Splanting: 3D plant capture with gaussian splatting

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Fig. 1. 3D gaussian splatting based capture of a canola plant in a laboratory environment with a visually cluttered background: (left-to-right) reconstruction including background splats (scrambled for privacy), isolated canola *splant* after removal of background splats by spatial clustering, isolated flower clusters after radiance-based segmentation, and zoomed-in flowers.

3D capture and characterization of plant shoot architecture is a grand challenge in plant phenotyping research, made difficult by plants' intricate 3D shape, composed of thin and flat sub-structures (stems, leaves, flowers, pods, etc.). In this paper, we show that 3D gaussian splatting is well-suited for capturing 3D plant representations, which we call *splants*. We report a simple and fast capture procedure and 3DGS processing software that is tailored to foreground object capture. Splant generation worked well across plant species and growth stages. Our preliminary results point to a promising future for splant phenotyping, which we expect will lead to a dramatic increase in the use of multi-view imaging and 3D analysis in plant pathology and plant breeding research.

$\label{eq:ccs} \texttt{CCS Concepts:} \bullet \textbf{Computing methodologies} \to \textbf{Point-based models}; \textbf{Reconstruction}; \bullet \textbf{Applied computing} \to \textbf{Agriculture}.$

Additional Key Words and Phrases: 3D gaussian splatting, radiance field rendering, image processing, 3D reconstruction, plant shoot architecture, plant phenotyping

1 Introduction

3D reconstruction and rendering of a plant's above-ground shoot architecture is a long-standing grand challenge in plant phenotyping. The shape and spatial arrangement of a plant's shoots determine its capacity to capture solar energy, its yield potential, its susceptibility to certain plant pathogens, and how easily it can be harvested. However, characterizing a plant's shoot architecture is notoriously difficult due to its intricate 3D shape, composed of lower-dimensional manifolds: 1D branches, stems, silique and 2D leaves, flower petals, pods. Reconstruction of complex and intertwined plant structures from images has been hampered by severe self-occlusion and self-similarity which makes traditional photogrammetry methods brittle, both in terms of feature matching and localization.

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3D gaussian splatting (3DGS) [Kerbl et al. 2023] has generated impressive visual results for natural scenes, but has yet to be evaluated for complex plant architectures with substantial self-occlusion. In this paper, we provide a preliminary evaluation and analysis of 3DGS-based plant reconstructions, which we call *splants*. We examine splants across a range of plant species and demonstrate that their optimized splats are representative of physical plant structures and that splat location, orientation and radiance can be used to localize and identify plant organs.

The contributions of our paper include: 1) an exemplar splant dataset including diverse samples across plant species and growth stages; 2) a simple splant capture procedure; 3) splant software for 3DGS processing (optimized for foreground object capture) and splat filtering, clustering, and contraction processing, towards skeletonization of a plant's architecture; 5) a demonstration of using splat features (location, orientation, radiance) for plant organ segmentation; and 6) a discussion of the implications of splant capture for plant phenotyping.

2 Related Work

3D Plant Capture. Prior to recent advances in radiance field rendering, a wide range of approaches have been proposed for 3D capture of plant shoot architecture. Plants have been subjected to multi-view photogrammetry, X-ray imaging [Dutagaci et al. 2020], and laser scanning [Schunck et al. 2021]. Of these traditional capture methods, 3D laser scanning has worked best [Young et al. 2024], but still requires fairly slow and careful manual or robot-controlled movement of the scanner to capture the thin, intricate, and self-occluded structures that comprise a plant's overall architecture.

Neural Radiance Fields for Plants. Radiance field rendering research is rapidly accelerating in many directions and quickly gaining interest in specialized domains, such as plant phenotyping. Neural Radiance Field (NeRF) capture of plants has shown promising initial results for specific species [Saeed et al. 2023] and even in challenging agricultural field conditions [Arshad et al. 2024].

3D Gaussian Splatting for Plants. 3DGS has been successfully used to measure stem length and count bolls in cotton plants [Jiang et al. 2024]. However, to the best of our knowledge, 3D gaussian splatting has yet to be evaluated for complex plant structures that include substantial self-occlusion and for multiple plant species. Concurrent to our work, 3DGS and NeRF reconstructions of wheat plants have been compared [Stuart et al. 2024]. One advantage of 3DGS compared to implicit representations, like NeRF, is that it directly generates an explicit representation of the plant as a 3D splat cloud that is suitable for downstream 3D plant simulation [Cieslak et al. 2022; Huang and Stavness 2022; Li et al. 2023].

3 Methods

We have designed a quick and effective workflow for splant generation that extends the canonical 3D gaussian splatting optimization procedure for capturing individual plants from multi-view images.

3.1 Splant Capture

Imaging. Our splant capture protocol uses a single 360° side-view video clip around the plant (Figure 2), from which we extract ~100 frames for 3DGS processing. Our pilot testing with different cluttered backgrounds, lighting conditions, and camera paths all worked well for static scenes. We found the traditional setup of a fixed camera and a rotating plant on a turn-table, as well as entirely blank backgrounds, were less effective because they degraded the initial structure-from-motion (SfM) step of the 3DGS process. As future work, fiducial markers may improve SfM initialization.

3DGS Processing. We use the original implementation of 3DGS for processing splants [Kerbl et al. 2023]. Although our interest is only in the foreground plant, background features assist in the SfM estimation of camera parameters and initial splat locations. However, to reduce processing time and focus 3DGS processing on the plant

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Fig. 2. Our splant capture protocol uses dense side-view images or a video clip around the plant.

species	growth stage	# images in	# splats out	shown
canola	rosette	77	48,264	Fig. 5
canola	flowering	97	214,598	Fig. 1
wheat	mature	90	104,433	Fig. 3
bean	vegetative	93	144,486	supp. video
chickpea	vegetative	82	150,080	Fig. 3
lentil	vegetative	83	431,979	Fig. 4
pea	vegetative	103	127,031	Fig. 7
pea	pods	79	187,807	Fig. 7
pea	mature	93	143,775	Fig. 7

Table 1. Exemplars from our *splant* dataset.

subject, we pre-process input images by cropping to the plant region-of-interest (ROI) using a COCO-pretrained YOLOv8n detector [Jocher et al. 2023]. For the pea splant, ROI image cropping reduced 3DGS processing time by approx. 25% and reduced the number of background splats by half (Table 2) although this was scene dependent. Tight ROI cropping, however did cause SfM failures in some cases, in which case we revert to the original images.

Background splat removal. The primary post-processing step is to isolate the foreground plant splats from the background splats and to remove spurious splats that are not associated with plant subjects. Using the Cloud Compare v2.11.3 CLI (CC), we first perform Statistical Outlier Removal (SOR) to remove the "floating" splats using a standard deviation multiplier threshold of 1 and 15 points for mean distance estimate. Next, we generate candidate splat clusters via connected components in CC, with one component representing the plant and the other components representing the background. We manually select the component that best represents the plant in CC, a step which we plan to automate in future work. After exporting the selected plant component, we merge back per-splat parameters, which is not yet supported in CC. Of note, the support surface for the plant was often included in the plant component, which generated rather pleasing renderings of wire shelves and carts (Figures 1a, 4), but had to be manually removed.



Fig. 3. Example renderings of wheat and pea splants, showing side, top-down, and zoomed-in views.

3.2 Splant Dataset

We have captured splants across a range of agronomically-important plant species, growth-stages, and backgrounds. While the present dataset is modest in size, we expect it to grow rapidly due to the ease and speed of capture, supported by our splant processing software. Exemplar samples from the dataset are summarized in Table 1.

3.3 Splant Analyses

We have developed preliminary analyses of splant renderings to demonstrate their potential utility for characterizing plant traits. In order to demonstrate that the optimized splats that comprise a splant are representative of plant structures, we explore the use of splat locations to skeletonize splants and compute splant volume, splat radiance to isolate canola flowers, and splant orientation to differentiate splat clusters belonging to different types of organs.

4 Results

Qualitative results. Figure 1 depicts a representative rendering of a canola splant at the flowering stage, first with the whole room background reconstructed, then with plant-related splats isolated, and finally with canola flower splats isolated based on their radiance. Additional exemplars include wheat and pea splants with visible spikes and pods (Figures 3), lentil and chickpea splants with small dense leaflets (Figure 4), and a pea splant over time (Figure 7).

Using splat location to skeletonize plant architecture. Using splant location alone is akin to point-cloud processing [Meyer et al. 2023]. Splant splat clouds are substantially sparser than traditional plant point clouds, but utilizing similar contraction-based processing of splant locations clearly reveals the stems and leaves in a rosette-stage canola splant with only a few missing regions due to self-occlusion (Figure 5). We also found that the concave hull of splat locations provided a tight fit for the chickpea splant and may be a good proxy for a plant's above ground biomass volume.

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Fig. 4. Example renderings of lentil and chickpea splants, showing side, top, and zoomed-in views. The small leaflets in these species are visible, but blurry when zoomed-in.



Fig. 5. A rosette-stage canola splant: (a) oblique perspective, (b) top-down perspective, (c) splat locations after contraction showing that splats are arranged near plant structures, (d) stem splats isolated using RANSAC.



Fig. 6. Orientation of splat principle axis relative to the overall object axis for three instances of artificial support sticks, canola stems, and wheat spikes (isolated from full plant captures). Splats are aligned to thin plant organs, and uniformly oriented for wheat spikes to capture the thin awns.

inp imag	ut ges	# splats total	<pre># splats background</pre>	# splats plant	3DGS processing time (minutes)
origi	nal	515,971	388,940	127,031	29
crop	ped	372,541	114,658	257,883	22

Table 2. Comparison pea splant generation with and without cropping input images to the plant ROI.

Using splat orientations to differentiate plant organ types. When optimized to render thin and flat plant structures (stems and leaves), we expect that splats will be oriented to align with thin structures. To characterize the optimized splant splats, we measure the distribution of their orientations for three canonical organ/objects within our splant dataset: wheat spikes, canola stems, and artificial support sticks. We first find the principal axis of the organ using principal component analysis on the isolated splat cloud. Each individual splat geometry is composed of its scaling factors and the rotation matrix created from the normalized given quaternions. Using this information, we can efficiently extract the principal axis of each splat by extracting a column from the rotation matrix that corresponds to the largest scaling factor. We found that splats were well aligned to the stick, mostly aligned to the stem, and more uniformly distributed for the wheat spike (Figure 6).

Using splat radiance to isolate canola flowers. Given their distinctive yellow color, canola flowers are the "lowhanging fruit" of plant phenotyping. However their dense arrangement into cylindrical clusters (racemes), makes it difficult to accurately count individual flowers from a single image. We isolate flowers in the canola splant by extracting the splats' 0th spherical harmonic band (base color) and thresholding for yellowness in the LAB colorspace. Additional filtering was done for low-opacity splats and by SOR to remove small parts of stems that appear yellow. We show isolated canola flowers in place and zoomed-in for nine examples (Figure 1).

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Fig. 7. The same pea plant captured across growth stages: vegetative, pod-development, senescence.

Number of input views required. We found that our side-view 360° capture procedure required a narrow separation between input images to work well. For most splants, less than ~30 images (12° separation) caused the initial SfM step to fail, and greater than ~100 images (~ 3.5° separation) did not improve the reconstructed splant.

5 Discussion and Implications

To conclude, we discuss the implications of *splants* for plant phenotyping research. Our preliminary attempts at generating 3D splants resulted in highly realistic 3D renderings for every species attempted, with few failures. Compared to laser scanning [Schunck et al. 2021], splant capture is faster (~15 sec vs. ~15 min), cheaper (\$100 camera vs. \$10,000+ scanner), generates smaller point clouds (~100k splats vs. ~1M+ points), and is less tedious. In our exemplar splants, we have observed some limitations: splats were missing in highly occluded regions, support structures (cart, tabletop) were not always removed with our simple SOR approach, and very dense plants did not admit a view of internal architecture and rendered somewhat fuzzy (Figure 4). However, by and large the splant exemplars we produced are rather wonderful at a whole plant level, and have sufficient splat density and detail to permit interrogation of whole plant architecture and individual plant organs.

The implications of splant capture for plant phenotyping are numerous. The speed and ease of capture would permit 3D phenotyping for modest size plant populations (100s-1000s of individuals) for plant breeding and pathology experiments. The computational resources required (a desktop GPU) are accessible to most plant science research groups around the world. The resulting photorealistic renders are likely sufficient for qualitative visual phenotyping and screening, which is common and often preferred by breeders. High-throughput and automated splant capture and analysis will require adapting phenotypic analysis approaches from 2D images and 3D point clouds to 3D splat clouds, but the compact representation afforded by splants will make analysis workflows more scalable. 360° splant capture is only feasible for certain plant experiment conditions and growth facilities, but these types of experiments, e.g. plant pathology, are the ones that require detailed phenotypic information potentially afforded by splants. Evaluating outdoor splant capture for crops grown in dense field plots is an important future step for expanding splant use in crop breeding programs. Finally, new 3DGS variants are rapidly improving efficiency and accuracy of reconstructions for general scenes, and optimizing these methods for *splanting* is a promising direction for future work.

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