebiotic potential
- Incorporation of the produced prebiotic mixture in food matrix or its production and further commercialization as a prebiotic “mix”
- Conversion of industrial by-products (waste) and renewable raw materials into added value food ingredients (prebiotics) → EU Green Deal



# Acknowledgements

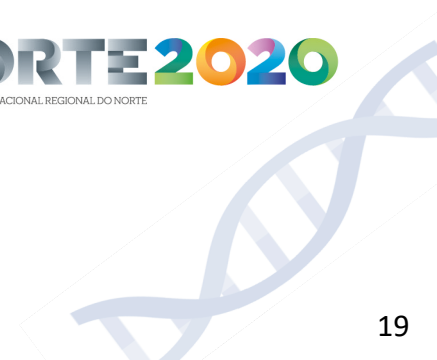


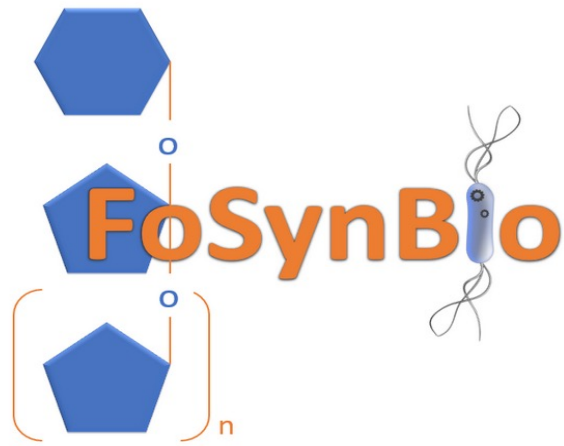
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para a Ciência  
e a Tecnologia

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2020

**PORTUGAL**  
2020

**NORTE2020**  
PROGRAMA OPERACIONAL REGIONAL DO NORTE





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