

Supplementary Material

Successive Fermentation of Aguamiel and Molasses by *Aspergillus oryzae* and *Saccharomyces cerevisiae* to Obtain High Purity Fructooligosaccharide

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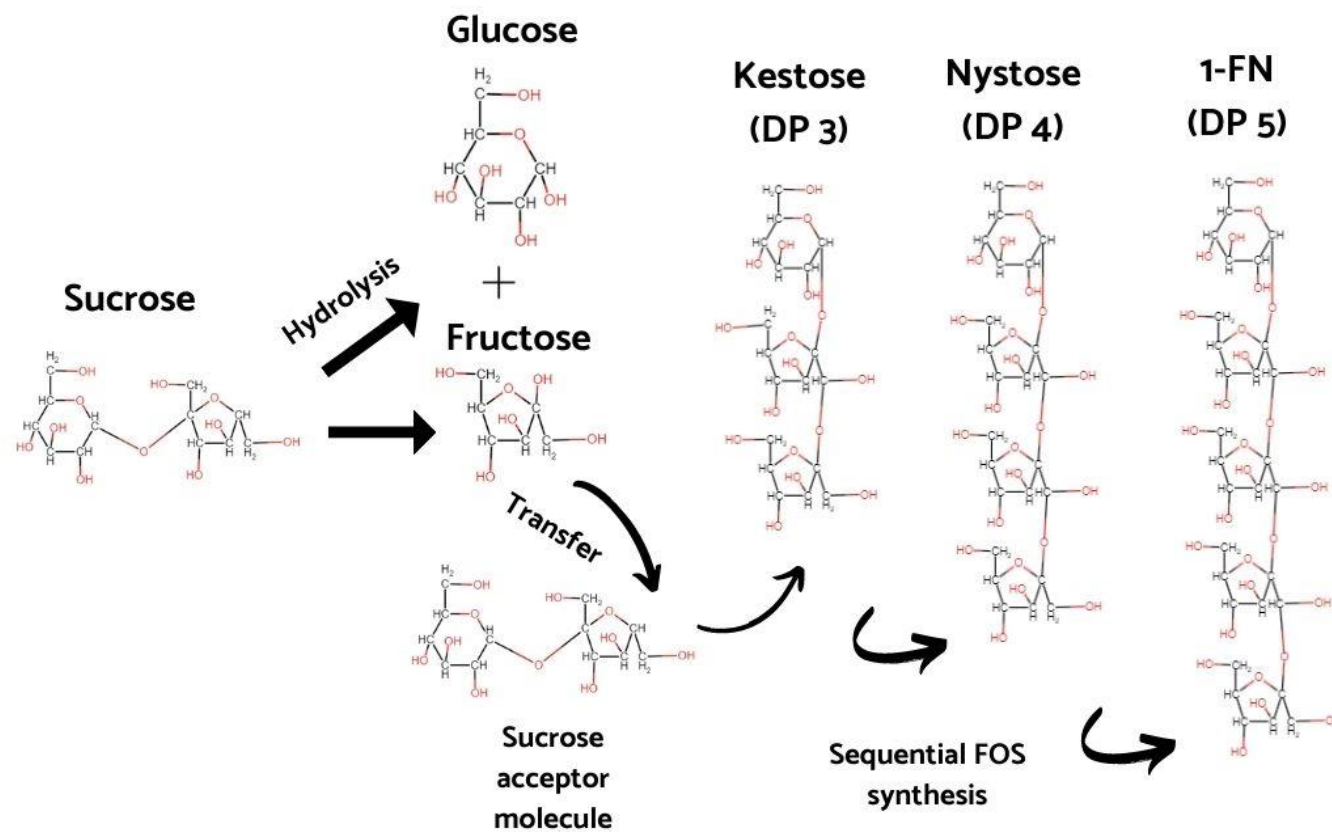


Figure S1. Schematic representation of fructooligosaccharides (FOS) structures obtained by synthesis during fermentation with strain *Aspergillus oryzae* DIA-MF. DP: degree of polymerization; 1-FN: 1-Fructofuranosyl nystose.

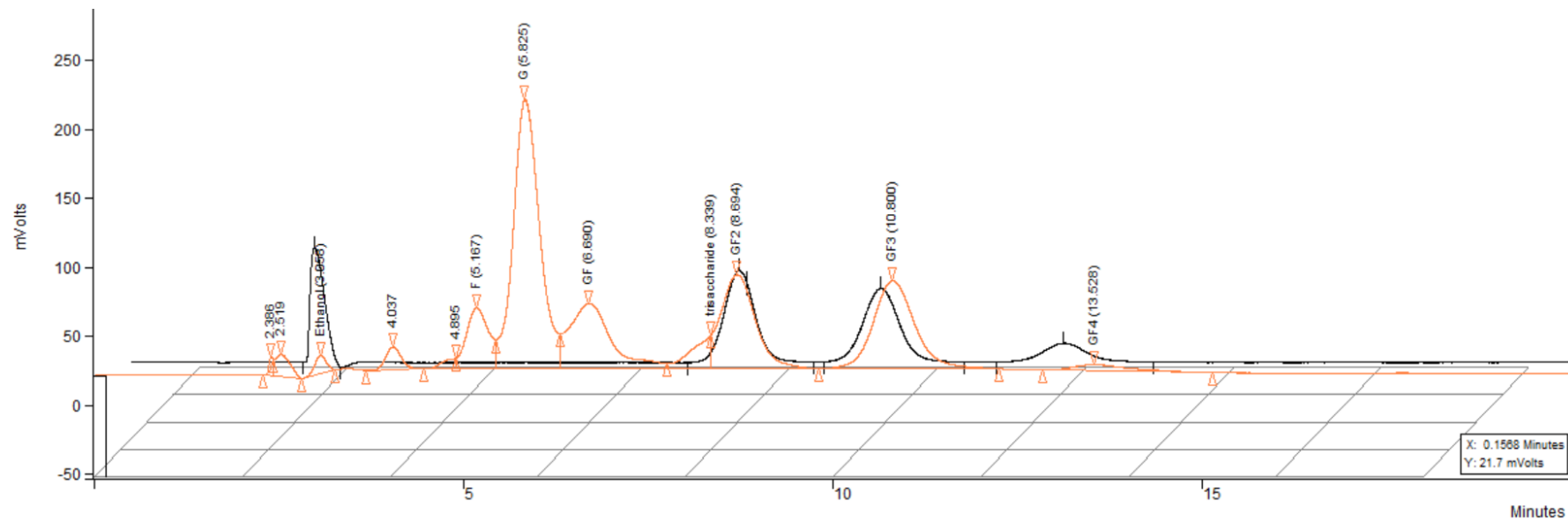


Figure S2. Chromatograms obtained by HPLC for: a mixture of standard fructooligosaccharides (FOS) (Black line) and for a fermentation broth sample obtained from the fermentation of Aguamiel and Molasses (AgMe) with *Aspergillus oryzae* (Orange line). Fructose (F), glucose (G), sucrose (GF), unidentified compound - possibly a trisaccharide, kestose (GF2), nystose (GF3), 1-Fructofuranosyl nystose (GF4).

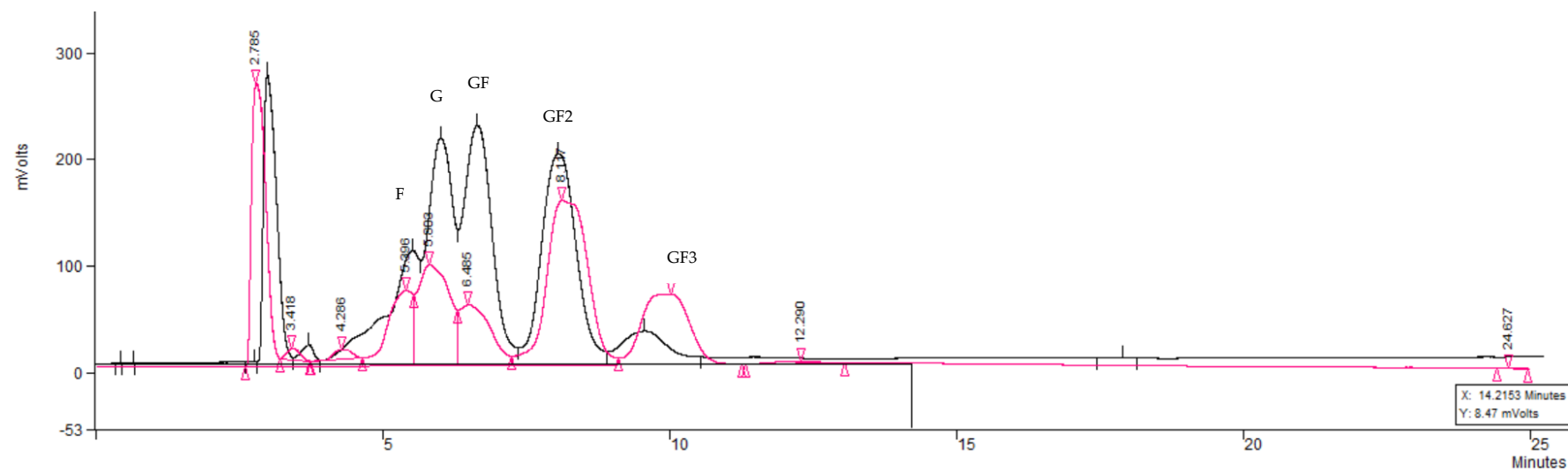


Figure S3. Chromatograms obtained by HPLC for: for a fermentation broth sample obtained from the fermentation of Aguamiel and Molasses (AgMe) with *Aspergillus oryzae* before (black line) and after (pink line) inoculation of *Saccharomyces cerevisiae* 227 (S227). Fructose (F), glucose (G), sucrose (GF), kestose (GF2), nystose (GF3).

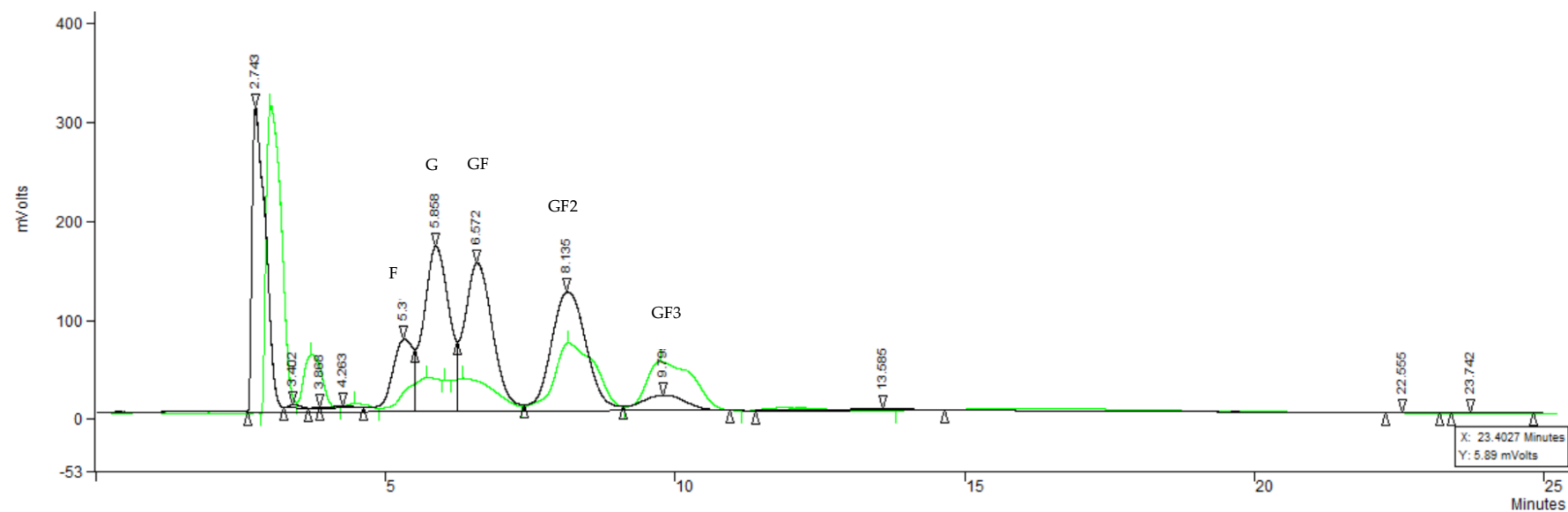


Figure S4. Chromatograms obtained by HPLC for: for a fermentation broth sample obtained from the fermentation of Aguamiel and Molasses (AgMe) with *Aspergillus oryzae* before (black line) and after (green line) inoculation of *Saccharomyces cerevisiae* 200 (S200). Fructose (F), glucose (G), sucrose (GF), kestose (GF2), nystose (GF3).

Table S1. Sugar concentration obtained on the aguamiel from *Agave salmiana* and on the sugar cane molasses used as raw materials for media formulation.

Raw material	Fructose (g/L)	Glucose (g/L)	Sucrose (g/L)
Aguamiel	4.17 ±0.21	4.33 ±0.39	98.56 ±9.77
Molasses	96.10 ±9.56	71.35 ±6.20	441.41 ±42.02

Table S2. Statistical analysis of the full Central Composite Design with two factors used for the optimization of fructooligosaccharides production.

Variables and interactions	Estimated effects	Standard Error	t-value	<i>p</i>
Mean/Interc	-4.580	1.307	-3.504	0.017*
(1) GFi (L)	0.005	0.004	1.165	0.296
GFi (Q)	0.000	0.000	-1.855	0.123
(2) Inoc (L)	1.348	0.371	3.631	0.015*
Inoc (Q)	-0.097	0.026	-3.711	0.013*
1L by 2L	0.000	0.000	0.474	0.656

GFi: Initial sucrose concentration, Inoc: Inoculum concentration

*Significant influence at 95% confidence level.