Erratum

FEBS 22442

Erratum to: Plant cell growth and differentiation may involve GAP regulation of Rac activity (FEBS 22204)

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In the original publication of this article, the top and right hand edges of Fig. 1 were missing. It is reprinted in full below. The publisher apologizes to authors and readers.

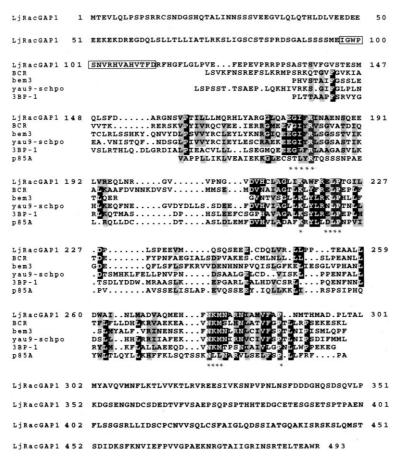


Fig. 1. The LjRacGAP1 amino acid sequence. The putative GAP domain is compared to five GAP domain sequences known from mammalian and fungal cells. Human BCR, AC: P11274, residues 1029–1221; yeast bem3, AC: P32873, residues 898–964 and 994–1099, *Schizosaccharomyces pombe* yau9, AC: Q10164, residues 1041–1238; mouse 3BP-1, AC: P55194, residues 183–362, and bovine PI3'K p85α, AC: P23727, residues 127–276. Points correspond to gaps that are introduced to maximize homology. Asterisks under the sequences designate core motifs of three GAP subdomains. The three *LjRacGAPs* sequences have been deposited at GenBank under accession numbers AF064787, AF064788 and AF064789.

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