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LIBRO DE RESÚMENES



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P4-05 The loss of function of the tomato homolog of *ROTUNDIFOLIA3* impairs leaf expansion and plant growth

<u>Teresa Barragán-Lozano</u>^a, Ricardo Lebrón^a, Benito Pineda^b, Begoña García-Sogo^b, Fernando J. Yuste-Lisbona^a, Vicente Moreno^b, Rafael Lozano^a

^aCentro de Investigación en Biotecnología Agroalimentaria (CIAMBITAL), Univ. de Almería. Almería, Spain. ^bInstituto de Biología Molecular y Celular de Plantas (UPV-CSIC), Univ. Politècnica de València, Valencia, Spain.

We characterized the tomato 2489etmm recessive mutant, which belong to our T-DNA insertional mutant collection (1). Homozygous mutant plants showed slower growth rate at greenhouse conditions, as well as wrinkled leaves and enlarged stem. Despite these developmental defects, 2489etmm mutant plants were able to develop flowers and yield seed-bearing fruits after almost ten months growing under optimal conditions, in contrast to the three months required by the wild-type plants. By combining mapping-by-sequencing and CRISPR/ Cas9 genome editing methods, we proved that a 2-bp deletion in the tomato homolog of the Arabidopsis ROT3 gene (SIROT3) was responsible for the phenotype observed in the 2489etmm mutant. We engineered knockout mutations at the SIROT3 locus by using the CRISPR/Cas9 system with a single guide RNA. Independent first-generation CRISPR lines homozygous or biallelic for edited mutant alleles showed slower growth, wrinkled leaves, and enlarged stem, a phenotype resembling that observed in the 2489etmm mutant. The ROTUNDIFOLIA3 (ROT3) gene encodes a cytochrome P450 family protein involved in regulating leaf length of the model species Arabidopsis thaliana. Specifically, ROT3 is required for the conversion of typhasterol to castasterone in the early C6-oxidation pathway of brassinosteroid biosynthesis (2). Brassinosteroids are a group of plant steroid hormones and playing a key function during plant growth and development. Together, our results suggest that *SIROT3* may act regulating tomato leaf expansion in a similar manner as occurs in Arabidopsis, by probably participating in brassinosteroid biosynthesis pathway.

1. Pérez-Martín et al. (2017). Plant Biotechnology Journal, 15: 1439-1452. 2. Ohnishi et al. (2006). The Plant Cell 18, 3275–3288.

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