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OPEN Author Correction: CLE peptideencoding gene families in Medicago truncatula and Lotus japonicus, compared with those of soybean, common bean and Arabidopsis

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This Article contains an error in the legend of Figure 5.

"Multiple sequence alignment of the prepropeptides of AtCLE18 and LjCLE34. CLE domains are highlighted with a red box and the CLEL domain is underlined in blue. Conservation between amino acid residues of the two sequences is represented by grey (partial) and black (100%) shading".

should read:

"Sequence alignment of LjCLE5 and MtCLE12. The LjCLE5 translation start site corresponding to that of MtCLE12 results in a truncated prepropeptide (denoted by the asterisk shaded in black). The blue box represents an alternative start codon that results in the CLE domain region of LjCLE5 (underlined in red) being translated in frame. Black highlighted nucleic acid bases are 100% conserved".

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