## STUDIES ON SOME SOURCES OF RESISTANCE FOR FUSARIUM WILT IN WATERMELON ECOTYPES

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## **ABBREVIATIONS**

**C.V** Coefficient of Variance

**TSS** Total Soluble Solids

**MP** Mid Parents

**GCA** General Combining Ability

**SCA** Specific Combining Ability

KVRF Kaha Vegetable Research Farm

**EP** Experimental Plot

RCBD Randomized Complete Block Design

**ADH** Average Degree of Heterosis

**BP** Better Parent

R Resistant

S Susceptible

PAL Phenylalanine Ammonia Lyase

PPO PolyPhenol Oxidase

PO PerOxidase

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**Title of Thesis:** Studies on Some Sources of Resistance for Fusarium Wilt in Watermelon

**Ecotypes** 

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#### **ABSTRACT**

This study was conducted during the period from 2011 to 2016 at Kaha Vegetable Research Farm (KVRF), Qalubia Governorate to evaluate some watermelon landraces collected from ten different local geographic regions of Egypt to increasing their homogeneity through inbreeding and selecting the best individual genotypes during four successive generations to produce 15 inbred lines. Seedling inoculation procedures were used to screen 15 lines of watermelon ecotypes for resistance to fusarium wilt. Of these ecotypes, 5 lines significantly showed resistance to fusarium oxysporum. And used four lines as female parent Suhag 1(P1) - Suhag 2(P2)-Sinaa (P3)-Bir El-Abd 2(P4); and the cultivars Giza1 (P5), Crimson Sweet (P6) and the line (Beni Swif) (P7) were used as male parents in a line X tester mating design to produce twelve hybrids. Parents and hybrids were evaluated to horticultural characters and resistance to fusarium wilt. Biochemical parameter such as phenolic content and some enzymes in three watermelon hybrids shown susceptible, moderately resistance and resistant to fungus fusarium oxysporum f.sp. niveum.

Estimates of coefficient of variance (C.V %) values showed that, most of the breeding lines were highly homogeneous and could be considered pure lines. All parents and crosses were significant with positive value for all traits studied. With respect to heterosis effects, cross P2 XP6 gave highest values with significant effect for plant fresh weight, total yield /plant and fruit weight for mid-parent and better parent. Crosses P2X P7, P3X P5 and P4 X P6 gave earliest female flower over midparent and better parent, cross P4XP6 gave highest values with significant effect for rind thickness for mid-parent and better parent. Cross P1XP7 gave highest values with significant effect for TSS for mid-parent. Results showed that the lowest mean performance of seven parents and their twelve crosses of watermelon for disease severity on the foliage growth were 24.8% and on the xylem vesicles were 19 % for parents line (Beni Swif) (P7) and the lowest disease severity on the foliage were 24.30 % for crosses (P4XP5) and on the xylem vesicles were 28 % for crosses (P4XP7). Data reveal that, in general, the three enzymes, i.e. PAL, PPO and PO and free, conjugate and total phenols were greatly increased in the shoot base of the hybrids by increasing the resistance of these hybrids.

**Key words:** landraces, Evaluation, *Fusarium* resistant, KVRF.

## INTRODUCTION

Watermelon, (*Citrullus lanatus, Thunp.*) is an important horticultural crop across wide areas of the world. Having 2n=22 chromosomes. Genetic information on watermelon could be obtained from inter-varital crosses which need to increase production, improved quality and resistance to diseases. Watermelon fruits mostly eaten fresh, but in Africa they are also cooked. The rind may be pickled or candied. (Robinson and Decker-Walters 1997).

Giza1 watermelon cultivated area reached 81485 fed, which yielded about 13.73 tons/fed, with a total production of 1118771 tons <sup>(1)</sup>, while Balady watermelon cultivated area reached 30341 fed, which yielded about 12.89 tons/fed, with a total production of 391261 tons <sup>(1)</sup>.

All over the world, many investigators are interested in collecting their landraces or accession lines and estimate their agronomic traits or pathogenicity test in order to raise a new genotype for breeding programs.

Fusarium wilt of watermelon is a soilborne disease caused by the fungus *Fusarium oxysporum* f. sp. *niveum*. The disease is a major problem in watermelon production in susceptible varieties. Presently, within the formae specialis *F. oxysporum* f. sp. *niveum* occurs in three races: races 0, 1 and 2 on the basis of their degree of aggressiveness on watermelon differential genotypes (Crall, 1963; Martyn, 1987, Martyn and Bruton, 1989; Netzer, 1976; Zhou and Everts, 2004).

<sup>1.</sup> Department of Agriculture and Economics and Statistics, Ministry of Agriculture A.R.E. 2016.