

Original research article

Genetic variation in the gravitropic response of maize roots to low temperatures

Andreas Hund

Institute of Plant, Animal and Agroecosystem Sciences, ETH Zurich, 8092 Zurich, Switzerland Corresponding author: A. Hund, Email: andreas.hund@ipw.agrl.ethz.ch, Phone: +41 44 632 3829, Fax: +41 44 632 1143

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Abstract: The distribution of roots in soil determines their acquisition of spatially varying resources. It may be altered by changing the response of roots to gravity. The aim of the study was to assess gravitropic set-point angles (GSAs) of maize (Zea mays L.) roots, their response to temperature and the feasibility to measure them in growth pouches. The GSAs of the primary, seminal and crown roots of a set of nine temperate inbred lines were measured. The lines were grown under controlled conditions in growth columns either at 15/13°C or 24/20°C (day/night) until the two-leaf stage (V2). The GSA was measured as the deviation of the initial 3 cm of root axis from the vertical zero. Low temperature resulted in a decrease in the GSAs of the crown roots by 10°, i.e. the roots oriented more vertically. The effect of the GSAs on the distribution of the roots was verified in wider columns using two extreme inbred lines. The proportion of roots in the upper 5 cm of the columns was 78% for the line S335 with the strongest tendency to horizontal root growth and only 39% for CM105 with almost vertical orientation of the roots. The differences in GSAs between these two genotypes were even more pronounced in growth pouches, thus proving the feasibility of this system for rapid screening. The results indicate that there is a huge genetic variability available to alter the growth direction of the seedling roots of maize. However, there was little effect of the temperature.

Keywords: corn, gravitropism, liminal angle, plagiogravitropism, root angle, *Zea mays* L.

Abbreviation: GSA, gravitropic set-point angle

Introduction

Variation in root architecture is essential for the adaptation of plants to target environments since it determines their efficiency in acquiring soil resources. One parameter determining root architecture and root occupancy of the soil is the gravitropic set-point angle (Digby and Firn 1995), sometimes also called 'liminal angle'. It is defined as the angle, at which an organ is maintained by gravitropism and is measured as the deviation from the vertical zero. Roots may grow vertical (orthogravitropic) oblique (plagiogravitropic) or horizontal (diagravitropic) in response to gravity. This response may be altered by environmental stimuli. Ge et al. (2000) illustrated the implication of changes in the GSA for the acquisition of phosphorus and/or water using the example of common bean (Phaseolus vulgaris L.). For this species, increased GSAs led to an increased uptake of phosphorus due to more intense foraging of the topsoil (Liao et al. 2001). For other species, decreased GSAs led to a better water acquisition due to deeper roots, for example in upland rice (Kato et al. 2006), sorghum (Tsuji et al. 2005), wheat (Manschadi et al. 2008) and maize (Hammer et al. 2009).

Apart from enhancing the acquisition of resources, greater GSAs may also facilitate the escape from low subsoil temperatures. It was observed already in the late 19th century that maize roots grew more horizon-tally early in the season than later (Three references cited by Onderdon and Ketcheso, 1973). While the authors of these early studies assumed that the temperature gradient was responsible for this effect, later studies identified temperature *per se* as the cause. GSAs were largest at about 18°C. They decreased with increasing temperatures from 18 to 36°C (Mosher and Miller 1972, Onderdon and Ketcheso 1973) and also with decreasing temperatures from 17 to 10°C (Onderdon and Ketcheso 1973). The temperatures during the early developmental phase, when the

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roots were less than 10 cm long, determined their later trajectory in the field (Tardieu and Pellerin 1991): the horizontal spread of these roots decreased exponentially with increasing soil temperature. This temperature effect may explain the decrease in the GSAs of crown roots from higher internodes, which develop in warmer soil later in the season (Araki et al. 2000, Kaeriyama and Yamazaki 1983, Tardieu and Pellerin 1990).

But are there genetic differences in the GSAs of maize roots and their response to temperature? There is not much information available since experiments were usually conducted with one genotype, only. Distinct phenotypes for nodal root angle were found by Irwin et al. (1985) for 11 hybrids. Quantitative trait loci for root angles were mapped in the context of lodging resistance of maize (Barriere et al. 2001, Guingo et al. 1998) and flooding tolerance of a maize × teosinte cross (Omori and Mano 2007). Furthermore, two maize mutants showed pleiotropic effects between branching pattern and gravitropic response (Taramino et al. 2007, Woll et al. 2005).

The lack of information about genetic differences for root architecture, including GSAs, may be attributed to the difficulty to assess the trait. This situation is rapidly changing: non-invasive imaging techniques open new possibilities for root phenotyping of large germplasm sets. Imaging is becoming more and more popular since the computer processing power, the resolution of imaging sensors and affordable disc space increased dramatically during the last decades. In parallel, two dimensional growth systems were developed allowing for a non-destructive imaging of the roots. For example, the distribution of roots in two dimensions was studied in gel chambers (Bengough et al. 2004), root observation chambers (Manschadi et al. 2006), on germination paper in pouches (Hund et al. 2009), using X-ray absorption (Pierret et al. 2003) or light transmission (Garrigues et al. 2006). Some of these systems with an image acquisition time in the order of minutes allow for enough throughput. For example, growth pouches in combination with digital image analysis allowed to map quantitative trait loci for root elongation (Ruta et al. 2010, Trachsel et al. 2009).

Sampling roots of temperate maize inbred lines in a cold-tolerance study (Hund et al. 2008), I observed differences in GSAs. Some roots did not grow vertically into the growth column but circled along its surface, sometimes even growing upwards. My objectives were, accordingly, to evaluate the GSAs of maize roots and their response to temperature. Additionally, I aimed to find out if GSAs could be determined equally well in growth pouches, which would allow for rapid screening.

Materials and Methods

Experiment 1 in narrow columns

A set of nine inbred lines, D171, CM105, ETHDeL3, ETHFIH1, ETHFIL8, Lo 964, Lo 1016, D167 and S335, was evaluated in narrow growth columns (5.6 cm in diameter and 50 cm high) at low 15/13°C (day/night) and high 24/20°C (day/night) air and soil temperature. For details concerning the lines see Hund et al. (2007). The lines were grown in growth chambers (PGW36, Conviron, Winnipeg, Canada) at a 12 h photo- and thermoperiod, 60/70% (day/night) relative air humidity and 500 μ mol m⁻² s⁻¹ photosynthetic photon flux density (PPFD). The growth substrate was a mixture of quartz sand (particle size 0.08 to 0.2 mm) and 5% w/w vermiculite powder (Vermex Pulver E, Vermica AG, Bözen, Switzerland) with a volumetric content of 15% modified Hoagland solution (Hund et al. 2007). Moist substrate was packed into the growth columns to a bulk density of 1.25 Mg m⁻³; after planting, the substrate was covered with a 1-cm isolation layer of Perlit (PePe[®] Pflanzen Perlit, Otto Hauenstein Samen AG, Rafz-Biberist-Landquart-Orbe, Switzerland). Seeds were surface-sterilised for 12 min with 2.5% NaOCl, pre-germinated at 25°C and planted at a depth of 2 cm with the primary root pointing downwards. Once the coleoptile had reached a length of ~ 1 cm, the growth columns were covered with aluminium foil to prevent warming of the upper zone of the substrate due to the light. The soil temperature was measured at 1 cm (in the isolation layer), 5 cm and 15 cm depth. It was constant throughout the substrate and about 1°C higher in the isolation layer.

Measurements

The genotypes were harvested when more than half the plants had reached the two-leaf (V2) stage. After removing the root system from the growth columns, the roots were rigid enough to allow measurement of their initial angle. For each root, i.e. the primary root, all seminal roots and all crown roots of the first and second shoot internodes, the initial gravitropic set-point angle was measured. The distances (about 3 cm) between the point of root emergence and the point, at which it touched the wall of the tube were used to calculate the initial GSA, further just termed GSA. It was calculated as the arctangent of the ratio between the horizontal distance (Fig. 1, a) and vertical distance (Fig. 1, b).



Fig. 1 Gravitropic set-point angles (α), of the inbred line S335 in narrow growth columns was measured as the deviation from the vertical zero as an arctangent of the ratio of distance a to b.



Experimental design and statistics

The experimental design was a randomised complete block design with two growth chamber replications at both low and high temperatures. Each replication consisted of four blocks, each containing a whole set of genotypes. The experimental unit within block and replication was one growth column with one plant. Thus, the experimental design was:

$$y_{ijklm} = g_i + a_j + t_k + ga_{ij} + gt_{ik} + at_{jk} + gat_{ijk} | r_i + b_{lm} + p_{ilm} + \varepsilon_{ijklm}$$
(1)

where the GSA of each individual root y_{ijklm} depended on the root type and internode it emerged from a_j (j=primary root, seminal roots and crown roots from the first and second internode), inbred line g_i (i=1,...,9), temperature t_k (k=15/13 or 24/20°C day/night) and their interactions. Each of the four root categories was nested within growth chamber run r_l (l=1,...,4; two runs per temperature), complete randomized blocks within growth chamber run b_{lm} (m=1,...,4), individual plant per genotype within complete block p_{ilm} and the random error ε_{ijklm} . Random effects are those to the right of the vertical line in Equation 1. Analysis of variance was carried out with the 'ASReml-R' package in R (Butler 2006).

I aimed to determine whether the GSAs of root types that develop early can be used to predict the GSAs of root types that develop later. Therefore, multiple mixed linear models based on the best linear unbiased predictors derived from equation 1 were fitted for each pair of root types:

$$y_{ik} = a + t_k + at_k \mid g_i + \varepsilon_{ik} \tag{2}$$

Fig. 2 Wide growth columns to measure the vertical and horizontal distribution of roots of contrasting inbred lines S335 and CM105. Plants were placed in a 2-cm cylinder to force the roots to grow vertically. Roots grew in their original direction when entering the wide column with a diameter of 20 cm. The positions at the mesh where the roots grew into the next sections were marked. The distance between the central imaginary shoot axis and the marked position presented the horizontal spread.

where y_{ik} is the GSA of a later developing root type (e.g. seminal roots or crown roots from the first and second internode) depending on the GSAs of an earlier developing root type *a* (e.g. primary root, seminal roots or crown roots from the first internode), the temperature t_k (k=15/13 or 24/20°C day/night), the interaction at_k and the random genotype g_i (i=1,...,9). The intercept was set to 45°, by subtracting the trait values of the earlier root type by 45.

Experiment 2 in wide columns

I aimed to determine whether a large initial GSA would lead to a significant change in the distribution of roots in soil. For this purpose two genotypes with contrasting GSAs (CM105 and S335) were grown in wider columns with, 20 cm in diameter and 20 cm high. Four plants per genotype (one per column) were grown to the V2 stage at 15/13°C. The same substrate and environmental conditions were used as in experiment 1. The columns were divided vertically into 5-cm sections separated by a 2-mm mesh (Fig. 2) to allow recording of the horizontal spread of the axile roots. At harvest, the distances between the radial axis (shoot axis) and the point at which the axile roots

passed through the mesh into the next column section were measured. After recording this horizontal spread, the roots in each column section were harvested and cleaned under running tap water. The root length within each column section was determined by digital image analysis: roots were spread in a tray in a thin layer of water and scanned with a scanner equipped with a top light (Epson, Expression 1640 XL, Epson America, Inc., USA) to acquire 8 bit images at a resolution of 23.7 dots per mm (600 dpi). The images were subsequently analyzed with WinRHIZO 2003b (Regent Instruments, Montreal, QC, Canada) to measure the length of the roots. The debris removal filter was set to remove objects with an area smaller than 0.005 cm² and a length/width ratio lower than 5.

Experiment 3 in growth pouches

Excavating roots from soil is laborious and does not allow for sufficient throughput for the screening of larger populations. Therefore, I wanted to know if the results obtained in sand could be reproduced in growth pouches, described by Hund et al. (2009). The two genotypes CM105 and S335 of Experiment 2, were evaluated in these pouches on blotting paper (21×29.5) cm; Anchor Paper, St. Paul, MN, USA). A total of nine plants per genotype grew under 24/20°C day/night until 10 days after germination. Their root systems were scanned using a Hewlett Packard Scanjet 4670 "See Thru Vertical Scanner" (Hewlett-Packard, Palo Alto, CA, USA). The GSAs of the seminal, primary and crown roots were measured using the "Ruler" tool of Photoshop (Photoshop CS3 extended, Adobe Systems Inc., San Jose, CA, USA). The initial GSA was measured by drawing a line (trace) from the origin of the root to the point where the roots changed the direction significantly. This endpoint was chosen in a way that the distance between the trace and the root was never larger than 1 cm (c.f. Fig. 5, arrows). The mixed linear model to

determine the difference in GSAs among root types and between genotypes was calculated with ASReml-R was:

$$y_{ijkl} = g_i + a_j + ga_{ij} | r_k + b_{ik} + p_{ikl} + \varepsilon_{ijkl}$$
(3)

where the trait value of each individual root y_{ijkl} depends on the root type a_j (j=1,...,3), the inbred line g_i (i=CM105 or S335) and their interactions. The root types were nested within growth chamber run r_k (k=1,...,3), randomly distributed genotypes within growth chamber run b_{ik} and individual plants per genotype within growth chamber run p_{ikl} (l=1,...,3).

Results

Inbred lines classified according to vertical and horizontal rooting

Genotypes were classified according to shallow and deep rooting, independent of temperature. An example of the orientation of the crown roots of inbred line S335 is shown in Fig. 1. Significant effects on the GSAs were detected for most explaining factors and their interactions. Only the inbred line-by-temperature interaction and the three-way interaction among the temperature, inbred line and root type was not significant (Table 1). Averaged over root types, the genotypes with the most vertically oriented GSAs were D171, Lo1016 and CM105, the genotypes with the most horizontally oriented GSAs were ETHDeL3, ETHFIL8 and S335. The GSAs of the seminal roots showed the largest variation among genotypes $(30^{\circ} \text{ to}$ 70°) followed by those of the primary root, $(5^{\circ} \text{ to } 30^{\circ})$, and the crown roots (40° to 60°).

Low temperature resulted in a more vertical orientation of crown roots by 10°

In general the primary roots grew more vertically with

Table 1 Anova results of the effect of the inbred line (Line), temperature regime (Temp) and root type (Root; primary, seminal, crown 1 and crown 2) on the gravitropic set-point angle in narrow growth columns

	Df.	Sum of Sq.	Wald statistic	P ^a (Chisq)
Line	8	17733	95	***
Temp	1	13067	70	***
Root	3	63368	340	***
Line: Temp	8	2532	14	
Line: Root	24	12583	67	***
Temp: Root	3	4109	22	***
Line:Temp: Root	20	5668	30	

^a Significance level at p < 0.1 (.) and 0.001 (***).

Inbred line	Gravitropic set-point angle (°)				
	Pr ^a	Se ^b	Cr 1 ^c	Cr 2 ^d	
D171	13.7	28.3	45.7	38.5	
Lo 1016	5.4	42.8	42.4	40.0	
CM105	23.6	37.0	42.2	NA ^e	
ETHFlH1	16.3	47.3	39.8	NA	
Lo 964	15.5	42.9	51.3	43.1	
D167	24.6	52.9	51.1	NA	
ETHDeL3	24.8	62.8	39.9	50.6	
ETHFIL8	32.2	47.2	53.2	49.1	
S335	25.4	71.6	58.8	NA	
avsed ^f	6.6				
Temp. (°C) ^g					
15/13°	12.7	48.9	42.3	43.2	
24/20	27.6	47.3	52.0	53.4	
avsed	3.8				
Mean	20.2	48.1	47.2	44.3	

 Table 2 Gravitropic set-point anlges in narrow growth columns as dependent on temperature (Temp.), inbred line and root type

^a primary root

^b seminal root

^c crown root from the first shoot internode

^d crown root from the second shoot internode ^e not estimated since low number of plants

advanced to Cr 2

^f average standard error of the difference

^g temperature day/night

mean GSAs of about 20°, while the successive developing seminal and crown roots grew more horizontally with mean GSAs between 44° and 48° (Table 2). Surprisingly, low temperature resulted in a more vertical orientation of primary axile roots (about 15°) and crown roots (about 10°)(Table 2). The responses of GSAs of the embryonic primary and seminal roots to temperature must be considered with

caution since these roots may have established at slightly higher temperature. The reason for this is that the final soil temperature was adjusted by covering the column with aluminium foil after the coleoptile emerged. For example, in the low temperature treatment, the temperature at 1 (in the Perlite), 5 and 15 cm column depth was about 19, 17 and 16°C, respectively, before covering the tubes and 16, 15 and 15°C thereafter.

The GSAs of seminal and early crown roots correlated with the GSAs of crown roots from the second internode

Can GSAs of early internodes be used to predict GSAs of later internodes and does temperature play a role? Using equation 2, the angles of the primary roots, which tended to grow vertically, could not be used to predict GSAs of later root types (data not shown). However, for other root types, predictions were possible (Table 3, a). The GSAs of the crown roots from the second internode could be predicted by those of the seminal roots (slope of 0.97) and the crown roots from the first internode (slope of 1.41). Temperature did not influence this covariance since no interaction was found between temperature and GSA (data not shown). Thus, the interaction terms (at_k) were dropped from the models. Temperature affected the intercept of the growth angles for those models where the seminal roots were the explaining variable (Table 3, t). Optimal temperature resulted in an estimated increase in the root angles of the crown roots of about 10°.

Variability in GSAs resulted in different soil exploration of two contrasting genotypes

The GSAs for the contrasting inbred lines, CM105 with a more vertical root orientation and S335 with a more horizontal root orientation, are shown in Fig. 3. The GSAs of the seminal and crown roots of S335 were much larger as those of CM105. The effect of the differences in GSAs on the distribution of roots in soil

 Table 3 Linear mixed model (equation 2) to predict GSAs of later developing root types as dependent on the temperature and GSAs of earlier developing root types

		Cr 1 vs. Se		Cr 2 vs.	Cr 2 vs. Se		Cr 2 vs. Cr 1	
	-	estimate	p ^a	estimate	р	estimate	р	
Intercept	(at 45°) ^b	41.19	***	38.44	***	45.98	***	
Root angle	<i>(a)</i>	0.29		0.97	**	1.41	**	
Optimal Temp.	<i>(t)</i>	9.45	*	11.78	*	-0.94	ns	

For abbreviations of root types see Table 2.

^a significance level at p < 0.1 (.), 0.05 (*), 0.01 (**) and 0.001(***).

^b The intercept was set at 45°.



Fig. 3 Gravitropic set-point angles (GSA) of contrasting inbred lines S335 and CM105 in narrow growth columns. Shaded areas indicate average GSAs including 95% confidence intervals of the primary (Pr), seminal (Se) and crown (Cr) roots of the first internode.

Table 4 Best linear unbiased estimates (equation 3) of
gravitropic set-point anlges (GSA) measured
in growth pouches as dependent on inbred line
and root type

	Factors				
	Line	Root	Line:Root		
ANOVA	***	***	ns		
	GSA of root type (°)				
Inbred Line	Pr ^a	Se ^b	Cr °		
CM105	14.3	29.0	43.4		
S335	19.4	53.5	68.2		
avsed ^d	8.2				

^a primary root

^b seminal root

^c crown root

^d average standard error of the difference

was evaluated using the wide growth columns (Experiment 2). The axile roots of S335 spread in a much more horizontal direction and, consequently, did not reach the lowest 15 to 20-cm section of the column (Fig. 4a). By contrast, the axile roots of CM105 grew much more vertically and reached the 15 to 20-cm section of the columns. This difference in vertical root distribution is corroborated by the relative distribution of the roots (Fig. 4b). The inbred line S335 had twice as many roots in the 0 to 5 cm section compared to CM105. This changed in the successive column sections where CM105 had at least twice as many roots.



Fig. 4 Root distribution of two contrasting genotypes in wide growth columns: S335 (triangles) with more horizontal rooting; CM105 (circles) with more vertical rooting. Distance from the shoot axis, at which axile roots passed into the next column sections (a; horizontal spread, *cf.* Fig. 2) and distribution of root length with column depth as percentage of total root length (b). Shaded areas indicate the average maximum and minimum horizontal spread of the axile roots. Error bars show one standard deviation.

Growth pouches are suitable for assessing GSAs

The differences in the GSA of CM105 and S335 were verified in growth pouches. The lines differed with respect to the GSA of their seminal and crown roots (Table 4). Compared to GSAs measured in the narrow columns, the differences between the genotypes were even more pronounced. At 24/20°C (day/night) in narrow columns, the differences between the genotypes were 19° for the seminal roots and 6° for the crown roots of the first whorl. In growth pouches these differences were around 30° for both root types. Two root systems of CM105 and S335, each, are shown in Fig. 5. The magnified part of the image illustrates how GSAs were measured.

The primary roots of the two S335 plants in Fig. 5 show a plagiogravitropic response changing from an almost vertical orientation to a more horizontal orientation. Since only the initial GSAs were measured, this later change was not recorded. Taking it into account may have resulted in pronounced differences also between the GSAs of the primary roots of the two genotypes.

Discussion

The GSAs of the inbred lines differed strongly, leading to pronounced differences in the distribution of roots in the wider columns. If classified according to the simplified root system shapes illustrated by Hammer et al. (2009), the inbred line CM105 had a more elliptical root system, while S335 had a standard circular root system. According to Hammer et al. (2009), the more elliptical, compact root system may reflect the root system of modern varieties adapted to



Fig. 5 Examples of the root system of S335 and CM105 (two plants each) grown in growth pouches. The magnified area illustrates the measurement of the GSAs of the primary root (red), seminal roots (green) and crown roots (blue). The roots were traced in a way, that the distance between the root and the trace (see arrows) was never larger than 1 cm.

high plant density. However, there are several reasons why this may be an over-simplification. In particular in cool or phosphorus-deficient soil, more horizontal orientation of some roots may be of greater benefit. The avoidance of low subsoil temperature by increased GSAs is well documented for maize (Mosher and Miller, 1972; Onderdon and Ketcheso, 1973; Tardieu and Pellerin 1991). However, this type of response was not verified in the present set of inbred lines. On the contrary, the crown roots grew in a slightly more vertical direction. Similar effects were observed by Onderdon and Ketcheso (1973) where the GSAs were largest at 17°C and lower when plants were grown at lower temperature. The initial GSAs were measured between 0 and about 3 cm from the shoot base in a substrate with uniform temperature (due to the isolation layer of Perlit and the coverage with aluminium foil). Therefore, it may be argued, that the roots would have responded if there had been a temperature gradient in the substrate. However, according to studies under controlled conditions, the temperature per se rather than thermotropism is

responsible for altering the GSA of roots in soil (Mosher and Miller, 1972; Sheppard and Miller, 1977).

Compared with the GSAs reported in older literature, the GSAs of the modern inbred lines used here, showed a striking difference: The primary root of modern inbred lines oriented almost vertical, while the primary (and seminal roots) of older genotypes grew almost horizontally, i.e. were plagiogravitropic or even diagravitropic (Feldman and Briggs 1987, Kisselbach 1999, Mosher and Miller 1972, Onderdon and Ketcheso 1973, Scott and Wilkins 1969, Suzuki and Fujii 1978, Weaver 1925). It is tempting to speculate that the more vertical orientation of modern inbred lines reflects breeders' selection during the last 30 years as suggested by Hammer et al. (2009). This raises the question as to whether roots of modern inbred lines have lost their ability to benefit from warming of the superficial soil layer in spring by more horizontal rooting.

One problem of testing the adaptive value of a more horizontally oriented seedling root system is the effect of soil temperature on the shoot apex of maize seedlings (Hund et al. 2008). Low apex temperatures cause a strong decrease in the photosynthetic efficiency of some lines, e.g. CM105. In order to test the beneficial effect of shallow rooting independent of this temperature effect on photosynthesis, one solution is to develop near-isogenic lines for this trait.

Another problem is that the initial GSAs may not be sufficient to characterize genotypes according to the distribution of their roots in soil. For example, plagiogravitropic behaviour of the primary root becomes obvious only after the roots grow to a considerable length in the soil (Nakamoto 1994). This behaviour could also be observed for the primary root of S335 in the growth pouches (Fig. 5). The plagiogravitropic response of these roots in the pouches may be quantified in using adequate software allowing to measure root trajectories. Such software is available for simpler root system, such as the one of *Arabidopsis thaliana* L. (Armengaud et al. 2009). Thus, growth pouches are feasible to measure both initial GSAs and successive gravitropic responses of the roots.

My major concern was that scattered light in the pouch may lead to a decrease in the GSAs (Feldman and Briggs, 1987). However, the GSAs deviated to an even greater extent from the vertical as in sand, and the effect of scattered light therefore seemed to be marginal. The feasibility of growth pouches to measure GSAs is also supported by Bonser et al. (1996) using them successfully to assess the response of GSAs of basal roots of common bean to phosphorus. Growth pouches or gel chambers are among the phenotyping systems with the highest throughput and are probably first choice for selection experiments or

QTL mapping approaches. More elegant methods to assess the root system in undisturbed soil, such as magnetic resonance tomography (MRI) (Van As 2007) or x-ray computer tomography (Gregory et al. 2003) have the disadvantage that they lack throughput. For example, the throughput of partly automated MRI screening systems is expected to be 10-15 individuals per day (Jahnke et al. 2009). For larger plants, excavating the roots in the field may be efficient to assess this trait. Working with larger plants would allow measuring the dynamics of the change in GSAs across successively developing internodes. The initial GSAs of maize decrease with increasing internode (Araki et al. 2000) but there may be differences among genotypes as indicated by the results presented here: Roots emerging from the first shoot internode may already exhibited a strong vertical orientation, as shown for CM105.

Conclusion

There was great variability in the GSAs among the tested inbred lines ranging from vertical to almost horizontal. The GSAs did not increase at low temperature suggesting that most studied genotypes would not be able to benefit from soil warming in spring. Given the importance of the GSAs for the acquisition of resources from soil, we need more insight into their genetic control. Growth pouches proved feasible for measuring GSAs at similar precision as in a more natural sand substrate. The comparably high throughput of these pouches enables the selection for GSAs or mapping of quantitative trait loci controlling them. This in turn enables us to elucidate the significance of GSAs for the adaption of plants to cool soil in spring.

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Dr. Andreas Hund's research interest is crop ecophysiology. His main focus is on the inheritance of root system architecture of maize and its adaptation to abiotic stresses.