LETTER

Underground gibberellin activity: differential gibberellin response in tomato shoots and roots

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Summary

GAs (Gibberellins) are growth-promoting hormones that regulate organ growth, mainly via cell elongation. Contradicting reports leave an open question of whether GA is as important for root elongation as it is for stem elongation.

Here we have addressed this question focusing on tomato (*Solanum lycopersicum*) primary-root elongation. We used a combination of physiological, molecular and genetic approaches to tackle this question.

Tomato has three GA receptors; GID1a, GID1b1 and GID1b2. The loss of all three receptors, strongly suppressed stem elongation and leaf expansion, but had a relatively minor effect on primary root elongation. The effect of GA on cell elongation and gene-expression was much weaker in roots, than in shoots, reaching saturation at lower hormone concentrations. Our results imply that this differential response to GA in shoots and roots is caused by the lower expression of the dominant GA receptor GID1a in roots.

We show that the differential activity of GA between shoots and roots affects root-to-shoot ratio, and speculate that this evolved as an adaptive mechanism to changing environments.

Plant organ growth is governed and modified by developmental programs and environmental cues. In most cases, these changes are mediated by the activity of phytohormones (Bradford and Trewaves, 1994; Verma *et al*., 2016). GAs (Gibberellins) are growth promoting hormones that regulate many developmental processes, including organ growth and elongation (Davière and Achard., 2013). GA affects elongation by promoting cell division and expansion (Ubeda-Thomas *et al*., 2009). The nuclear DELLA proteins inhibit all GA-elongating responses (Locascio *et al*., 2013) and GA binding to the GID1 (GIBBERELLIN-INSENSITIVE DWARF1) receptor leads to DELLA degradation and activation of growth (Ueguchi-Tanaka *et al*., 2005; Ueguchi-Tanaka *et al*., 2007).

While GA plays a central role in stem elongation (Sun and Gubler, 2004), its general significance for root elongation is less clear, with numerous conflicting reports (Torrey, 1976; Feldman, 1984; Phinney, 1984; Tanimoto, 2005; Tanimoto and Hirano, 2013). It is well established that Arabidopsis (*Arabidopsis thaliana*) root elongation depends on GA. Several studies demonstrate the central role of GA in Arabidopsis primary root elongation (Achard *et al*., 2009; Ubeda-Tomás *et al*., 2008; Ubeda-Tomás *et al*., 2009; Rizza *et al*., 2017). The Arabidopsis GA deficient mutant *ga1-3* exhibits shorter primary root, which is rescued by GA application or loss of DELLA activity (Fu and Harberd, 2003). Ubeda-Tomás *et al*. (2008) showed that inhibiting GA signaling specifically in the endodermis of Arabidopsis roots is sufficient to disrupt root elongation, indicating that the endodermis is the key site for GA action in the regulation of root elongation. This was supported by Shani *et al*. (2013) that found the accumulation of exogenous bioactive tagged-GAs in the endodermis of the elongation zone. Rizza *et al*., (2017) showed that endogenous bioactive GA levels correlate with cell length in Arabidopsis roots. Tanimoto and Hirano (2013) suggest that roots are very sensitive to GA and therefore respond to extremely low GA concentrations. For instance, while root elongation of the *ga1-3* Arabidopsis mutant was strongly induced by low concentration of GA_4 (10^{-10} M), this treatment had no effect on leaf expansion (Arizumii *et al.*, 2008).

In other plant species however, the role of GA in root elongation is indistinct; while Whaley and Kephart (1957) show that GA application to maize (*Zea mays*) promotes root elongation, Svensson (1972) reported that the hormone has no effect on maize root growth. Similarly, Butcher and Street (1960) show that elevating GA concentrations progressively promote tomato (*Solanum lycopersicum*) root elongation, whereas

Tognoni *et al*. (1966), showed that application of GA inhibits tomato root elongation in a concentration-dependent manner. Recent work by Fonouni-Farde *et al*., (2019) showed that GA-treated *Medicago truncatula* plants display shorter primary roots compared to untreated plants, and treatments with the GA-biosynthesis inhibitor Paclobutrazole (PAC) increased root length. These suggest that in Medicago GA inhibits root elongation.

Here we used GA tomato mutants to study the role of GA in primary root elongation. We first examined primary root elongation in the GA deficient mutant *gib-2* (Koornneef *et al*., 1990). We treated 14-day-old WT plants and *gib-2* mutants with 10^{-5} M GA_3 and after two-weeks we measured stem and root length. Untreated *gib-2* stems were almost three times shorter than WT. GA application induced strong stem elongation in both genetic backgrounds, and their final length was similar (Fig. 1a). Non-treated *gib-2* primary roots, however, were not significantly shorter than those of the WT and GA treatment had only a very mild effect on root length (Fig. 1a). This observation suggests that either GA has no effect on tomato root elongation, or that root elongation is highly sensitive to GA and reaches saturation at very low levels of GA. Since *gib-2* exhibits residual GA activity (Illouz-Eliaz *et al*., 2019), this may be sufficient to allow normal root growth. We therefore tested primary root elongation in the $gid1^{TRI}$ (*gid1* triple) mutant that lacks any GA activity due to the loss of all three GA receptors, GID1a, GID1b1 and GID1b2 (Illouz-Eliaz *et al*., 2019). We first measured the elongation rate of primary roots and hypocotyls for 10 days following germination. Hypocotyl elongation rate of $gid1^{TRI}$ was 10-times lower than that of WT (Fig. 1b), but root elongation rate of the mutant was only three-times lower (Fig. 1c). After ten days, primary roots of $gid1^{TRI}$ were ca. 2.5 times shorter than those of the WT, whereas the mutant hypocotyls were 10 times shorter (Fig. 1c). Root-to-shoot ratio was ca. 4 times higher in $gid1^{TRI}$ (Fig. 1d). Since the $gid1^{TRI}$ exhibits very slow growth, we also examined mature plants of the same physiological age (similar number of leaves in $gid1^{TRI}$ and WT). Primary roots of $gid1^{TRI}$ were only two times shorter than those of WT, whereas $gid1^{TRI}$ stems were ca. 10 times shorter than those of WT (Fig 1e). Thus, the lack of GA activity, strongly affects shoot development, but only partially affects primary root elongation. We cannot exclude the possibility that the inhibition of root elongation in $gid1^{TRI}$ can also be a results of limited assimilate supply by the extremely small canopy.

We further tested the effect of exogenous GA on root and stem elongation. To this end, 14-day-old WT seedlings, were treated with 2 mg/L PAC followed by the application of elevating GA_3 concentrations (from 10^{-7}M to 10^{-3}M). After 15 days, we measured stem and primary root length. WT stems exhibited a strong and increased elongation response to rising GA_3 concentrations, reaching saturation at very high concentrations (above 0.5 mM, Fig. 1f). In contrast, primary root length was hardly affected and exhibited a bell-curve response (Fig. 1g). To examine how GA affects cell length, we treated WT seedlings with PAC or PAC with GA_3 (10^{-6}, 10^{-4}, and 10^{-3} M), and after 10 days we measured epicotyl and primary root epidermal cell length using confocal microscopy. Stem epidermal cell elongation strongly responded to GA treatments and cell length of stems treated with PAC and 1 mM GA_3 were four times longer than those treated only with PAC (Fig 1h). Primary root epidermal cells exhibited a very mild response to PAC and GA_3 (Fig 1i) and the effect of the hormone was saturated already at 10^{-6} M. GA-treated cells were only 1.2 times longer than the PAC treated cells.

It was previously suggested that roots are more sensitive to GA than stems (Tanimoto and Hirano, 2013). To further test root sensitivity to GA, we analyzed the response of various known GA-regulated genes, including GA biosynthesis (*GA20-OXIDASE*s and *GA3-OXIDASE*) and signaling (*GID1s*) genes that are downregulated by the hormone (Middleton *et al*., 2012; Illouz-Eliaz *et al*., 2019) and the GA-induced gene, *GAST1* (*GIBBERELLIC ACID STIMULATED TRANSCRIPT1*, Shi *et al*., 1992). Seedlings were treated with 2 mg/l PAC for 10 days and then with several GA_3 concentration (10^{-8} to 10^{-5}M), and three hours later, gene expression in elongating stems and roots was analyzed by qRT-PCR. We found a clear and significant response to 10^{-7}M, but not to 10^{-8}M GA_3, for all genes, in both elongating stems and roots (Fig 2a-f), suggesting similar sensitivity to GA. Moreover, the intensity of the molecular response in roots was weaker and saturated at lower concentrations than in stems. These results are consistent with our cell elongation results (Fig. 1h and i).

We previously showed that GID1a is the dominant GA receptor in tomato stems, due to its high affinity to DELLA and the fact that it is not inhibited by the feedback response to GA (Illouz-Eliaz *et al*., 2019). Its presence alone, in the absence of GID1b1 and GID1b2 activity, can induce the strong stem-elongation response to exogenous GA that is saturated only at very high concentrations. GID1b1 and GID1b2 exhibit a weak GA response that is saturated at low concentrations. To examine if the differential

response to GA in stems and roots results from differential activity of the different GID1s, we tested the effect of GA application on the three tomato *gid1* double mutants. The GID1 family in tomato is composed of three members, therefore each double mutant contains, one active GID1. In stems, only *gid1b1 gid1b2* with active GID1a exhibited strong elongation response, similar to WT (Fig. 2g, Illouz-Eliaz *et al*., 2019). In contrast, roots of this double mutant did not show elongation response (Fig. 2h), suggesting that GID1a does not promote strong GA response in roots as it does in stems. Previously we showed that *GID1a* and *GID1b1* are highly expressed in elongating stems, while *GID1b2* exhibits relatively low expression (Illouz-Eliaz *et al*., 2019). We analyzed available public data of tomato root transcriptome (Koenig *et al*., 2013; Zouine *et al*., 2017; Góra-Sochacka *et al*., 2019; Gray *et al*., 2020) and found very low expression of *GID1a* compared to *GID1b1* and *GID1b2* in all datasets (Fig. 2i presents the data analyzed from Góra-Sochacka *et al*., 2019). This raises the possibility that the minor effect of GA on root elongation, is caused by low *GID1a* expression in roots. Further research regarding the activity of GID1a as a disjunctive component of GA signaling in above-and underground organs is thus highly warranted. An example for such studies could be exploring the effect of highly expressed GID1a under a root-specific promoter.

To conclude, our results suggest that unlike shoots, tomato roots can grow rather well without GA and the hormone has only a minor role in the regulation of primary root elongation. Although previous studies suggest that roots are more sensitive to GA than shoots, this is probably not the case in tomato. We found however that very low GA concentrations are sufficient to saturate root elongation, but not stem elongation.

Plants adjust root-to-shoot ratio to adapt to changes in the environment. Under drought conditions root-to-shoot ratio increases to reduce transpiration and increase water uptake (Xu *et al*., 2015). GA accumulation is inhibited under osmotic stresses, such as drought and salinity (Achard *et al*., 2006; Colebrook *et al*., 2014). Thus, the reduced GA levels strongly affects shoot but not root growth. It is possible that this mechanism evolved as a strategy to modify root-to-shoot ratio under stress conditions.

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Figures legend

Figure 1. GA-induced elongation response spatially differentiates in tomato roots and shoots (**a**) Stem and root elongation of 14-days-old WT and *gib-2* mutants, treated continuously (or not) with 10^{-5} M GA_3. Values are mean of 4 biological replicates ± SE. Each set of letters above the columns represents significant differences (Tukey–Kramer HSD, $P < 0.05$). (**b** and **c**) Hypocotyl (b) and root (c) elongation rate of WT and *gid1*^{TRI} seedlings grown on MS plates. Values are mean of 7 plants of each line and the shadow presents the statistical mean. (**d**) Root-to-shoot ratio of 10-day-old WT seedlings and *gid1*^{TRI} mutants. (**e**) Representative image of WT and *gid1*^{TRI} at the same physiological age. Numbers present the average length of 3 biological replicates ± SE. (**f** and **g**) Stem (f) and primary root (g) length of 14-day-old WT seedlings treated with 2 mg/L PAC followed by GA_3 (10^{-7} to 10^{-3}M). Values are mean of 10 plants ± SE. Each set of letters above the columns represents significant differences (student's t test, $p < 0.05$). (**h** and **i**) Epidermal cell length of elongating stems (h) and primary roots at the elongation zone (i), was measures after 10 days of treatment with PAC or PAC + GA_3 using confocal microscopy and analyzed by imageJ. Each set of letters above the columns represents significant differences (student's t test, $p < 0.05$).

Figure 2. qRT-PCR analysis of (**a**) GA20ox-1 (**b**) GA20ox-3 (**c**) GA3ox-1 (**d**) GID1b1 (**e**) GID1b2 and (**f**) GAST1 expression in elongating stems and roots treated with 2 mg/L PAC. After 10 days with PAC, plants were treated with different GA_3 concentrations and RNA was extracted three hours later for expression analysis. Values (normalized to ACTIN) are means of four biological replicates ± SE. (**g** and **h**) Stem (g) and primary root (h) length of 14-day-old WT seedlings treated with 2 mg/L PAC followed by GA_3 application (10^{-4}M). Values are mean of 9 plants ± SE. Each set of letters above the columns represents significant differences (Tukey–Kramer HSD, $P < 0.05$). (**i**) Tomato *GID1a*, *GID1b2* and *GID1b2* expression in roots taken from transcriptomic data by Góra-Sochacka *et al.*, 2019. Values are mean of 3 plants ± SE. Each set of letters above the columns represents significant differences (student's t test, $p < 0.05$).

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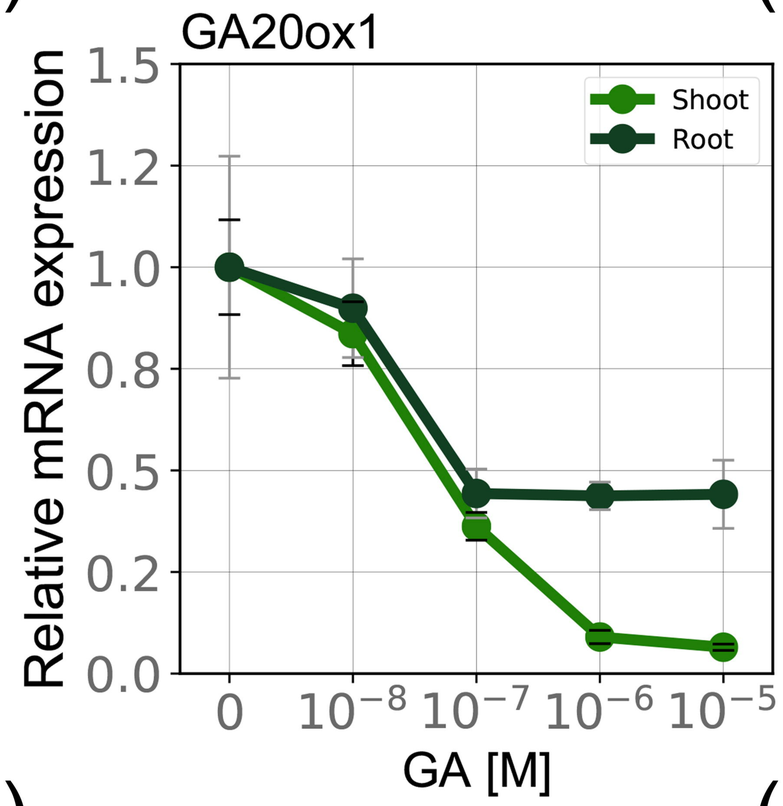
(a)
Stem
Root
Mock
GA
Length (cm)
WT
gib-2
(b)
Stem
WT y = 0.2675x - 0.0403
gid1TRI y = 0.0265x - 0.0215
Length (cm)
Days
(c)
Root
WT y = 0.8963x + 0.0051
gid1TRI y = 0.3163x - 0.065
Length (cm)
Days
(d)
Root-to-shoot ratio
14.1
3.4
WT gid1TRI
(e)
30 ± 0.6
2.8 ± 0.2
20.6 ± 3.4
43.6 ± 2.5
WT
gid1TRI
(f)
Stem length (cm)
GA [M]
(g)
Root length (cm)
GA [M]
(h)
Stem
Cell length (µm)
PAC
GA [M]
(i)
Root
Cell length (µm)
PAC
GA [M]

(a)



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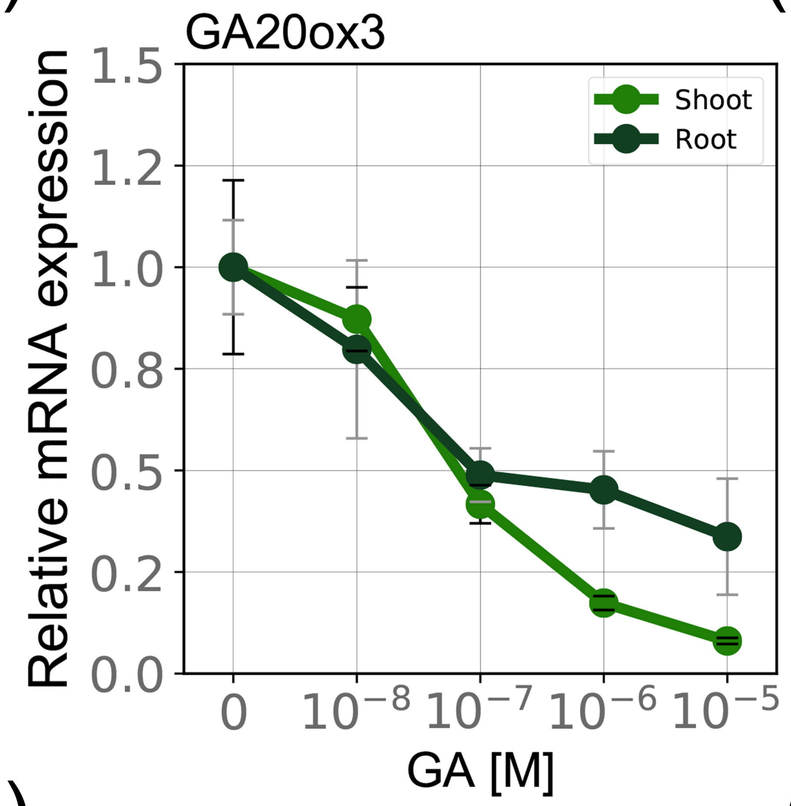

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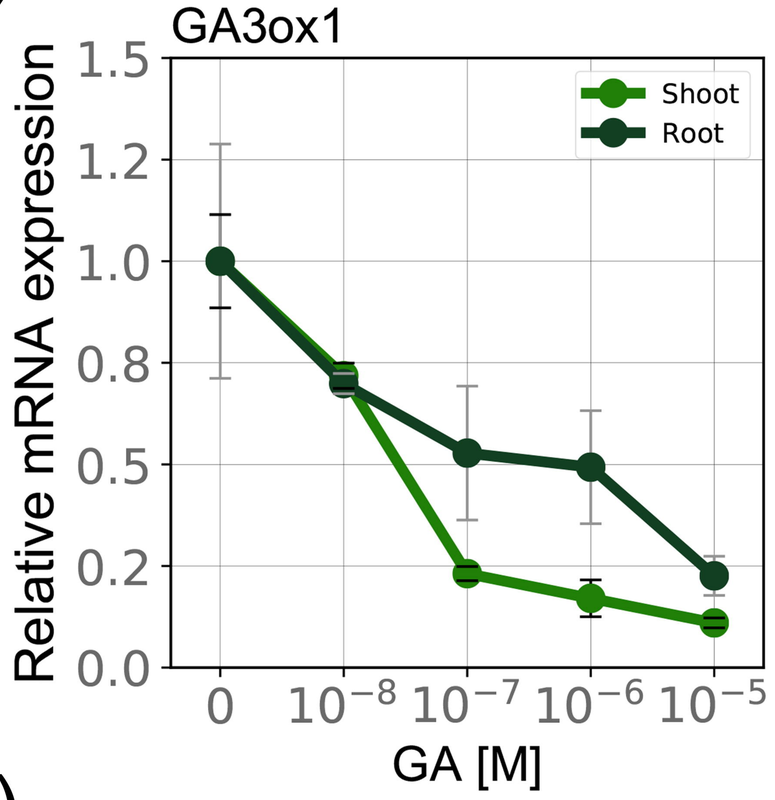

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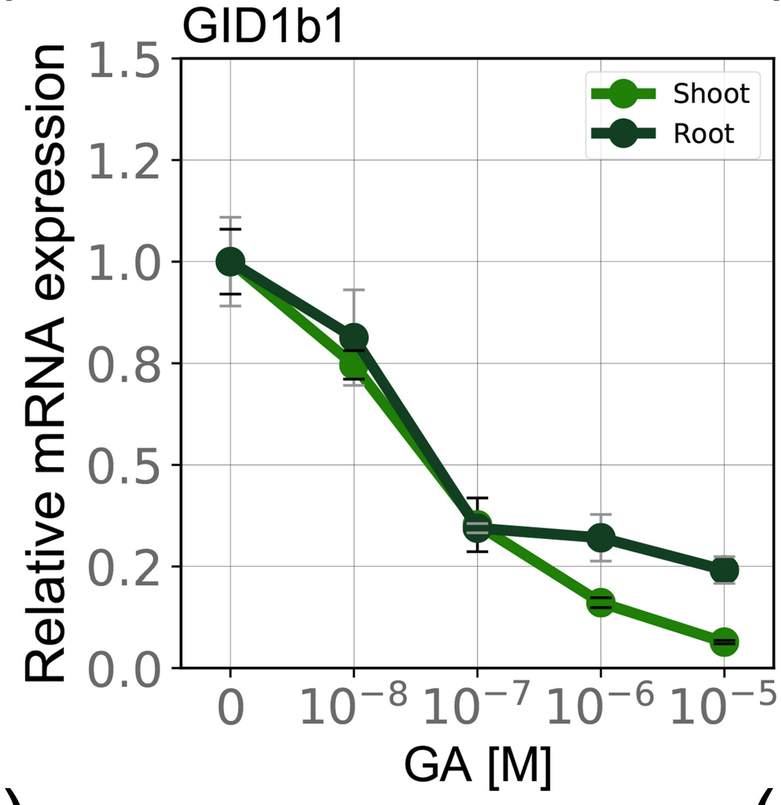

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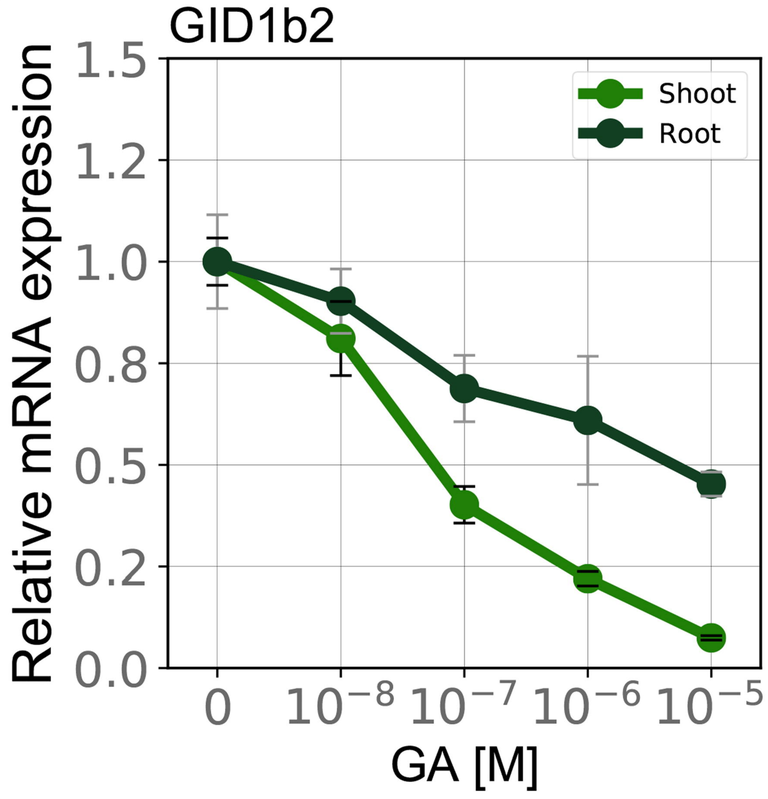

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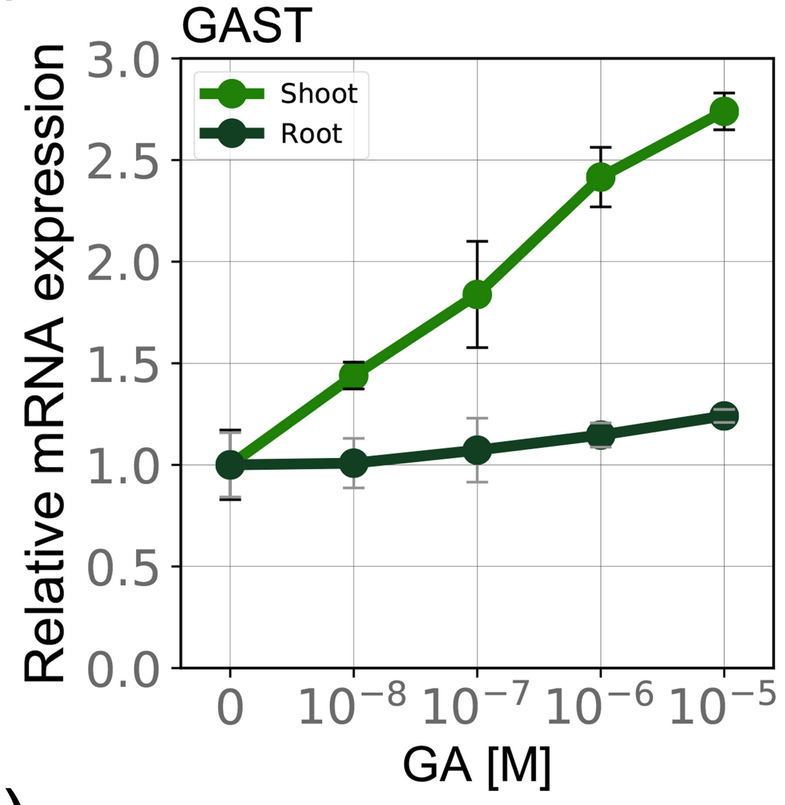
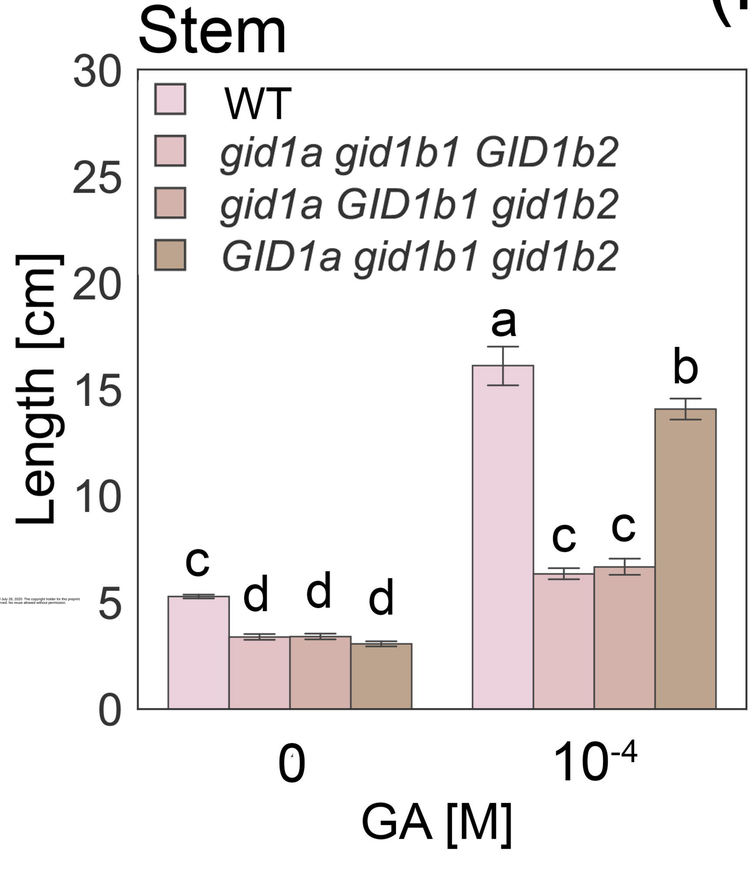
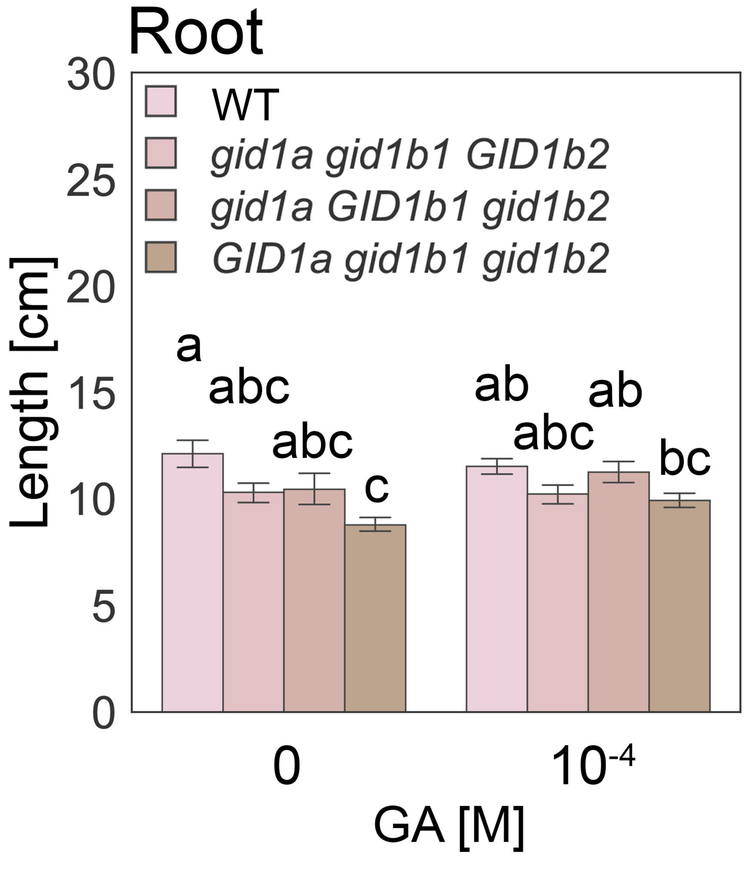


(g)



(h)



(i)