



# Hydrogel-based transparent soils for root phenotyping in vivo

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**Root phenotypes are increasingly explored as predictors of crop performance but are still challenging to characterize. Media that mimic field conditions (e.g., soil, sand) are opaque to most forms of radiation, while transparent media do not provide field-relevant growing conditions and phenotypes. We describe here a “transparent soil” formed by the spherification of hydrogels of biopolymers. It is specifically designed to support root growth in the presence of air, water, and nutrients, and allows the time-resolved phenotyping of roots in vivo by both photography and microscopy. The roots developed by soybean plants in this medium are significantly more similar to those developed in real soil than those developed in hydroponic conditions and do not show signs of hypoxia. Lastly, we show that the granular nature and tunable properties of these hydrogel beads can be leveraged to investigate the response of roots to gradients in water availability and soil stiffness.**

soil | transparent | hydrogels | plants | microbiome

Growing plants for research is constrained by an apparently necessary compromise. On one hand, media that are representative of field soil (e.g., soil, sand) are opaque to most forms of radiation (1) and offer limited control over heterogeneities that affect the development of roots (e.g., gradients in water availability, nutrient concentrations, mechanical properties, porosity). On the other hand, transparent media (e.g., hydroponics, aeroponics, gels) do not provide field-relevant phenotypes and growing conditions (2).

Media that have air-filled, connected pores display several physiologically relevant characteristics of soil, such as aeration and physical interfaces (3). Unfortunately, these porous media are usually opaque to most electromagnetic radiation because each interface changes the direction of propagation of photons, due to refraction and reflection. The magnitude of these deflections increases with the difference between the refractive indices (a physical property of matter dependent on electronic density and susceptibility) of the medium and the material contained in the pores (4). Therefore, a porous medium can become transparent to light if it is fully saturated with a fluid whose refractive index matches that of the porous medium (5).

Index matching of granular materials, including hydrogels, was used successfully to study hydrology, soil physics, and fluid dynamics in porous media (6, 7). Nonetheless, the use of this approach to study root development is subject to numerous complex constraints that have made this task notoriously challenging. The medium must be (i) produced simply and inexpensively in large quantities (hectoliters), (ii) nontoxic to plants, (iii) transparent enough in common nutrient solutions to allow for the phenotyping of a whole root system in vivo, and (iv) strong enough to not collapse under its own weight. Furthermore, it should provide water and nutrition to the growing plant and have a fully connected porosity to prevent the formation of air pockets. A recent pioneering work by Downie et al. (5, 8) reported Nafion, a sulfonated tetrafluoroethylene-based fluoropolymer–copolymer, as a promising

material for this purpose, given the similarity of its refractive index to that of water. Unfortunately, the material is currently very expensive (~\$1,000/kg), it must be chemically processed before use with plants, it does not absorb water or nutrients (some parts of the root system must be saturated with nutrient solution), and its index matching solution has significant concentrations of sorbitol (0 to 13%, wt/vol), which can cause osmotic stress in plants (9).

We here describe a porous medium that allows for the imaging of unconstrained root systems in vivo by both photography and microscopy and the time-resolved phenotyping of roots. The medium consists of interconnected pores that are surrounded by nutrient solution, held into spherical beads of hydrogel. These beads have controllable size and hardness and are produced simply, rapidly, and inexpensively by dropping a solution of gellan gum and alginate into a stirred solution of MgCl<sub>2</sub>. Temporary saturation with nutrient solution (a treatment comparable to rainfall or watering) makes this medium sufficiently transparent to allow imaging of a 20 × 20 × 20 cm volume. This medium outperformed hydroponics in producing field-relevant root phenotypes in *Glycine max* (six of seven key phenotypes were not significantly different from field soil's, instead of two). Similarly, a key gene involved in response to root hypoxia [nonsymbiotic Hemoglobin (nsHB); Glyma.11G121800 (10)] was significantly overexpressed in hydroponics but was not

## Significance

**Imaging of plant roots is severely limited by the opacity of soil media. Hydroponic (or gel) conditions provide transparency but nonphysiological root phenotypes. Here, we develop a “transparent soil” with high transparency, good mechanical stability, tunable pore sizes, low cost, and easy scalability. This porous media can support root growth in the presence of air, water, and nutrients, and allows for the imaging of unconstrained root systems in vivo by both photography and microscopy. Our study provides evidence that the roots of soybean developed in this medium are significantly more similar to those developed in real soil than those developed in hydroponic conditions and do not show signs of hypoxia.**

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Conflict of interest statement: L.M. and L.C. are inventors on a patent application (US 16/107,512) submitted by Iowa State University Research Foundation, Inc. that covers methods of making hydrogel-based transparent soil.

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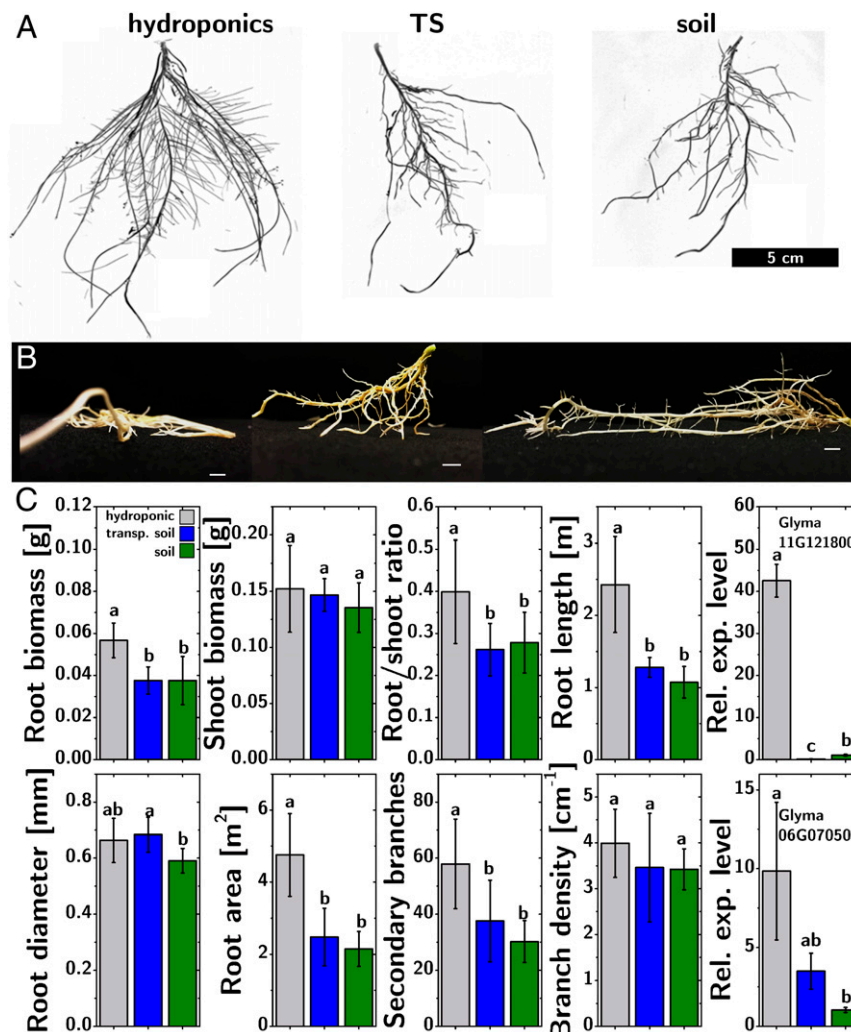
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**Fig. 3.** Root phenotyping in TS. (A and B) Comparison of (A) *G. max* roots and (B) their mechanical stiffness, after growth in hydroponics (Left), TS (Center), and sterilized field soil (Right). (Scale bar in B, 1 cm.) (C) Comparison of biomass, root morphology traits, and gene expression (Glyma.11G121800 and Glyma.06G070500) in *G. max* plants grown in hydroponics, TS, and sterilized field soil. Error bars for biomass and root morphology traits indicate SD ( $n = 9$ ); error bars for gene expression represent SD ( $n = 4$ ). Different letters above the histograms (a, b, c) indicate significant differences between treatments ( $P < 0.05$ , one-way ANOVA followed by Tukey's test).

of lateral roots and root tips (Glyma.06G070500, Glyma.08G133800, Glyma.02G043400) in plants (root tissues of *G. max*, IA2102, 12 d old,  $n = 4$ ) grown in hydroponics, TS, and field soil, using soil extract as a nutrient medium, as described for the phenotyping study. Relative expression was calculated by the  $2^{-\Delta\Delta Ct}$  method by considering soil treatment as the calibrator (18). Genes associated with nutrient or water deficiency (Glyma.12g221500, Glyma.08g053500, Glyma.04G203300) were not differentially expressed across treatments, suggesting that the TS does not deprive plants of nutrients or water, compared with either field soil or hydroponics. Two of the genes associated with root development (Glyma.08G133800, Glyma.02G043400) were also not differentially expressed in the three treatments.

Two genes were found to be differentially expressed (cf. Fig. 3C). The first (Glyma.11G121800, nsHB) is strongly associated with response to hypoxia in soybean through the synthesis of leghemoglobin, and was strongly overexpressed in the hydroponic treatment (42-fold compared with field soil,  $P < 0.001$ ) and underexpressed in the TS treatment (0.14-fold compared with field soil,  $P < 0.001$ ), strongly indicating that TS, in contrast to hydroponics, does not cause hypoxic stress in roots. The lower hypoxic stress in TS and field soil would be consistent with the

increased curvature observed in the root phenotypes (19). The other differentially expressed gene (Glyma.06G070500, GTP-binding nuclear protein Ran) is associated with signal transduction and stress response and was overexpressed in hydroponics (10-fold compared with field soil,  $P = 0.007$ ). This gene is found to be highly expressed in lateral roots, and the expression qualitatively mimics the dependency of SRN on the treatments, suggesting that the observed differential expression is not due to a stress response but due to a different root structure.

This TS is also compatible with in vivo microscopy of plant roots. *Arabidopsis thaliana* plants (with a GFP-tagged plasma membrane) were grown in Petri dishes filled with TS and fitted with a glass coverslip window (cf. Fig. 4A). Confocal microscopy (cf. Fig. 4B) shows root segments at different depths (0 and 2.4 mm, behind a hydrogel bead) into the TS. Similarly, fluorescence microscopy allows optical and fluorescence imaging of roots behind hydrogel beads, several millimeters (2.9 mm) inside the TS (cf. Fig. 4C).

Lastly, this TS can be easily modified to visualize chemical changes caused by the roots or study the effect of soil heterogeneities on root development in vivo. Fig. 4D shows *G. max* roots growing in TS, where a yellow overlay indicates local



much lower cost in the same amount of time by implementing very simple automation (e.g., cameras on rails coupled with peristaltic pumps for watering/draining).

Root phenotypes are nonlinear functions of large numbers of correlated input variables. Therefore, their study is a statistics-dependent problem where data quantity and quality are key. Using a GxE screen as an example, our TS medium could enable the initial screening of relevant traits over very large genotype/environment sets, and allow a highly statistically informed decision on what genotypes to explore in an X-ray CT/MRI, data/time/expertise/cost-intensive manner.

X-ray CT is often regarded as the state-of-the-art technique for root phenotyping, but it is not devoid of limitations. Besides throughput and cost, X-ray tomography (i) generally requires moving the plant to the instrument [which can affect plants (27)], (ii) requires small pot sizes [ $\sim 8$  cm in diameter, according to the review by Metzner et al. (1)], (iii) has difficulty identifying all roots [it reliably identifies 60 to 70% of the root system, according to Metzner et al. (1)], and (iv) is significantly affected by the moisture, type, and heterogeneity of soil (28).

These limitations affect its natural-substrate relevance: Limiting the pot size can expose plants to thigmotropic responses with the walls of the container (1, 28, 29), using soils suited for CT limits the ability to test the phenotypes in other soils, and missing 20 to 30% of the root system could add unknown systematic errors to phenotype characterization (e.g., in the characterization of average root length for genotypes with different average root diameter). In summary, when all methodologies have limitations, it is usually best to develop complementary

techniques with different limitations. In this sense, this TS fills a much-needed niche in the spectrum of techniques available for root phenotyping.

Of course, as outlined above, this TS medium has limitations. (i) The transparency and the mechanical properties limit the size of the root volume to about  $20 \times 20 \times 20$  cm. (ii) The size of the beads is currently limited to between 0.5 mm and 5 mm. (iii) Methods to produce hectoliter amounts of TS are not detailed here (a scale-up approach to produce 50 L/d from a single setup is outlined in *SI Appendix*). (iv) The surface chemistry of the TS beads is significantly different from that of soil.

There are reasons to believe that many of the above limitations can be overcome: Hydrogels have been a material of choice for tissue engineering and drug delivery, due to the endless possibilities they offer for functionalization and control of mass transport (30–33). We expect that this application of hydrogels will provide, through synthetic polymers and their functionalization, broad possibilities for the quantitative study of GxE interactions in root development and the modeling of the rhizosphere, as well as in other fields of science such as animal science (34–36), robotics (37), soft matter physics (38), and biomechanics (39).

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