

Supplementary Material

Identification and functional analysis of calcium sensor calmodulins from heavy metal hyperaccumulator *Noccaea caerulescens*

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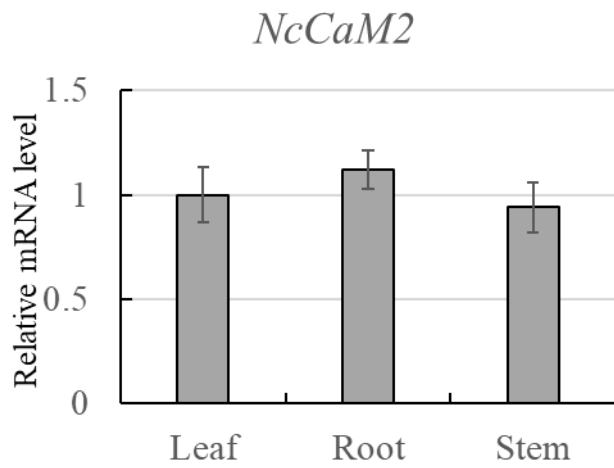


Figure S1 **Tissue-specific expression analysis of *NcCaM2***. The expression of *NcCaM2* in leaf was set as 1. The relative expression levels of *NcCaM2* in different tissues were calculated compared with that in leaf.



Figure S2 **Genetic transformation of NcCaM2 in tobacco seedlings.** (a) The growth process of tobacco materials after Agrobacterium-mediated genetic transformation. (d) GUS staining of the root from positive tobacco lines expressing NcCaM2:GUS fusion protein. (c) GUS staining of the leaf from positive tobacco lines expressing NcCaM2:GUS fusion protein. (e)

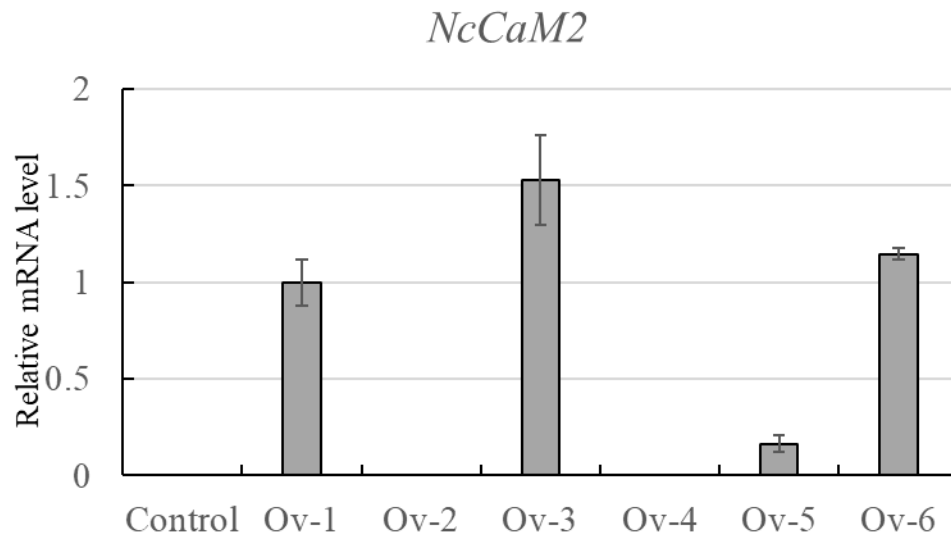


Figure S3 qRT-PCR checking the expression level of *NcCaM2* in tobacco seedlings. The expression of *NcCaM2* in wild type seedlings was treated as negative control. The relative expression levels of *NcCaM2* in Ov-1 line was set as 1. The expression level of *NcCaM2* in different over-expressing lines was calculated compared with that in Ov-1. ‘Ov’ means over-expressing line.