Supplementary Information

J. Mex. Chem. Soc. **2025**, *69*(2)
Regular Issue

ISSN-e 2594-0317

Fusarium oxysporum Tolerance assay in Strawberry (*Fragaria x ananassa*) Varieties and Analysis of *FaPAL* Gene Expression in an *In vitro* System

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Received November 2nd, 2023; Accepted Febraury 23rd, 2024.

DOI for the article: http://dx.doi.org/10.29356/jmcs.v69i2.2162

Supplementary Information

Table S1. Severity scale values assigned to the *Fof* tolerance in 'Nikté' and 'Camino Real' strawberry plants, according to the infection severity scales (Table 1).

	'Nikté'		**'Camino Real'**	
	Control	***Fof***	**Control**	***Fof***
Leaves	1 ± 0 b	2.67 ± 1.21 a	1 ± 0 b	2.7 ± 1.12 a
Roots	1 ± 0 b	3.47 ± 1.11 a	1 ± 0 b	3.9 ± 0.92 a
Crown	1.13 ± 0.35 b	3.07 ± 1.11 a	1 ± 0 b	2.87 ± 1.07 a

Table S2. Sequences of the used primers to amplify specific sequences.

Gene	**NCBI acc. number**	**Annealing temperature (Tm, °C)**	**Sequence (forward/reverse) Orientation 5' to 3'**	**Amplicon (pb)**	**Reference**
FaEF1α	BK009992.1	57	TGGATTTGAGGGTGACAACATGA GTATACATCCTGAAGTGGTAGACGGAGG	145	[46]
FaGAPDH2	AF421145	60	CCCAAGTAAGGATGCCCCCATGTTCG TTGGCAAGGGGAGCAAGACAGTTGGTAG	117	[46]
FaPR1	AB462752.1	60	ACATGGGATGCCAATCTAGC CCACAGGTTCACAGCAGATG	150	[25]
FaMBL1	KF962716.1	57	AAACCAACACGGCCAATAAG GTCTGTCGGGTAGTCGAAGC	116	[48]
FaWRKY1	EU727547	60	ACAGCAGTAAGATTAGGGATGAAGAAGGGAG GCTTCTTCACATTGCAACCCTGATGCGTG	196	[46]
FaCyf1	AJ845186	60	GCCAAGTTCAACATGCTACTC TGCTCCACCTCCATCTGAT	116	[47]
FaChi3	AF134347.1	60	ACCAAGTTCAGCTCGCAGAT TCCTAATGGCCTTGAAGTGG	177	[25]
FaPAL1	KX450226.1	57	CTTCCTCAAGATTGCTGCTT AATTCCTCCCTCACAAACCT	159	This work
FaPAL2	KX450227.1	57	CATTACTCCCTGCTTGCCTCTCC CCTGAGCTGATACCGACTTGTTCG	171	This work

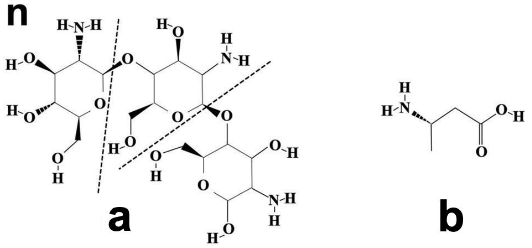

Fig. S1. Structure of chitosan **(a)**, and BABA **(b)**. Chitosan is a polymer of β-(1–4)-linked D-glucosamine repeated "n" times, in comparison to BABA structure.

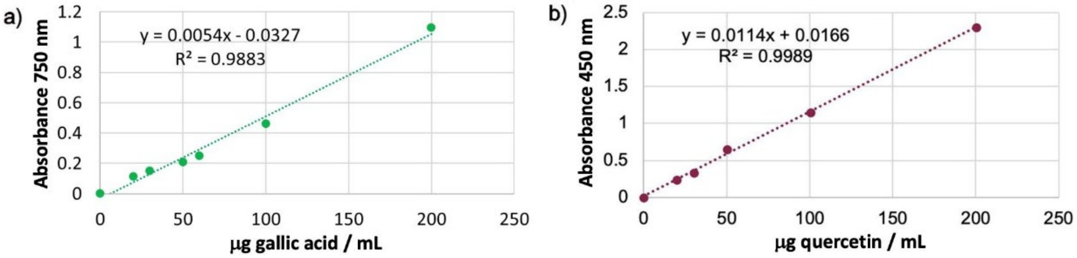

Fig. S2. Standard curves of GA and quercetin to quantify phenols and flavonoids in strawberry tissues.

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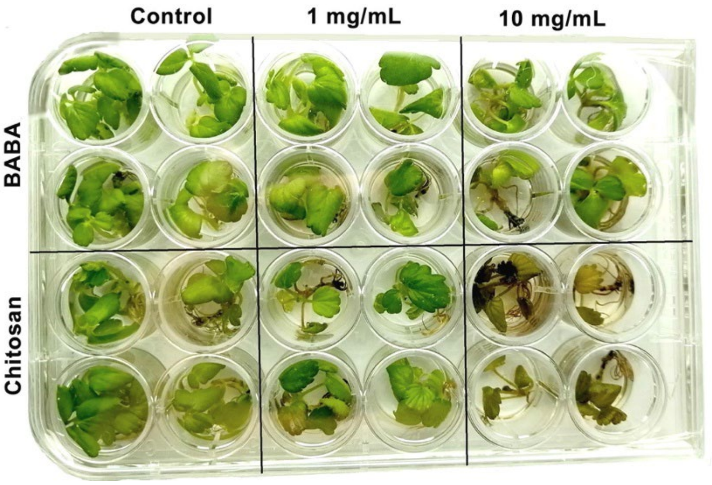

Fig. S3. BABA and chitosan treatments of strawberry plantlets. Results of chosen concentrations of BABA 10 mg/mL and chitosan 1 mg/mL after 3 days, are shown. Controls were diluted acetic acid pH 5.8 for chitosan, and DDW for BABA.

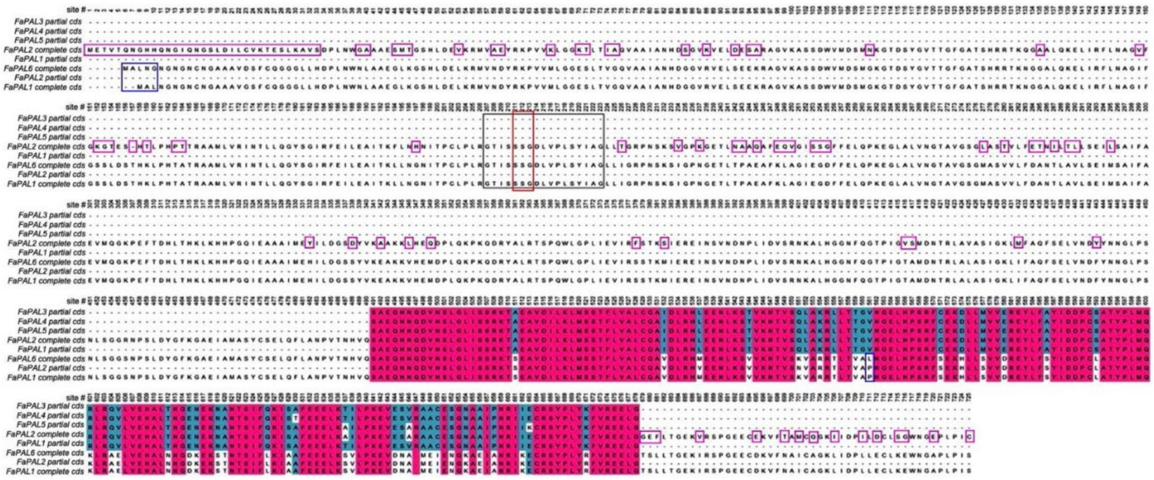

Fig. S4. Complete amino acid sequences of FaPAL1, FaPAL2 and FaPAL3. Magenta boxes are the differences between FaPAL 2, and FaPAL1 - FAPAL6. Blue boxes are differences between FaPAL1 and FaPAL6. Black box is the active site and red box is the catalytic triad. Pink and green boxes are the conserved sequences of the partial *FaPAL1, FaPAL2, FaPAL3*, *FaPAL4* and *FaPAL5*.