## REGULAR ARTICLE



# Phenotypic response to soil compaction varies among genotypes and correlates with plant size in sorghum

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## **Abstract**

Aims Soil compaction is a major yield-reducing factor worldwide and imposes physico-chemical constraints to plant growth and development. Facing limitations, roots can adapt and compensate for loss of functioning through their plasticity. Being primarily a belowground challenge, tolerance to soil compaction needs to be associated with root phenotype and plasticity. It is therefore of importance to distinguish between size-related apparent and size-independent adaptive plasticity. We determined the above- and belowground plasticity of sorghum genotypes varying in overall plant size.

*Methods* We quantified plasticity as the degree response (adaptive and apparent plasticity) to soil compaction and conducted two experiments with sorghum and two soil density levels (1.4 and 1.8 Mg m<sup>-3</sup>). First, we quantified the shoot biomass plasticity of 28 sorghum genotypes. Second, we studied the root plasticity of six genotypes varying in shoot size and tolerance to soil compaction.

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J. Correa · J. A. Postma · T. Wojciechowski (⊠) Forschungszentrum Jülich, Wilhelm-Johnen Straße, 52425 Jülich, Germany e-mail: t.wojciechowski@fz-juelich.de Results Plasticity was correlated with plant biomass with larger genotypes responding earlier and more intensely. Soil compaction affected roots more than shoots and plasticity was expressed foremost in nodal root number and fine root length. Impeded plants produced 35 and 47% less root mass and length, respectively.

Conclusions Plasticity to soil compaction varies among genotypes, but less-sensitive lines are in general smaller-sized genotypes. The association between tolerance and plant biomass may pose challenges to crop production; however, vigorous genotypes with unresponsive shoots to soil compaction do exist. Maintaining shoot growth relatively stable while the root modifies its structure can be an important adaptation mechanism to soil compaction.

**Keywords** Plasticity · Genotype-by-environment interaction · Plant phenotyping · Root system architecture · Soil bulk density · Soil strength · Phenotypic variation

## Introduction

When a genotype is able to express different phenotypes in different environments, it is called plastic (Bradshaw 1965; Correa et al. 2019; Palmer et al. 2012; Pigliucci et al. 2001; Sultan 1987; Via et al. 1995). In general, the phenotypic responses of plants can be classified into true or apparent plasticity. The



distinction between these two types of plasticity may be challenging. Apparent plasticity is phenotypic variation that is associated with plant size and age (allometry). For example, changes in biomass allocation usually are a function of plant biomass during growth and development (allometric) (Correa et al. 2019). In contrast, plasticity may have an adaptive value, providing tolerance to several environmental constraints (Bradshaw 1965; Des Marais et al., 2013). True plasticity (or just plasticity) can encompass stress, damage, and adaptive responses. Thus, plants are thought to be able to face environmental constraints through their plasticity (Bradshaw 1965; Des Marais et al. 2013; Palmer et al. 2012; Via et al. 1995).

In crop production, stress is any condition that decreases yield (Wallace 1986). Tolerant genotypes are those that survive and produce with comparatively low yield reductions (Negin and Moshelion 2016). Since phenotypic plasticity can be associated with environmental tolerance (Bradshaw 1965; Des Marais et al. 2013), the plasticity for specific phenotypic traits is likely adaptive. The mechanism of tolerance, however, might not be readily observed when they involve belowground or physiological responses. Consequently, tolerant genotypes may be (mistakenly) seen as non-plastic because they hardly express evident phenotypic changes. Plasticity can be described as environment dependent phenotypes of a genotype. Different plant traits can vary in their plasticity to different environments and, depending on the trait, these phenotypic changes may or may not be adaptive. Apparent plasticity is given by ontogenetic effects (the phenotype depends on the size or the state of development of the plant), is relatively easily observed for size-related plant traits (Correa et al. 2019). When the environment affects plant growth, all the size-related plant traits may be influenced concordantly. This correlation between traits and plant size is called apparent plasticity or allometry. Examples for size-dependent apparent plasticity/allometry are found for drought (Blum and Sullivan 1997; Blum et al. 1997), heat (Blum et al. 1997), boron deficiency (MacInnes and Albert 1969), and soil compaction (Coelho Filho et al. 2013). The negative effect of soil compaction on shoot growth was more severe on plants harboring the tall Rht allele than semidwarf and dwarf lines (Coelho Filho et al. 2013) or commercial cultivars (Jin et al. 2015) of wheat. For example, Coelho Filho et al. (2013) showed that the leaf elongation of the taller line was more affected by mechanical impedance of roots than the semi-dwarf line. A GA-insensitive severe dwarf NIL did not show any reduction in leaf elongation by root impedance while having the smallest value of root dry mass among the *Rht* lines.

Soil compaction is a global issue affecting millions of hectares of agricultural lands (Oldeman et al. 1991). Specifically, soil compaction, often caused by heavy traffic of farm equipment, can negatively affect root growth and thereby nutrient and water uptake (Arvidsson and Håkansson 2014; Bengough and Mullins 1990; Bengough et al. 2011; Passioura 2002; Unger and Kaspar 1994). The degree of compaction of a particular soil depends on many soil properties, such as texture, structure, organic matter content, water potential, porosity, etc. (Atkinson et al. 2020; Kolb et al. 2017; Lucas et al. 2019; To and Kay 2005). Furthermore, root systems also affect soil by compacting the surrounding soil as they grow and leaving behind a dense net of biopores as they are degraded (Lucas et al. 2019). Yield losses by compaction have been estimated to be approximately 20 and 25% of total yield in barley and wheat, respectively (Arvidsson 1999; Barken et al. 1987). The lower yields result from reduced uptake of water and nutrients, and lower biomass, which in turn are consequences of soil mechanical impedance on root growth and development (Grzesiak et al. 2014; Håkansson et al. 1988; Lipiec and Stępniewski 1995; Passioura 2002; Stirzaker et al. 1996). The major phenotypic response of the root system to soil compaction is a reduced total root length with a coinciding increase in root diameter (Bingham et al. 2010; Colombi and Walter 2017; Eavis 1972; Goss 1977; Grzesiak et al. 2002; Pfeifer et al. 2014; Popova et al. 2016). Although some observed plasticity may be linked to negative (stress or damage) consequences for plant growth, we suggested that other changes might be adaptive and represent a possible mechanism of crop tolerance to soil compaction. To breed tolerant lines to soil compaction, tolerance needs to be associated with root system phenotype and be adaptive rather than apparent plasticity. Because little is known about the role of plasticity on the phenotypic responses to soil compaction, we studied whether the genotypic diversity in the degree of responses to soil compaction is more dependent on true plasticity than on plant



biomass (apparent plasticity). Additionally, we examine whether those responses are expressed mainly above- or belowground. To answer these questions, we conducted two experiments in controlled environments. As a first step, we carried out a simple experiment to assess how phenotypically diverse a panel of 28 genotypic lines is in terms of shoot biomass and response to compaction. Then, in a larger and longer experiment, we investigated both the above- and belowground response of six genotypic lines which varied in overall size and tolerance to compaction.

#### Materials and methods

We carried out two greenhouse experiments: (1) an initial screen of 28 sorghum genotypes and (2) an in-depth phenotyping of 6 selected genotypes. In the first experiment, we grew the 28 genotypes in compacted and loose soil for 25 days to screen for differential shoot biomass responses to soil compaction and determine the relation between overall plant size (biomass) and the relative response of plant size to compaction. In the second experiment, we studied the above- and belowground phenotypic apparent and true plasticity to soil compaction in six different genotypes. These genotypes differed in their degree of shoot mass plasticity to soil compaction in the first experiment. The two experiments were carried out in the greenhouse facilities of the Institute of Bio- and Geosciences (IBG-2) at the Forschungszentrum Jülich GmbH, Germany (50° 54′ 36" N, 6° 24′ 49" E). A loam field soil (10% clay, 38.6% silt, and 51.4% sand) was uniformly mixed, air dried until constant mass was reached, sieved to 2 mm, and used as substrate. 28 genotypes of sorghum (Sorghum bicolor L. Moench) with diverse geographical, breeding status, and genetic origin were cultivated (Supplementary Table S1). Surface-sterilized seeds were pre-germinated at 21 °C in Petri dishes on moistened filter paper for 48 h. We transplanted those healthy seedlings that had an intact radicle.

## Experiment 1

For experiment 1, 28 genotypes were grown under two soil density treatments, loose and compacted soil. Eight replicates and two plants per

replicate were used for each genotype. Seedlings were planted in pot trays composed of 60 small containers of 0.25 L each. Each container was filled with dry soil according to the following densities: 1.4 and 1.8 Mg m<sup>-3</sup> for loose and compacted treatments, respectively. The soil was compacted homogeneously throughout the pot using a hand hammer and compacted until the required amount of soil would fit in the container. This resulted in a penetration resistance of 0.4 and 1.8 MPa, respectively for dry soil (measured with a hand penetrometer for top layers IB, Eijkelkamp, The Netherlands). After wetting the penetration resistance in the compacted treatment dropped to 1.6 MPa. We transplanted in each container two seedlings of the same genotype. Containers were placed in trays. Each tray had one soil density level and 28 containers with in total 56 plants of 28 genotypes. 16 trays (8 reps, two treatments) were placed in the greenhouse according to a randomized split-block design (see Supplementary Fig. S1A).

Plants grew for 25 days from the 3rd to the 28th of July (from seedling transplanting to harvest). In addition to natural light during the day, supplementary illumination was supplied by mercury lamps (SON-T AGRO 400, Phillips, The Netherlands) every time that light intensity outside the greenhouse was  $< 400 \mu mol m^{-2} s^{-1} during 16 h between$ 06:00 and 22:00 h local time. Environmental conditions during the experiment were: day length of 16 h, day/night air temperatures of ca.  $26.2 \pm 0.03$  $/20.3 \pm 0.02$  °C, and day/night air relative humidity  $47.1 \pm 0.11/66.9 \pm 0.01\%$ . The trays were watered once a week by capillary action from the bottom to the top by putting them on a bigger tray with water until they reached an on average 90% of field capacity (46 and 39 cm<sup>3</sup> per container of gravimetric water content for loose and compacted soil, respectively) with distilled water. Water content at field capacity was determined by weighing trays after they were wet to near saturation and then drained for 48 h. At harvest, the shoots were collected and dried in an oven at 65 °C until constant shoot mass was reached and recorded. Later, six genotypes differing in shoot dry mass and degree of plasticity to soil compaction were selected for the second experiment in which we studied root growth plasticity in relation to shoot responses.



# Experiment 2

In experiment 2, we investigated the above and belowground. In experiment 1, six genotypes were selected that differed in their phenotypic plasticity to soil compaction in terms of shoot biomass. This is to ensure sufficient genotypic and phenotypic variation to be observed in the second experiment using a smaller number of genotypes, a larger pot size, and applying higher levels of soil compaction.

We grew the plants according to a two factorial completely randomized design with six genotypes, two soil conditions, and 12 replicates (n=144 plants). We filled cuboid-shaped pots (volume 4000 cm<sup>3</sup>) at densities of 1.4 and 1.8 Mg m<sup>-3</sup>, for loose and compacted soil, respectively. The soil was compacted with a manual bolt press (Holzmann Dop 3000, Holzmann Maschinen GmbH, Austria). The soil compression yielded penetration resistances of 0.4 and 3.1 MPa, for loose and compacted soil respectively (measured in dry columns with a hand penetrometer for top layers, Eijkelkamp, The Netherlands).

We transplanted one seedling per pot. Plants grew for 45 days from the 4th of September to the 18th of October. In addition to natural light during the day, supplementary illumination was supplied as previously indicated for experiment 1. Environmental conditions during this experiment were: day/night air temperatures of ca.  $22.7 \pm 0.01/19 \pm 0.01$  °C, and day/ night air relative humidity  $50.2 \pm 0.05/63.6 \pm 0.07\%$ . To track shoot development over time, the shoot projected area was measured non-destructively two times per week for each plant individually. The 'ScreenHouse' automated phenotyping platform of IBG-2 was used for this task (more details are found in Nakhforoosh et al. 2016, Supplementary Fig. S1). The soil water content was determined gravimetrically, and the plants were automatically watered throughout the experiment using the 'ScreenHouse' platform. Plants were irrigated with distilled water at 90% of field capacity, 730 and 620 cm<sup>3</sup> of water per pot for loose and compacted soil, respectively. Water content at field capacity was determined by weighing pots after they were wet to near saturation and then drained for 48 h. The volumetric water content at saturation was 0.46 and 0.30 for loose and compacted soil, respectively.

At harvest (45 days old plants), plant height and number of tillers and leaves were evaluated. Then,

the shoot was cut off from the rest of the plant at the substrate surface. Leaf area and stem projected area were measured using a LI-3100C area meter (LI-COR, Inc., Nebraska, USA). Afterward, roots were carefully separated from the substrate and the rest of the soil particles were washed away from the roots with water. Images of scanned roots were analyzed using WinRHIZO Pro image analysis system (Regent Instruments, Inc., Quebec, Canada) to estimate total distribution of root length by root diameter class. The root length was recorded in 25 root diameter classes equally distributed between 0 and 2.5 mm. The root, stem, and leaf dry mass were obtained after drying it in an oven at 65 °C until constant mass. Specific leaf area and specific root length were calculated per plant as the ratio of leaf area to leaf biomass and root length to root dry mass, respectively (for a complete list of traits, see Supplementary Table S2).

# Statistical analysis

A two-way Analysis of Variance (ANOVA) and an Analysis of Covariance (ANCOVA) for each trait were performed. In both analyses, genotype, soil compaction treatment and their interaction (genotypeby-treatment interaction or G×T) were used as factors. For ANCOVA, plant dry mass was added to the ANOVA model and considered as a covariable. We defined the relative effect of a factor on the phenotype of a given trait as the proportion of the total phenotypic variance explained by this factor. This proportion was calculated based on the mean squares of each factor and error according to the ANOVA or ANCOVA model. Additionally, to analyze the allometric relationship between root and shoot biomass, an additional ANCOVA was computed considering the natural logarithm of root dry mass and shoot dry mass as the dependent variable and the covariable, respectively. Before these analyses, the assumptions of normality and homoscedasticity of variances of residuals were evaluated by the Shapiro-Wilks and the Levene tests, respectively. Variables that failed to meet these assumptions were transformed to natural logarithm, after which the tests were repeated successfully. Significant differences among genotypes were compared by the Fisher's test (P < 0.05)using the R package "agricolae" (Mendiburu 2012). Additionally, to test how significant was the treatment within each genotype a two-sample t-test was



performed. To analyze the relationships among the traits, a correlation analysis was carried out based on Pearson's correlation coefficient between traits.

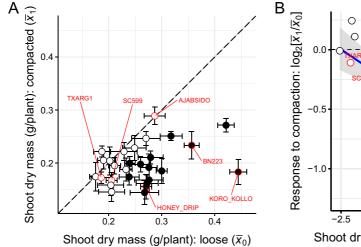
We used the fold change of the logarithm base 2 (FC2) as plasticity index for a simpler analysis of the correlation between plant biomass and the phenotypic plasticity to soil compaction. FC2 has been used for analysis and visualization especially in genomics and bioinformatics (Love et al. 2014). Thus, the phenotypic plasticity of a specific trait was expressed as the FC2 of the ratio of mean value in compacted  $(\bar{x}_1)$  to that in loose soil  $(\bar{x}_0)$  soil for each genotype and trait (plasticity =  $log_2|\bar{x}_1/\bar{x}_0|$ ). This value represents a plasticity index. The FC2 values are close to zero mean no or low plasticity whereas large negative or positive values mean strong plasticity. For example, if the plasticity value for a given trait equals -1, it indicates that the phenotype of that specific trait under compacted conditions was half that of loose conditions. On the other hand, an FC value equal to +1 means that the phenotype of a trait under compaction is twice as large as that found under loose soil. Finally, hierarchical clustering of traits was performed using the R package "ClustOfVar" (Chavent et al. 2017). We analyzed all data using the R statistical programming language (R Core Team 2018). All the graphs were drawn using the "ggplot2" package of R (Wickham 2009).

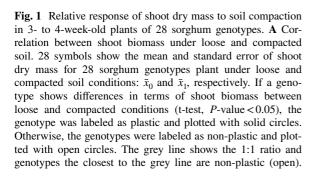
# **Results**

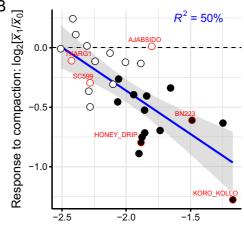
## Experiment 1

Soil compaction reduced the shoot dry mass of most but not all genotypes (Fig. 1A). The compaction treatment explained 89% of the phenotypic response in shoot biomass, whereas genotype and the  $G \times T$  explained 6 and 4% respectively (Table 1).

Genotypes with a significant treatment effect on shoot biomass (t-test, P < 0.05) were labeled 'plastic' (black dots) whereas the others were labeled as







Shoot dry mass (g/plant): loose ( $log_2[\overline{x}_0]$ )

Plastic lines are far below the grey line (solid circles). **B** Correlation between response to soil compaction and shoot biomass. Phenotypic response to soil compaction (y-axis) is the fold change of the logarithm base two of the ratio of mean value in compacted and loose soil for each genotype giving the plasticity index. Negative numbers indicate biomass is smaller under compacted conditions, and -1, indicates biomass in compacted conditions was half that of loose conditions. The blue curve with its confidence interval in grey (at 95%) is the fitted linear regression model between the response and the logarithm base two of the mean value of each genotype under loose conditions



Table 1 Effect of soil compaction on shoot mass of 3- to 4-week-old plants of sorghum (screening experiment)

Treatment	Shoot dry mass (g)			CV (%)	n	Factor R <sup>2</sup> (%)		
	Mean ± SEM	min	max			Genotype	Treatment	G×T
Loose	$0.259 \pm 0.008$	0.035	0.857	47.7	224	5.6***	88.7***	3.6*
Compacted	$0.198 \pm 0.005$	0.012	0.507	41	224			

SEM, min, max, CV, n: standard error of the mean, minimum, maximum, coefficient of variation; number of observations (pot with 2 plants), respectively

R<sup>2</sup>: determination coefficient according to mean square results from two way ANOVA, GXT: Genotype-by-treatment effect. Significant codes (*P* value): <0.001 '\*\*\*', 0.01-0.05 '\*'

'non-plastic' (white dots). Plastic genotypes had on average 28% reduced shoot dry mass compared with the responsive ones (Fig. 1). In Fig. 1B, we expressed the plasticity to soil compaction for each genotype using the proposed plasticity index. The plasticity index is 0 when plants are unresponsive, and negative numbers indicate that biomass is smaller under compacted conditions, and -1, indicates that biomass in compacted conditions was half of measured in loose conditions. Figure 1B shows that response to soil compaction expressed by the plasticity index correlates negatively with shoot biomass in the loose-soil control: The, under controlled conditions, larger-sized genotypes had greater reductions in shoot biomass compared to smaller-sized ones. This association between plant biomass and plasticity was especially clear for 'KORO\_KOLLO' and 'TXARG1': 'KORO\_ KOLLO' had relatively large plants and expressed the highest shoot biomass reduction, while 'TXARG1' was a non-plastic genotype with a relatively small shoot mass. However, 'AJABSIDO' had relatively large shoot but did not respond to compaction and thereby was an exception to the trend.

Based on experiment 1, we selected six genotypes with varying plasticity and shoot mass (Fig. 1, Supplementary Table S1, Figs. S1, S2). For the highly plastic genotypes, we selected the relatively small 'HONEY\_DRIP' (shoots of 0.271 g in loose soil and 43% smaller under compacted soil in experiment 1) and the relatively large 'KORO\_KOLLO' (shoots of 0.444 g in loose soil and 59% smaller under compacted soil). For the intermediate responsive genotypes, we chose the relatively small 'SC599' (shoots of 0.205 g in loose soil and 19% smaller under compacted soil) and the relatively large 'BN223' (shoots of 0.356 g in loose soil and 35% smaller under compacted soil). Finally, for the non-plastic unresponsive

(unresponsive) genotypes, we selected the relatively small 'TXARG1' (shoots of 0.186 g in loose soil and 7% smaller under compacted soil) and the relatively large 'AJABSIDO' (shoots of 0.287 g in loose soil and 0% smaller under compacted soil).

# Experiment 2

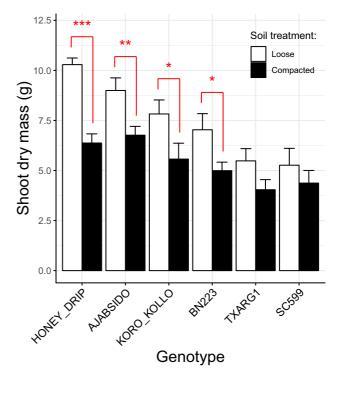
In this second experiment, we grew the plants 20 days longer and, consequently, the shoot mass was on average 25 times greater than in experiment 1. The responsive genotypes in the first experiment were also responsive in the second: 'HONEY DRIP', 'KORO\_KOLLO', and 'BN223'. The only exception was 'AJABSIDO', which was unresponsive in terms of shoot mass in experiment 1 while in experiment 2 it was one of the most responsive lines (reduction in shoot biomass). 'TXARG1' and 'SC599' had the smallest plants and did not respond to soil compaction in either experiment. Importantly, the relationship between plant size and true or apparent plasticity to soil compaction was also strong in the second experiment (Figs. 2, 3, 4, 5; Table 2, Supplementary Fig. S2): the higher the shoot dry mass in loose soil the higher the effect of soil compaction on shoot biomass.

## Plasticity of the root and shoot traits

To summarize the phenotypic plasticity, we plotted a heatmap based on the plasticity index (normalized mean response) for each genotype and trait (Fig. 3). We sorted the traits using a cluster analysis of variables. This yielded four clusters (Fig. 3 top panel, Supplementary Table S2). C1 is mainly made of biomass traits, e.g.: leaf, shoot, stem, and root Dry



Fig. 2 Response of shoot dry mass to soil compaction in 6-week-old plants of six sorghum genotypes. For each genotype, white and black boxes indicate the mean value of shoot dry mass for plants growing in loose and compacted soil. The genotypes are sorted on the x-axis and ranked according to their phenotypic mean under loose conditions, from the largest (left) to the smallest (right). Error bar is the standard error of the mean. Significant results are highlighted in red according to the t-test between loose and compacted conditions. Significance codes (P-value): < 0.001 '\*\*\*'; 0.001-0.01 \*\*\*; 0.01-0.05 \*\*\*



Mass. C2 comprised length of coarse roots (diameter  $\geq$  2.3 mm), root diameter, and collar traits (number of nodal roots, root to shoot ratio, collar dry mass, etc.). C3 contained those variables of root length whose diameter was 1.1–1.9 mm. C4 comprised length of roots with diameter less than 1.1 mm, leaf area, and the number of tillers. The average plasticity indices for these clusters were 0.18, -0.34, -0.47, and -0.48 for C1, C2, C3, and C4, respectively.

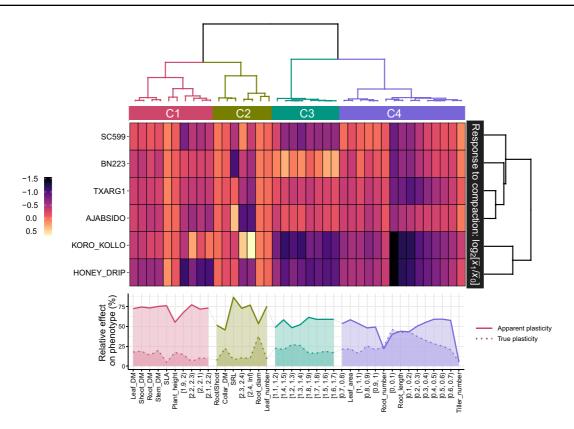
Depending on the genotype, most of the traits were affected by soil compaction (as indicated by negative values of plasticity index and dark colors in Fig. 3) and all of them had a significant genotypic effect (Supplementary Table S3). Genotypes were sorted based on their plasticity index by hierarchical clustering (right side). Sorted by their average plasticity index among all the traits, genotypes ranked as follows: (1) 'HONEY\_DRIP', (2) 'KORO\_KOLLO', (3) 'TXARG1', (4) 'SC599', (5) 'AJABSIDO' and (6)'BN223' (with -0.6, -0.5, -0.36, -0.3, -0.3, and – 0.2, respectively). Cluster C4, comprised of the lengths of roots with diameter classes  $\leq 0.2$  mm, had the most responsive traits (the darkest colors in the heatmap) with an average plasticity index of  $\sim -1.0$ .

Apparent and true plasticity

Based on the ANCOVA (Supplementary Table S4), we plotted to what extent the trait's plasticity was explained by plant biomass (allometric or apparent plasticity), and to what extent it was independent of plant biomass and thereby true plasticity (bottom panel of Fig. 3). Within cluster C4, the length of very fine roots (diameter < 0.1) had a true plasticity effect greater than the apparent plasticity. For example, this trait had both a plasticity index of -1.5 in 'HONEY\_DRIP' and 'KORO\_KOLLO'.

On average, very fine roots under compacted soils were 54% shorter than under loose conditions. On the other hand, biomass-related traits, SLA, plant height, root average diameter per plant, and length of thicker roots (diameter > 1.9 mm) (clusters C1 and C2) were less responsive to soil compaction than very fine root traits. C2 and C1 were the clusters with relatively high apparent plasticity (explaining ~70 and 62% of the total variance, respectively) and the lower true plasticity (explaining ~13 and 15% of the total variance, respectively). While C4 and C3 had lower apparent plasticity





**Fig. 3** Response to soil compaction of shoot- and root traits in 6-week-old plants of six sorghum genotypes. The heatmap shows the mean plasticity (degree of response) expressed as the standardized fold change of the logarithm base two of the ratio of mean value in compacted  $(\bar{x}_1)$  to that in loose soil  $(\bar{x}_0)$  soil for each genotype (rows) and trait (columns). Dark and light colors indicate high and low plasticity, respectively. The relative effect on phenotype of plasticity and apparent plasticity (line graph at the bottom) is based on ANOVA for each trait considering genotype, compaction treatment, plant dry mass (as a covariable) and their interactions as factors. The relative

(explaining ~ 45 and 53% of the total variance, respectively) and higher true plasticity (explaining ~ 28 and 21% of the total variance, respectively; Fig. 3; Supplementary Table S4).

# Root/shoot ratios

Root biomass was correlated with shoot biomass, root/shoot ratios, leaf area, and root length both in loose (r > 75%; Supplementary Table S5) and compacted soil (r > 75%; Supplementary Table S6). Under compaction, root biomass was on average reduced by 35% compared to the loose control.

effect is calculated by using the mean squares of each of these factors. Thus, plasticity is the sum of the importance of treatment and treatment-by-genotype interaction effects; apparent plasticity (allometric effect) is the sum of the importance of plant dry mass and all their interactions with treatment, genotype, and genotype-by-treatment interaction effects. Traits are sorted according to a variable clustering located on the top of the heatmap. Genotypes are sorted according to hierarchical clustering of their response to soil strength, displayed on the right side of the heatmap

Genotypes differed in root biomass but the  $G \times T$  interaction was not significant (Supplementary Table S3).

Compaction reduced root/shoot ratios by 11% on average and the largest proportion of variation was explained by genotypic effect (Supplementary Table S3). The log-log (allometric) relationship between shoot and root biomass across replicates and genotypes (Table 3) was significantly different between soil conditions. For every one percent change in shoot mass, there was a 1.37 and 1.7% change in root mass for compacted and loose soil treatments, respectively (see slopes in Table 3).



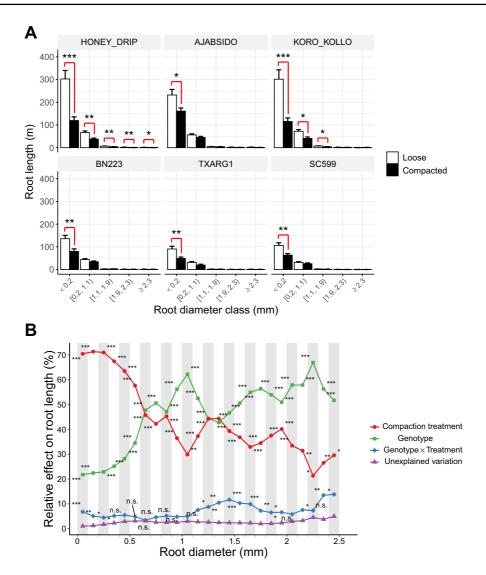


Fig. 4 Response of root length to soil compaction in 6-week-old plants of six sorghum genotypes. A Distribution of root length by root diameter class. The root length was recorded in five root diameter classes ranging from 0 to 2.5 mm. These five classes were based on cluster analysis (see Fig. 3). Genotypes are sorted according to Fig. 1. White and black boxes: mean of root length (mm) for each diameter class in loose and compacted soil conditions, respectively. Error bar: standard error of the mean. The significant results are highlighted

in red according to the t-test between loose and compacted conditions. **B** Relative effect of genotype and soil compaction treatment on root length for each diameter class. Root length in 25 root diameter classes ranging from 0 to 2.5 mm. Diameter classes are indicated as grey or white vertical bands. The relative effect is calculated by using the mean squares of each of these factors. Significance codes (P-value): <0.001\*\*\*\*; 0.001-0.01\*\*\*; 0.01-0.05\*; 0.0

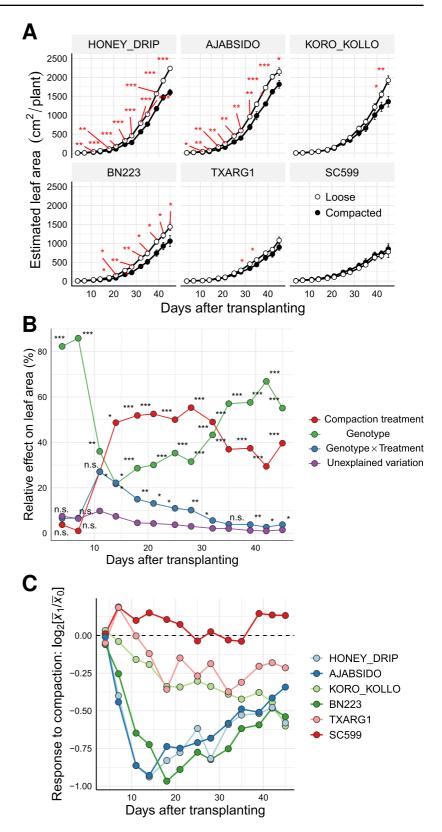
## Length of fine and coarse roots

Root length (Table 4) ranged from 383 m of 'KORO\_KOLLO' under loose soil to only 70 m of 'TXARG1' in compacted soil. Soil compaction reduced root length of all genotypes (46% of

reduction on average). For example, roots of 'KORO\_KOLLO' and 'HONEY\_DRIP' were almost 58% shorter under compacted than in loose soil conditions. Even unresponsive shoot genotypes to compaction such as 'SC599' had an important reduction in root length (35% shorter roots under compacted soil).



Fig. 5 Response of leaf area to soil compaction over time in 6-week-old plants of six sorghum genotypes. A Estimated leaf area over time for the six selected genotypes growing in loose and compacted soils. The measurements were done twice a week yielding in total 13 date points (days after transplanting). Genotypes are sorted according to their leaf area at harvest under loose conditions (from the largest to the smallest): 'HONEY\_DRIP', 'AJAB-SIDO', 'KORO\_KOLLO', 'BN223', 'TXARG1', and 'SC599'. White and black circles: mean of estimated leaf area (cm<sup>2</sup>) for loose and compacted soil conditions, respectively. Error bar: standard error of the mean. The significant results are highlighted in red according to the t-test between loose and compacted conditions. B Relative effect of genotype and soil compaction treatment on leaf area for each measurement date. The relative effect is calculated by using the mean squares of each of these factors. C Variation over time of the leaf area response to soil compaction for each genotype. The response is the fold change of the logarithm base two of the ratio of mean value in compacted and loose soil for each genotype and date point. Significance codes (*P*-value): < 0.001 '\*\*\*'; 0.001-0.01 "\*\*"; 0.01-0.05'\*'; n.s. not significant (P > 0.05)





**Table 2** Genotypic diversity of plant dry mass (g) in loose and compacted soil

Genotype	Loose		Compacte	Within-		
	Mean ± SEM <sup>1</sup>		Mean ± SEM		genotype treatment effect <sup>2</sup>	
HONEY_ DRIP	13.1±0.58	a	7.8±0.63	ab	***	
AJABSIDO	12.9±0.95	a	9.3±0.73	a	**	
KORO_ KOLLO	10.7±1.05	ab	7.8±1.24	ab	n.s.	
BN223	$8.7 \pm 1.07$	bc	$6.2 \pm 0.58$	ab	n.s.	
TXARG1	$6.6 \pm 0.77$	c	$4.7 \pm 0.61$	b	n.s.	
SC599	$6.2 \pm 1.02$	c	$5.0 \pm 0.71$	b	n.s.	

<sup>&</sup>lt;sup>1</sup>Different letters indicate means  $\pm$  standard error of the mean (SEM) with statistically significant differences among genotypes according to Fisher's least significant difference test (P<0.05).

Additionally, total root length was significantly correlated with plant biomass (r=72 and 83% in loose and compacted soil, respectively) and 38% of the observed phenotypic variation in root length was explained by the variation in plant biomass, and the slopes of the regression depended significantly on genotype and treatment (see for ANCOVA in Supplementary Table S4).

We split root length into five root diameter classes (Fig. 3 for cluster analysis; Fig. 4 for root length). The classes with smaller diameters had much greater root lengths than those with thicker diameters (Fig. 4A). We refer to "fine roots" as the class of roots with diameters ≤0.2 mm. Unlike leaf area development, all the genotypes responded to the soil compaction treatment with a reduced root length in one or more root diameter classes: especially, all genotypes had reduced length in the fine root class in response to soil compaction. Genotypes 'HONEY\_DRIP' and 'KORO\_KOLLO', had large shoot dry mass response, also the greatest root length decrease

Table 3 Relative contribution of soil treatment and shoot biomass to the variation of root biomass

Regression coefficients for each treatment				Effects on root dry mass (%) <sup>2</sup>			
Treatment (T)	$R^2 (\%)^1$	intercept	slope	T	Shoot_DM	T ×Shoot_DM	
Loose	85.4	-3.07	1.70	8.0***	90.9***	1.0*	
Compacted	76.6	-2.39	1.37				

 $<sup>^{1}</sup>R^{2}$ : determination coefficient according to linear regression model within each treatment:  $ln(Root\_DM) = intercept + slope \times ln(Shoot\_DM)$ 

Table 4 Genotypic diversity of total root length (cm) in loose and compacted soil

Genotype	Loose		Compacted	Within-genotype treatment effect <sup>2</sup>	
	$Mean \pm SEM^1$		Mean ± SEM	Mean ± SEM	
KORO_KOLLO	38329.7 ± 4976	a	16239.1 ± 2253	ab	**
HONEY_DRIP	$37877.2 \pm 4588$	a	$16299.7 \pm 2089$	ab	**
AJABSIDO	$29674.6 \pm 2891$	ab	$21251.0 \pm 1847$	a	*
BN223	$18374.1 \pm 1882$	bc	$11808.5 \pm 1411$	bc	*
SC599	$14010.4 \pm 1512$	c	$9046.5 \pm 1047$	c	*
TXARG1	$12416.1 \pm 1628$	c	$7012.9 \pm 1016$	c	*

<sup>&</sup>lt;sup>1</sup>Different letters indicate means  $\pm$  standard error of the mean (SEM) with statistically significant differences among genotypes according to Fisher's least significant difference test (P<0.05) within each soil treatment level.



<sup>&</sup>lt;sup>2</sup>The significant codes are according to the t-test between the means under loose and compacted conditions for each genotype (P value): <0.001 '\*\*\*';0.001-0.01 '\*\*\*'; n.s.: notsignificant (P > 0.05).

<sup>&</sup>lt;sup>2</sup>Determination coefficient according to mean square results from ANCOVA

<sup>&</sup>lt;sup>2</sup>The significant codes are according to the t-test between loose and compacted conditions within each genotype (*P* value): 0.001-0.01 '\*\*', 0.01-0.05 '\*'

under compacted soil: about 57–58% shorter roots, and showed responses to soil compaction for almost all the root diameter classes. On the other hand, the smaller genotypes, 'BN223', 'TXARG1' and 'SC599', had significant effects on fine roots only. This was also true for the larger 'AJABSIDO'. Overall, the plant biomass explained 33–38% of the phenotypic variation of the fine root length. Root length of thicker roots correlated stronger to plant biomass: plant biomass explained 65% or more of the phenotypic variation in the length of roots with diameters > 2 mm (Supplementary Table S4).

Figure 4B shows the relative effect of soil compaction and genotype on the root length for each diameter class. Treatment had a relative effect of 50–70% for roots whose diameters were less than 0.6 mm. In the case of fine roots, this effect explained about 70% of the variation in root length.

The second most important explanatory variable was genotype, especially for thicker roots. This interaction was significant in almost all the diameter classes ranging between 5 and 15% of the total variation. Similarly, this interaction explained ~8% of the total variation in root length.

## Leaf area development

To track the effect of soil compaction on the development of shoot, we estimated leaf area development non-destructively based on color images of our plants. Green pixel count was validated against measured leaf area at harvest ( $R^2 = 99\%$ , RMSE=156.3 Supplementary Fig. S3). Figure 5A shows the increase in estimated leaf area over time based on this validation. Genotypes had different total leaf areas at harvest in loose soil (from the largest to the smallest): 'HONEY\_DRIP', 'AJABSIDO', 'KORO\_KOLLO', 'BN223', 'TXARG1', and 'SC599'. The four genotypes with a significant effect on shoot dry mass (Fig. 2) also responded to soil compaction in terms of leaf area. In addition to being the biggest genotypes in terms of leaf area and shoot dry mass, 'KORO\_ KOLLO' and 'HONEY DRIP' had the largest reduction in leaf area (32 and 29% smaller values under compacted soils; Fig. 5A) and shoot biomass (28 and 38% smaller values under compacted soils; Fig. 2).

Leaf area variation was mainly influenced by genotypic differences during the first days of growth explaining almost 80% of the total variation. The treatment effects on larger genotypes became evident during the second week after transplanting (Fig. 5A, B). At this time, the treatment became the more important source of variation explaining almost 50% of the variation in leaf area. Genotype became again the most explanatory factor during later stages.

As in Figs. 1 and 2, larger genotypes had the higher leaf area responses (Fig. 5C). 'KORO\_KOLLO' and 'TXARG1' only had significant treatment effects 30 days after transplanting. For these two genotypes, the treatment effect accelerated almost constantly until harvest time (Fig. 5C). On the other hand, Fig. 5B and C show that the degree of leaf area response decreased with time from 20 days after transplanting onward, specifically for 'AJABSIDO', 'BN223' and 'HONEY\_DRIP'. On the contrary, 'SC599' did not respond to compaction and consequently, its plasticity index was very close to zero (Fig. 5C).

#### Discussion

Overall, compaction reduced plant size but large variation among genotypes existed. Additionally, the plasticity to compaction was correlated to plant size. Finally, we found that the  $G \times T$  explained a small portion of the observed variation next to the effect of plant biomass. We have observed genetic variation in both above- and belowground plasticity to soil compaction. Some of this variation may be explained by plant size (allometric), and thereby a form of apparent plasticity (Correa et al. 2019; Weiner 2004). Phenotypic plasticity varied among genotypes and depended on biomass, with larger genotypes responding at earlier developmental stages and more intensely. We observed not only genetic variation for phenotypic response/plasticity in shoot size-related parameters, but also for plasticity in various root traits. Although all traits were correlated to plant size, especially the number of nodal roots and fine root length had strong true plasticity. Larger trials, however, are necessary to determine the heritability of true plasticity to soil compaction of those traits.



# Responses associated with plant size

In the second experiment, we grew the plants 20 days longer in bigger pots and, consequently, the shoot biomass was on average 25 times higher than in experiment 1. Plastic genotypes in the screening also were plastic in experiment 2 but shoot dry mass was sometimes inconsistent between experiments 1 and 2 (Fig. 2; Supplementary Fig. S2) At harvest, the roots of experiment 2 were able to colonize the entire pot, which may have reduced the impact of compaction for a small proportion of roots that were growing in areas with decreased penetration resistance, e.g. at the pot walls boundaries. Despite this, it is important to note that the relationship between plant size and response to soil compaction was also observed in both experiments (Figs. 1, 2, 3, 4, 5): the higher the shoot dry mass in loose soil the higher the effect of soil compaction on shoot biomass.

In general, larger genotypes were more responsive than those genotypes with smaller plants (Figs. 1, 2, 3, 4, 5). Selecting genotypes with different degrees of response had as a consequence that different plant sizes were also co-selected.

It is difficult to distinguish true or apparent plasticity because root to shoot ratio decreases as soil density increases and is correlated with plant mass. It is known that smaller or younger plants generally have comparatively greater root/shoot ratios (McConnaughay and Coleman 1999; Weiner 2004). On the contrary, we found that plants under compaction had lower root/shoot ratios despite being smaller than plants in loose soils. Additionally, the number of leaves on the main axis was not significantly affected by compaction. The log-log relationship between shoot and root biomass (Table 3) showed soil treatments had different slopes indicating different allocation patterns: an indication for true plasticity (Reich 2002). The slopes mean that a one percent decrease in shoot mass was associated with a 1.37 and 1.7% decrease in root mass for compacted and loose soil treatments, respectively. Therefore, plants growing in compacted soil have proportionally less root than shoot mass compared with non-impeded plants. This decrease in root/shoot ratios is accentuated by the fact that the plants are smaller and that smaller plants normally have increased root/shoot ratios. A decreased root/shoot ratio means that genotypes showed different biomass partitioning in favor of shoots under compacted soil.

# Root responses

Almost 75% of the total root length was represented by fine roots whose diameters were less than 0.2 mm. Fine roots were not only the most responsive component of the total length of the root to soil compaction (Figs. 3, 4) but also the most responsive trait (Fig. 5; Supplementary Table S3). Given their functional importance, we may assume that the reduction of fine root length has a great impact on root functioning. Due to their greater surface area per unit volume, fine roots are the principal pathway for nutrient and water uptake (Comas et al. 2013; Eissenstat 1992). Additionally, they have significantly higher rates of respiration, associated with higher N concentrations, compared with thicker roots (Eissenstat and Yanai 1997; Pregitzer et al. 1998) and a relatively short lifespan, rapid turnover, and quick decomposition (Jackson et al. 1997). Although fine root production is likely mechanically impeded in compacted soils, they may also be hampered by the low availability of soil resources such as N and oxygen (Bengough et al. 2011; Håkansson et al. 1988; Passioura 2002; Tubeileh et al. 2003). Based on that, the observed reduction of fine roots may be related to an optimization strategy of carbon and/or soil resources and as such adaptive plasticity. Plastic and tolerant genotypes could avoid producing fine roots not only because of their high cost under impeded conditions but also because they may be less efficient under compaction. Plants likely have mechanisms to compensate for a shorter root system by increasing the root uptake efficiency especially in those genotypes with unresponsive shoots. Further studies are needed to assess how greater resource acquisition efficiency can compensate for the loss of fine root length.

# Shoot responses

A common response to soil compaction is the reduction in shoot mass and leaf area of plants (Beemster and Masle 1996; Grzesiak et al. 2014; Masle and Passioura 1987). The same was found in our experiments (Tables 1, 2; Figs. 1, 2, 5). The soil strength levels applied in this experiment (>3 MPa) are considered highly limiting for root growth (Bengough et al. 2011; Passioura 2002; Pierce et al. 1983) and affected shoot growth of plants younger than 4 weeks (Figs. 1, 2; Supplementary Fig. S2).



Shoot plasticity to soil compaction varied among genotypes (Figs. 1, 2, 5), but we did not find a clear association between observed shoot phenotype and the genotype's origin or breeding status (ANOVA: *P*-value>0.05). The genotypic variation in shoot responses to soil compaction was correlated to plant size under loose soil (Figs. 1, 2). Hence, genotypes with larger plants under loose soil had greater reductions in leaf area than smaller sized genotypes under soil compaction.

The effects on shoots were evident from the second week after transplanting onward (Fig. 5). This early response is in agreement with previous observations on seedlings and young plants of wheat and barley growing in compacted soil (Goss and Russell 1980; Masle 1992; Masle and Passioura 1987; Rebetzke et al. 2014). The early response may be a factor to be considered as early vigor is key to the establishment and survival of seedlings especially under conditions of topsoil compaction. The increased soil strength by soil crusting, i.e. a formation of a seal at the soil surface, affects negatively both seedling emergence and establishment (Awadhwal and Thierstein 1985; Nortjé et al. 2012). Thus, seedling establishment of highly responsive lines may be severely reduced and may need to be compensated for by higher seeding rates. This is especially relevant for sorghum since it is said to be sensitive to crusting (Awadhwal and Thierstein 1985).

## Genetic diversity in response to soil compaction

Genetic diversity in the plasticity of plants to soil compaction has been documented in other studies (Arvidsson and Håkansson 2014; Colombi and Walter 2017; Grzesiak et al. 2014; Materechera et al. 1992; Rebetzke et al. 2014). If plasticity is adaptive and has a clear genetic basis, it can be useful for breeding. Therefore, the study of both genetic variation and the contribution to crop yield of these root responses is essential. Sorghum has wide genotypic diversity (e.g. Sinha and Kumaravadivel 2016). By using varieties with different geographical and genetic origin and breeding status, we expect that our panel represented this diversity (Supplementary Table S1). We found that the genotypic effect accounted for a large portion of the observed phenotypic variation (Tables 1, 2, 3; Supplementary Tables S4, S5). We expected that different sorghum genotypes differ in their plasticity to soil compaction, and indeed found a significant  $G \times T$  for several traits, notably root length. Other traits with significant  $G \times T$  were the number of nodal roots and tillers. In our second experiment, roots were more affected than shoots (Fig. 3). For example, the observed reduction in both root length and biomass in response to soil compaction was approximately 50 and 35%, respectively. While the reduction observed in terms of shoot biomass and leaf area was less than 30%. Therefore, sorghum genotypes differ in their plasticity to soil compaction  $(G \times T)$  and this might be exploited in future breeding programs.

Unresponsive lines growing in compacted soil have a relatively lower root/shoot ratio and a smaller proportion of fine roots. A reduced proportion of fine roots could be a sign of 'damage' but may be true adaptive plasticity if reduced investment into fine roots is accompanied by increased investment into other resource acquisition strategies resulting in greater uptake efficiency. However, more studies are needed to establish whether greater root absorption efficiency would compensate for the reduction in fine root length and its relationship with yield. Maintaining shoot growth relatively stable while the root modifies its structure can be an important adaptation mechanism to soil compaction.

The degree of response to soil compaction was correlated with plant biomass (Tables 1, 2, 3; Figs. 1, 2, 3, 4, 5). In general, larger genotypes such as 'KORO\_KOLLO' and 'HONEY\_DRIP' were the more plastic and displayed the higher and earlier response to soil compaction in terms of length of fine roots and leaf area than smaller plant genotypes such as 'TXARG1' and 'SC599'. On the other hand, 'AJABSIDO', a "drought-tolerant landrace" from Sudan (Supplementary Table S1), was a genotype that was relatively large and had intermediate plasticity. Shoot and root biomass of 'AJABSIDO' in compacted soil were reduced by 25 and 35% compared to the loose control, respectively. We know that penetration resistance increases as soil water potential decreases (Bengough et al. 2011; Whalley et al. 2005). It is likely that tolerance to compaction and drought evolved together and share some physiological or structural mechanisms of the plant to adapt to higher soil resistance conditions. For example, it has been observed that some maize and triticale genotypes that were tolerant to soil compaction were also drought tolerant (Grzesiak et al. 2014). 'AJABSIDO'



may thus be an interesting "ideotype" for both compaction and drought tolerance.

In general the genotypes of larger plants are proportionally more affected than smaller ones by soil compaction. One might conclude that tolerance to soil compaction cannot be exploited if it is associated with overall reduced vigor. However, high yields can be obtained using less vigorous genotypes by growing them at higher planting densities, assuming harvest index of the less vigorous lines is not reduced. We suggest that the extreme tolerance of 'TXARG1' and 'SC599' may still be exploited in breeding.

Root system plasticity to soil compaction is complex and involves dynamic changes and several interactions among root traits that were not addressed in our study. For example, greater production of thicker and steeper roots may be of advantage (Correa et al. 2019). This allows a better soil exploration that could be even more efficient if the roots are able to find those paths and patches with the least resistance to penetration. In those soil patches, some resources, such as oxygen and nitrogen, may be more available than in their surroundings. The proliferation of roots in these patches can be a way to compensate for the loss of root length.

## Conclusions

As long as the plasticity is adaptive and has a clear genetic basis, it will be useful for breeding. Sorghum genotypes differ in plasticity to soil compaction  $(G \times T)$  and, indeed, we found a significant  $G \times T$  for several traits, particularly root length. This genetic diversity in plasticity could be exploited in future breeding programs. However, plant size explained a larger proportion of the trait variation than  $G \times T$ , which can pose challenges in plasticity-based breeding and requires further research, especially if the tolerant but smaller sized genotypes can have high yields when grown at greater plant densities. To the best of our knowledge, this is the first study that illustrates how the phenotypic responses to soil compaction correlate with the potential plant size of a genotype in sorghum. Finally, the observed phenotypic changes in response to soil compaction are complex, both apparent plasticity (allometry) and plasticity are involved but apparent and true plasticity are distinguishable and measureable across genotypes. The association between tolerance and plant size can pose challenges in achieving high yields through breeding and requires further research. In general, root traits were more plastic than shoot traits. Plasticity was expressed foremost in nodal root number and fine roots, whereas thick root length was much less affected and more correlated with plant biomass. Fine roots were the most plastic component of root total length and the most plastic overall plant traits.

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Plant responses to soil compaction can be explained to a large degree by allometry that is the responsive and unresponsive genotypes are relatively small and large, respectively. Nevertheless, true plasticity (size-independent responses) was observed especially for the number of nodal roots and root length of fine roots, but also for biomass allocation patterns.

'AJABSIDO' was a genotype that was relatively large and had intermediate plasticity responses showing the best growth under compacted conditions and therefore, may serve as an 'ideotype' in breeding to mitigate the constraints of compacted soils.



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#### **Declarations**

**Conflict of interest** The authors declare no conflict of interest.

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