ICARHSE

October 15th 2022

 $International\ Conference\ on\ Advance\ Research\ in\ Humanities,\ Sciences\ and\ Education$

MALASIA CONFERENCE

https://confrencea.org

UDC 633.511:631.52:575.126:632

CREATION OF NEW COTTON VARIETY RESISTANT TO VERTICILLIUM DAHLIAE USING DNA MARKERS

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Abstract. This article presents the results of research on the creation of a new cotton variety C-6602, resistant to pathogens of the *Fusarium* and *Verticillium*, using modern methods of molecular genetics, biochemistry and phytopathology.

Keywords: *G.hirsutum L., F.oxysporum, V.dahliae*, phenotypes, DNA marker, peroxidase, morphology, pathogenicity, strains, variety, hybrid, generation, molecular genetics, selection,

Pathogens of the genera *Fusarium* and *Verticillium* are the most harmful pathogens among those affecting harvest and product quality in cotton. Species of the genus *Fusarium* Lk. cause damage to seeds after sowing, root rot at the germination stage, and additional damage to seedlings and young cotton plants. The harmful effects of seed and seedling damage are exacerbated by the fact that 80% of the future cotton yield is determined within 30 days from the date of seedling emergence.

Purpose of research: in this study, the molecular genetic diversity of several cotton breeding lines was analysed using DNA markers, and phytopathological and biochemical evaluations of lines resistant to *Fusarium* and *Verticillium wilt* were carried out. Four hybrids were identified as having the most potential with regard to the trait "resistance to *V.dahliae* and *FOV*" after evaluation: F_{11} [F_6 (L-101 x L105) x J-106], F_{11} [F_6 (L-105 x L-106) x L-104], F_7 [L-175/276 x Namangan-102] and F_7 [Jarkurgon x Namangan-34]. Eight predefined SSR markers were used for the DNA analysis. These DNA markers were identified in previous studies as being correlated with resistance to Fusarium and Verticillium wilt. The genotyping results, especially those for three primer pairs (BNL4003, CM209, TMB0161), showed that the best prospective hybrid was F_{11} [F_6 (L-101 x L-105) x L-106] because the largest number of individuals within this line were identified as having candidate loci linked to wilt resistance traits. Based on a biochemical analysis using marker proteins as well as a cytochemical study of the roots of various cotton lines, the hybrids F_{11} [F_6 (L-101 x L-105) x L-106] and F_7 [Zharkurgon x Namangan-34] were characterized as the most resistant to *V. dahliae*.

The results of molecular and biochemical research under laboratory conditions revealed that from the analysed selection material, line L-1077 exhibited high resistance to V.dahliae and FOV. This line was separated from the complex, interlinear hybrid combination $F_{11}[F_6(L-101 \text{ x L-105}) \text{ x L-106}]$. L-1077 produces fibre that meets the type IV fibre quality standards, produces high fibre yields and is highly resistant to V.dahliae and Fusarium. Moreover, an application (NAP 2020 0007 from 31.01.2020) for a patent for this line has been issued by the Intellectual Property Agency of the Republic of Uzbekistan under the cotton variety name C-6602.

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