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ВЛИЯНИЕ МУТАЦИИ АУТОИНГИБИТОРНОГО ДОМЕНА СDPK1 ARABIDOPSIS THALIANA НА ИНДУЦИРОВАННОЕ ТЕПЛОВЫМ СТРЕССОМ СТАРЕНИЕ В ТРАНСГЕННЫХ РАСТЕНИЯХ ТАБАКА Effect of mutation in autoinhibitory domain of CDPK1 Arabidopsis thaliana on heat-induced stress senescence in transgenic tobacco plants EFFECT OF MUTATION IN AUTOINHIBITORY DOMAIN OF CDPK1 ARABIDOPSIS THALIANA ON HEAT-INDUCED STRESS SENESCENCE IN TRANSGENIC TOBACCO PLANTS



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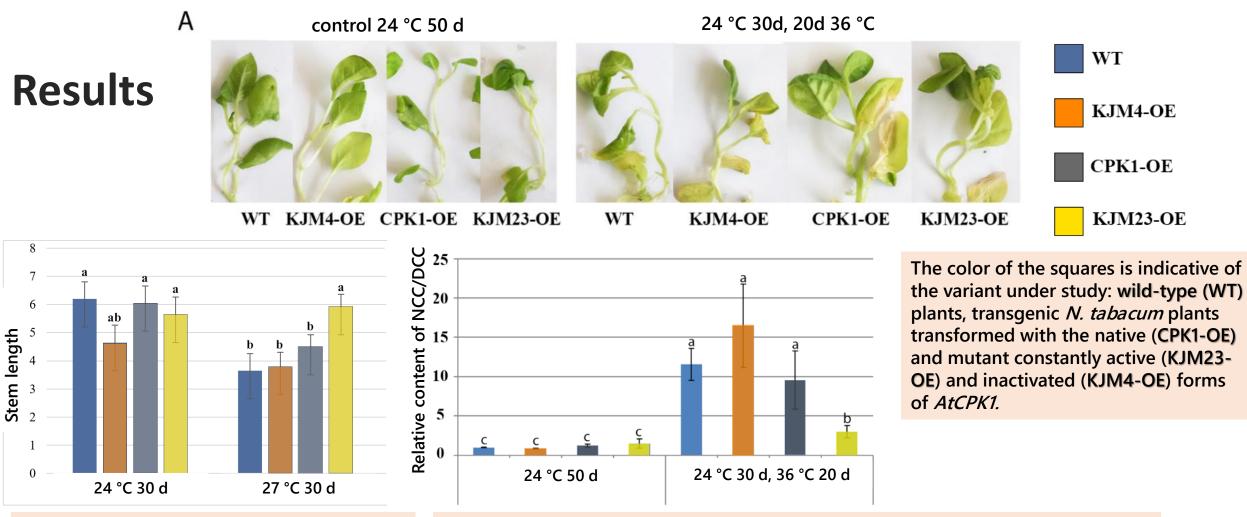
This study aimed to investigate whether a mutation in the junction domain of the CDPK1 in *Arabidopsis thaliana*, which renders it independent of fluctuating cytoplasmic calcium ion levels, affects the tolerance of *Nicotiana tabacum* L. plants to high temperatures and heat-stress-induced senescence.

Arabidopsis CPK1 gene were transferred by agrobacterial transformation. Domain structure of CDPK: N-terminal variable domain, responsible for localization They correspond to the name of the mutant forms that were obtained in the and specific recognition; C-terminal Ca²⁺-binding domain; K, kinase and J junction, 90s by Harper and Huang while studying the mechanism of domain, which inhibits kinase activity in the absence of calcium ions pseudosubstrate inhibition [1]. N AtCPK1-Wt B The native form WT of CPK1 functions only - Ca²⁺ activity 1.5% **VDGVAPDKPLDSAVLSRMKQFSAMNKFKKMALRVI** in the presence of calcium; + Ca²⁺ activity 100% The mutation KJM4 has been shown to lead to a reduction in enzyme activity to almost AtCPK1-KJM4 ******* zero, irrespective of calcium stimulus. AtCPK1-Wt - Ca²⁺ activity 0.6% The KJM23 mutation, a 6-amino acid AtCPK1-KJM4 AtCPK1-KJM23 + Ca²⁺ activity 1.2% substitution in the central part, had the D opposite effect and made the enzyme AtCPK1-KJM23 activity close to 100%, regardless of the * * * * * * * * * * * * * * * * * * * AtCPK1-Wt - Ca²⁺ activity 86% calcium stimulus. AtCPK1-KJM23 AtCPK1-KJM4 + Ca²⁺ activity 97% ruler 1240.....1250.....1260.....1270.....1280.....1290.....1300.....1310.....1320... Modeling the effects of short-term heat stress Modeling the effects of prolonged heat stress 40-day-old in vitro plants 20/30 day-old plants in vitro Stem length HPLC and mass Spectrophotometric detection 37 °C **ABA** extraction **ELISA** measurement 🛸 spectrometry 30 days 1 hour 20 days Target gene: XX 24 °C NCED1 Methanol extraction **cDNA** NCED3 **RNA** extraction ACD6 synthesis **SAG12**

[1]Harper, J.F. Genetic identification of an autoinhibitor in CDPK, a protein kinase with a calmodulin-like domain / Harper J.F., J.-F. Huang, S.J. Lloyd // Biochemistry. – 1994. – № 33. –P. 7267–7277.

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Real-time PCR



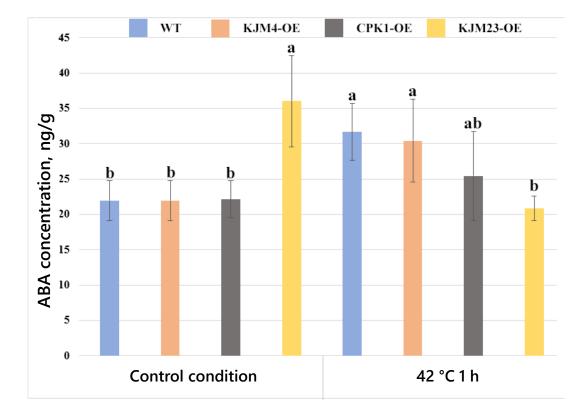
Prolonged exposure to heat stress was observed to result in a decrease in growth rate for the plants in this study. However, the KJM-23 transgenes exhibited no such change, maintaining a growth rate that was equivalent to that of the plants under control conditions. Externally, KJM23-transgenic plants exhibited a reduced susceptibility to the negative effects of heat stress, as evidenced by a decrease in leaf yellowing. When analysing the content of chlorophyll catabolites, no differences were observed between the studied forms and the wild type under control conditions. However, under prolonged heat stress, the content of chlorophyll catabolites increased 12-fold for the two studied forms and the wild type. Only a two-fold increase in degradation products was observed for the transgenes with the permanently active form, indicating strong defence mechanisms activated in these plants.

Results

	A		B	
	NCED1	NCED3	ACD6	SAG12
WT	1.36±0.11	1.05±0.03	3.21±0.79	8.55±0.21
C KJM4-OE	1.25±0.07	1.67±0.11	3.42±0.28	9.61±1.38
E CPK1-OE	1.06±0.17	1.63±0.07	2.77±0.10	16.21±0.94
UKJM23-OE	1.45±0.20	7.53±0.17	17.51±2.34	44.34±4.47
E WT	1.38±0.03	2.36±0.04	1.98±0.01	12.18±0.14
KJM4-OE	1.29±0.01	2.50±0.04	1.23±0.11	11.94±0.71
	1.27±0.04	3.64±0.03	6.77±2.55	16.92±0.65
[◀] KJM23-OE	1.18±0.03	1.05±0.14	1.22±0.04	1.11±0.15
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Relative expression level

Abscisic acid (ABA) is directly related to the aging process, so it is important to study the expression of its biosynthesis genes and the expression of proteases gene involved in cell death. In KJM-23 transgenic plants, an increased expression of the NCED3 gene under control conditions was observed. The transformation of KJM23 also activated the expression of SAG12 and ACD6 genes. However, under heat stress conditions, there was a complete blockage of the expression of both biosynthesis genes and agingrelated proteases. The native form of the gene didnt have this effect, and its expression pattern was similar to the wild type.



It was necessary to confirm the expression of data by analyzing abscisic acid content. As a result of this analysis, it was possible to reliably confirm an increase in the content of abscisic acids in KJM-23 transgenic plants under control conditions, as well as a dramatic decrease in concentration of abscisin to wild-type levels under control conditions in response to stress.

Conclusions

- In the presented work we showed that the transformation of tobacco plants to a constitutively active modified form of the AtCPK1 gene (KJM23-OE plant) allows to significantly mitigate such negative heat stress effect as degradation of chlorophyll.
- Detailed molecular-biochemical analysis showed that KJM23-OE plant resistance to long-term heat stress is accompanied by a decrease in the level of ABA biosynthesis and a decrease in ABA-dependent processes of stress-induced aging, presumably due to CPK-dependent on the activation of the SA signal system, which is the antagonist of the ABA signal system.
- The overexpression of the native form of the gene had no effect on resistance to long-term intense heat stress.

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Thank you for your attention