a Mp4g22840 (NRT2.1 ortholog)



Supplementary Figure S1 Transcript levels of genes related to nitrate assimilation and transport in the Mp*CKX2*-overexpressing line and the Mp*rrb* knockout mutant. (a) Mp4g22840 (*NRT2.1* ortholog), (b) Mp8g18730 (*NRT2.4* ortholog), (c) Mp4g16300 (*NRT3.1* ortholog), (d) Mp3g24720 (*NIR1* ortholog), and (e) Mp6g08310 (*AMT1* ortholog) were examined by qRT-PCR. mRNA levels were normalized to that of Mp*EF1a*. Data are presented as mean \pm SD (n = 3). Significant differences from the wild-type (Tak-1 or Tak-2) were determined by the Student's t-test as follows: *P < 0.05; **P < 0.01.

b Mp8g18730 (*NRT2.4* ortholog)



Supplementary Figure S2 Promoter activity of Mp*GCAM1* in the presence of exogenous cytokinin. 20day-old thalli of Mp*GCAM1pro:GUS* were treated with or without 50 μ M of tZ for four hours and subjected to GUS staining. Bar represents 2 mm.



Supplementary Figure S3 Gemma number per gemma cup in the Mp*rra* knockout line. Data are represented as mean \pm SD (n = 7). Significant difference from the wild-type (Tak-2) was determined by the Student's t-test: n.s., not significant.



Supplementary Figure S4 Morphological phenotype of the Mp*gcam1* Mp*rra* double mutants grown in the presence of exogenous cytokinin. (a) Thallus apices of the two Mp*gcam1* Mp*rra* double mutants (#15 and #17) were treated with (+tZ) or without (mock) 50 μ M tZ for 14 days. The right panels are magnified images of apical notches. Bars represent 2 mm. (b) Magnified image of gemma cups produced near the apical notch of tZ-treated plants (#15). Arrowheads indicate gemma cups. Bar represents 500 μ m.