Name: Regeneration1 (Rg1)

Accessions: Br1

Gene ID:

Map position: chromosome 3 (short arm)

Gene function:

Gene effect: Controls the phase of acquisition of competence for cell differentiation, which determines the capacity for both shoot and root formation in vitro from different explants.

Phenotypes: MT-*Rg1* presents high *in vitro* regeneration, and excessively shoot branching. 10% of MT-*Rg1* seedlings present bifurcate of tri cotyledons.

Comments:

Description of accessions available: MT-*Rg1* is a BC6Fn introgressed from LA4136.

Figures:



MT-Rg1 (right) showing excessively branching (A). The yellow fruits are due to the presence of *yellow flesh* (*r*) allele linked to Rg1. (B) high *in vitro* shoot regeneration from cotyledonary explants in Rg1. The plates in the bottom contain kanamycin for transgenic shoot selection upon infection with Agrobacterium.

Bibliography

Boiten H, Azmi A, Dillen W, De Schepper S, Debergh P, Gerats T, Van Onckelen H, Prinsen H (2004) The *Rg-1* encoded regeneration capacity of tomato is not related to an altered cytokinin homeostasis. New Phytologist 161:761-771.

Koornneef M, Bade J, Hanhart C, Horsman K, Schel J, Soppe W, Verkerk R, Zabel P (1993) Characterization and mapping of a gene controlling shoot regeneration in tomato. Plant Journal 3:131-141.

Lima JE, Benedito VA, Figueira A, Peres LE (2009) Callus, shoot and hairy root formation in vitro as affected by the sensitivity to auxin and ethylene in tomato mutants. Plant Cell Reports 28:1169-77.

Lima JE, Carvalho RF, Tulmann Neto A, Figueira A, Peres LEP (2004) Micro-MsK Micro-MsK: a tomato genotype with miniature size, short life cycle, and improved in vitro shoot regeneration. Plant Science 167:753-757.

Pino LE, Lombardi-Crestana S, Azevedo MS, Farinha TB, Borgo L, Quecini V, Figueira A, Peres LEP (2010) The *Rg1* allele as a valuable tool for genetic transformation of the tomato Micro-Tom model system. Plant Methods 6:23.