

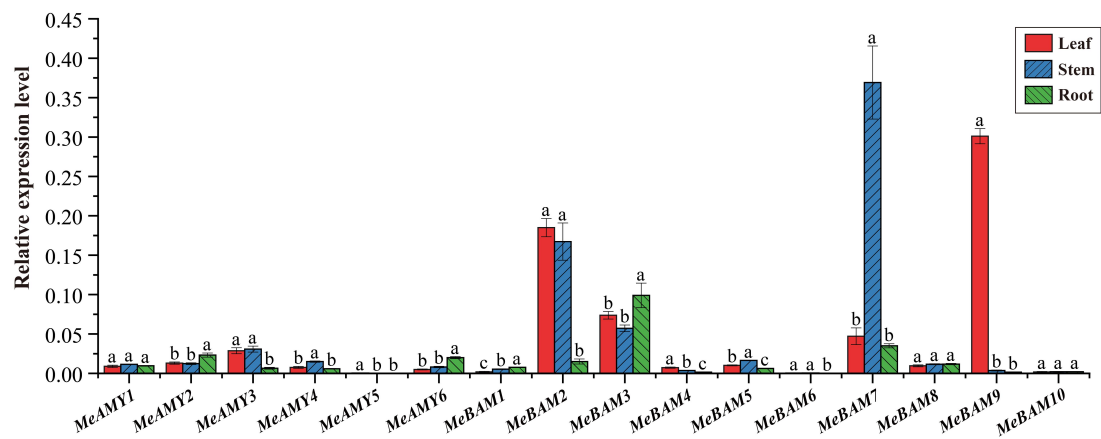


Fig. S5 Expression patterns of *MeAMY* and *MeBAM* genes in different parts of cassava under well watering condition. Expression levels were determined using qRT-PCR and calculated using the $2^{-\Delta C_t}$ method under the control of the *actin* housekeeping gene. Data are expressed as the mean $\pm$ SE of three independent replicates. Different letters indicate significant differences among different tissues of cassava ($P < 0.05$).