

Evalution of the progeny from transgenic watermelon resistance to Zucchini yellow mosaic virus and p

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ABSTRACT

Zucchini yellow mosaic virus (ZYMV) and Papaya ringspot virus type W (PRSV W) are major limiting factors for production of watermelon in Taiwan. For the effective control of these two viruses by transgenic resistance, an untranslatable chimeric construct containing truncated ZYMV coat protein (CP) and PRSV W CP genes was transferred to commercial watermelon cultivars by Agrobacterium-mediated transformation. Greenhouse evaluation of the selected 10 transgenic lines of "Feeling" cultivar revealed that two immune lines conferred complete resistance to ZYMV and PRSV W, from which virus accumulation were not detectable by western blotting four weeks after inoculation. The transgenic transcript was not detected but small interfering RNA (siRNA) was readily detected from the two immune lines and T1 progeny of line ZW 10 before inoculation, indicating that RNA-mediated post-transcriptional gene silencing (PTGS) is the underlying mechanism for the double-virus resistance. The segregation ratio of T1 progeny of the immune line ZW10 indicated that a single inserted transgene is nuclearly inherited and associated with the phenotype of double-virus resistance as a dominant trait.

Keywords : transgenic watermelon、coat protein、PTGS

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