

METHODOLOGY

Open Access



Developing non-invasive 3D quantificational imaging for intelligent coconut analysis system with X-ray

Yu Zhang¹, Qianfan Liu¹, Jing Chen², Chengxu Sun^{3*}, Shenghuang Lin², Hongxing Cao³, Zhaolin Xiao⁴ and Mengxing Huang^{1,5}

Abstract

Background As one of the largest drupes in the world, the coconut has a special multilayered structure and a seed development process that is not yet fully understood. On the one hand, the special structure of the coconut pericarp prevents the development of external damage to the coconut fruit, and on the other hand, the thickness of the coconut shell makes it difficult to observe the development of bacteria inside it. In addition, coconut takes about 1 year to progress from pollination to maturity. During the long development process, coconut development is vulnerable to natural disasters, cold waves, typhoons, etc. Therefore, nondestructive observation of the internal development process remains a highly important and challenging task. In this study, We proposed an intelligent system for building a three-dimensional (3D) quantitative imaging model of coconut fruit using Computed Tomography (CT) images. Cross-sectional images of coconut fruit were obtained by spiral CT scanning. Then a point cloud model was built by extracting 3D coordinate data and RGB values. The point cloud model was denoised using the cluster denoising method. Finally, a 3D quantitative model of a coconut fruit was established.

Results The innovations of this work are as follows. 1) Using CT scans, we obtained a total of 37,950 non-destructive internal growth change maps of various types of coconuts to establish a coconut data set called "CCID", which provides powerful graphical data support for coconut research. 2) Based on this data set, we built a coconut intelligence system. By inputting a batch of coconut images into a 3D point cloud map, the internal structure information can be ascertained, the entire contour can be drawn and rendered according to need, and the long diameter, short diameter and volume of the required structure can be obtained. We maintained quantitative observation on a batch of local Hainan coconuts for more than 3 months. With 40 coconuts as test cases, the high accuracy of the model generated by the system is proven. The system has a good application value and broad popularization prospects in the cultivation and optimization of coconut fruit.

Conclusion The evaluation results show that the 3D quantitative imaging model has high accuracy in capturing the internal development process of coconut fruits. The system can effectively assist growers in internal developmental observations and in structural data acquisition from coconut, thus providing decision-making support for improving the cultivation conditions of coconuts.

Keywords Intelligent coconut analysis, Non-invasive, Point cloud, Quantitative imaging model

*Correspondence:

Chengxu Sun

hnsuncx@qq.com

Full list of author information is available at the end of the article



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

Introduction

Coconut palms (*Cocos nucifera* L.) generate oil-bearing seeds which are essential for the production of vegetable oil in the marketplace. The coconut fruit is a fibrous drupe, not a nut. A drupe is a simple fleshy fruit containing one seed, derived from one ovary of a flower. A thin skin covers the outermost layer of the ovary wall. The coconut has a thick, fibrous middle layer of the ovary wall. The inner layer of the ovary wall is hard and contains one seed. A seed nut planted in a coconut nursery is a fruit. So the fruit wall consists of the exocarp, mesocarp, endocarp (coconut husk), solid endosperm (coconut meat), liquid endosperm (coconut water) and embryo (Fig. 1). The coconut fruit has evolved to have three layers of peel: the exocarp is leathery, thin and smooth, providing effective protection against external moisture; the mesocarp is fibrous, and protects the fruit from breaking when dropped; the peel is bony, hard and not easily deformed, and can effectively protect the embryo and endosperm of the coconut.

The palm has distinctive characteristics emerge throughout the germination and seedling growth processes. The seeds have a tiny embryo and a large quantity of endosperm. The distal section of the embryo expands in size to form a haustorium, which remains within the seed and expands significantly when the endosperm disappears. When the haustorium develops, the outer surface, which is closely attached to the endosperm, extends significantly. A numerous, minute, and protuberant structure is particularly developed on the attached surface. The undulating structure is pale yellow in color. Endosperm breakdown products saturate the surface and lodge in the troughs caused by invaginations. The

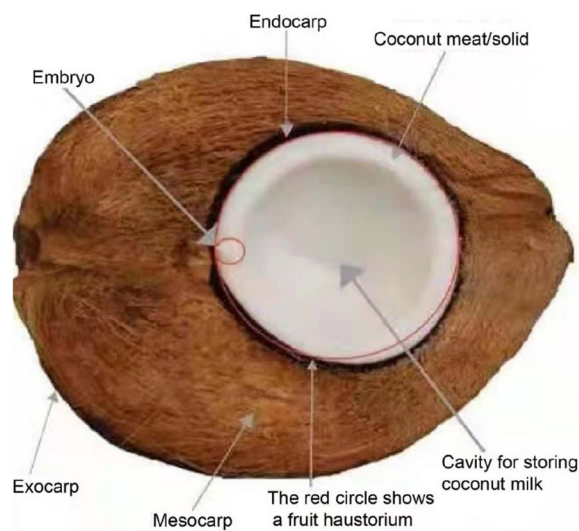


Fig. 1 The sectional structure of coconut fruit

epithelium, the haustorium's outermost layer, is composed of rectangular cells. During haustorium development, cells of the inner parenchyma progressively grow in size.

Coconut palms are divided into two types based on their physical traits and breeding habits: tall coconut palms and dwarf coconut palms. Tall coconuts cross-pollinate spontaneously, and the populations exhibit various degrees of heterozygosity [1]. Dwarf coconut, on the other hand, is naturally self-pollinating and exhibits typical morphological traits, including dwarf height, a thin trunk, a smaller crown, and small-sized nuts with relatively low copra content. Because of the coconut fruit's unique structure and growth, cultivators find it difficult to examine the inside, making non-destructive inspections even more crucial to ensuring natural growth. To overcome this issue an innovative radiographic imaging method known as X-ray computed tomography allows for non-destructive, non-invasive three-dimensional (3D) imaging at higher resolutions as well as utilizing a mathematical technique, researchers have examined the internal structure of the same fruit from several perspectives to determine the fruit's three-dimensional volume [2, 3].

The images were obtained by X-ray from the Computed Tomography (CT) machine has been proposed in different fruit. For instance, Cantre et al. [4] utilized CT to examine the microstructure of epidermal and sub-epidermal tissue in kiwi fruit, whereas Ting et al. [5] used CT to analyze the microstructures of various apple cultivars. On the other hand, CT were used to quantify and describe the internal structure of intact pomegranate fruit [6]. Few research were carried out to evaluate internal fruit disorders such mealiness in pears [7], internal browning in apples [8, 9], watercore in apples [10], and cracked stones in Japanese plums, CT has been used several times [11]. The results of all this research have contributed to an understanding of how internal disorders impact fruit microstructure and how these abnormalities might be prevented. More recently, CT was used to investigate the distribution of thermo-physical characteristics such porosity and thermal conductivity in Japanese apricot [12]. Janssen et al. [13] investigated the 3-dimensional (3D) pore structure of 'Braeburn' apples employing CT.

Therefore, we propose a CT-based non-invasive 3D quantitative imaging system for coconuts. As a consequence, in this research, we used the coconut as the object and developed a method for constructing a 3D model of the coconut using CT scan images. The purpose of this paper is to investigate the internal stratified physiological structure of coconut fruit while identifying its growth pattern through quantitative imaging model. The established model of coconut is highly

accurate and intuitive, forming a reliable non-destructive evaluation method, which can improve the screening rate of effective coconut fruits, reduce the presence of bad fruits, and improve the breeding efficiency of seedlings. It has a high application value in the field of coconut production.

Materials and methods

The process of establishing a non-invasive 3D quantitative imaging system using CT as a method for monitoring coconut fruit development (Fig. 2). We use CT axial scanning to obtain a set of cross-sectional images of coconut fruit [14]. Then the coordinates of the points in the cross-section image group are extracted, and the corresponding 3D point cloud images are generated by array transformation and stacking. The graph is then processed by K-nearest neighbor denoising. After fitting the outline of the outer surface, the system shows a complete three-dimensional quantitative model of the coconut fruit. Initially, CT coronal scanning of the coconut fruit was used to produce a series of cross-sectional images of the coconut fruit, which was then used to construct the point cloud model by extracting the 3D coordinate data of the cross-sectional image group and the RGB value data of the point. After that, a 3D quantitative model of coconut fruit was developed.

Cross-sectional image acquisition of the coconut fruit by CT coronal scan

In order to continuously observe the growth process of the coconut fruit, we first number the sample coconut fruit, then mark the posture position of the coconut fruit to ensure that it was in a fixed position in each scan. Then we used CT scanning to obtain the corresponding coronal scan image of each coconut fruit (Fig. 2b).

The coconuts were placed on a plastic shelf. Text and arrow markers were used so that the orientation of the image remained consistent throughout the cycle. The image was obtained using a 256 dual source CT scanner (Scanning instrument: SIEMENS SOMATOM Definition Flash) with the following parameters: tube voltage: 120 kV, tube current: 250 mas, temperature: 24 °C, Humidity: 50%. Use a set of coconut samples were scanned by a CT scanner(Fig. 3).

Point cloud model

After obtaining the CT image of the coconut, the next goal was to convert them into a point cloud model [15]. In MATLAB (MATLAB version: R2020b), the program first calls the ID of the target coconut, which is compressed and reconstructed into an array of 512*512*6. According to the Definition 1, the 6 columns are (x, y, z, r, g, b) . The converted data of each image are stacked, the points are drawn according to the corresponding coordinate

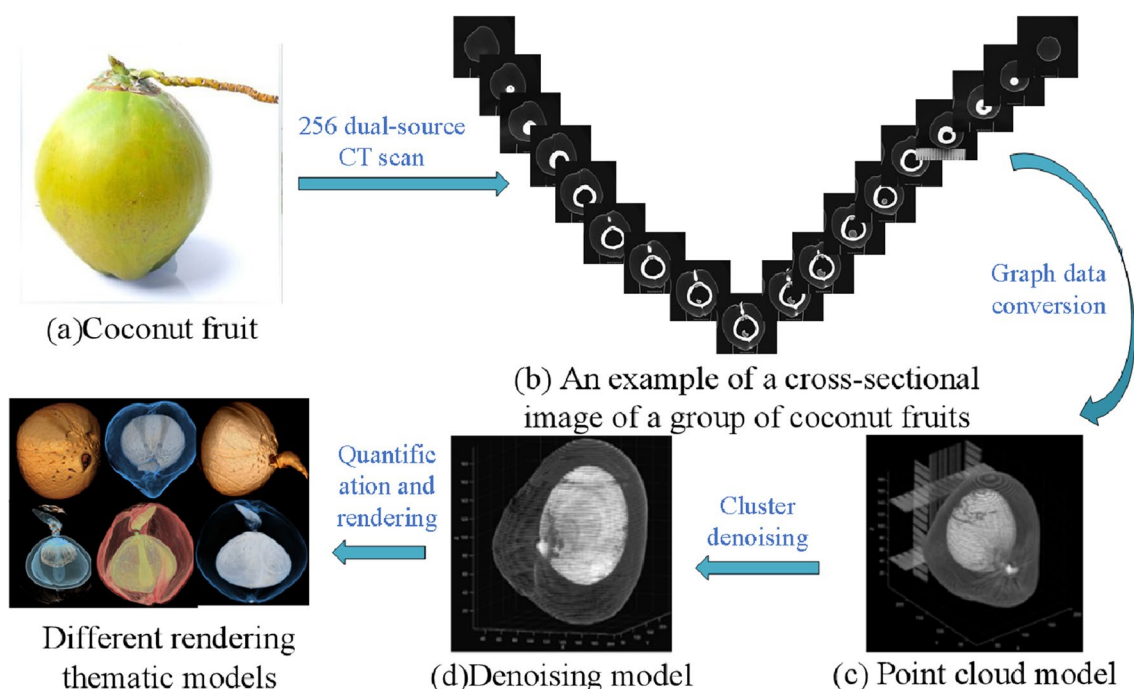


Fig. 2 An overview of coconuts reconstruction via 3D Quantificational Imaging

information values, and the 3D point cloud is displayed (Fig. 2c).

Definition 1 A coconut fruit point cloud model is a tuple $\langle X, Y, Z, R, G, B \rangle$, where X, Y and Z are the 3D coordinate data whereas, R, G and B are the Red, Green and Blue value respectively.

Algorithm 1: The calculation process of coconut CT images segmentation threshold

```

Input: Coconut CT images:  $D_C$ , Initial threshold:  $T_{initial}$ ;
foreach  $I_i \in D_C$  do
  Get Region  $D1, D2$  by split  $I_i$  with  $T_{initial}$ ;
  ( $D1$ : The gray value was greater than  $T_{initial}$ ,  $D2$ : The gray value was less than or equal
  to  $T_{initial}$ );
  Compute average gray value  $m1 = mean(D1)$ ,  $m2 = mean(D2)$ ;
   $T_i = (m1 + m2)/2$ ;
  while  $|T_{initial} - T_i| > 0.5$  do
    Get Region  $D1, D2$  by split  $I_i$  with  $T_i$ ;
    Compute  $m1$  and  $m2$ ;
     $T_i = (m1 + m2)/2$ ;
  end
   $T_{initial} = T_i$ ;
end

```

The algorithm 1 computes the segmentation threshold for coconut CT images. We implement the algorithm with Matlab. First, an initial threshold T is calculated based on manual experience, and then the image is segmented with T . $D1$ comprises all pixel components with gray values larger than T , whereas $D2$ includes all pixel components with gray values less than or equal to T . The average gray value $m1$ and $m2$ in each region are then computed, and the average value is used as the next threshold, and so on until the optimal threshold is established.

The equation for determining images after thresholding are

$$g(x, y) = \begin{cases} 1 & f(x, y) > T \\ 0 & f(x, y) < T \end{cases} \quad (1)$$

where $f(x, y)$ represents the original image, T is the finalized threshold, pixels marked with 1 correspond to objects and those marked with 0 are considered as background and invalid information.

(3) Conversion and merging of point cloud map data:

- a. The reshape function was used to reconstruct a specific matrix B to fit the dimension of the coconut point cloud, $B = reshape(A, size)$, where A

- (1) Pre-processing: The CT images derived from the direct scan of the CT machine are large, so they are compressed into 512*512 for ease of processing.
- (2) Image thresholding segmentation: recognition extracts the coconut part and distinguishes the effective information domain from the background of the CT images.

represents its own array of grayscale maps, and size was the reconstructed array dimension. In our work the value of size was 6.

- b. Repelem was a function that creates an array of repeating elements b . $b = repelem(A, r_1, \dots, r_N)$ repeats each element of A by r_1, \dots, r_N , returning an array where each element of r_1, \dots, r_N must be a scalar or vector with the same length as A in the corresponding dimension [16]. This was due to the fact that in the 3D coordinate system, one quantity was the same on each of the x, y , and z dimensions.
- c. Repmat was used to build a large matrix containing multiple repetition matrices B , $B = repmat(A, r)$, using the row vector r to specify the repetition scheme. $B = repmat(A, [mnp\dots])$ $B = repmat(A, n)$, with the contents of A stacked in $(M \times N)$ matrix B . The size of the B matrix was determined by $M \times N$ and the contents of the A matrix, with the same coordinates on the same axis, matching the 3D array created $B = repmat(A, n)$ returns an array containing n copies of A in its row dimension and column dimension. when A is a matrix, the size of B is $size(A) \times n$. $B = repmat(A, r_1, \dots, r_N)$, specifying a list of scalars r_1, \dots, r_N that describe how the copies of



Fig. 3 CT machine scans coconut samples

- A are arranged in each dimension. When A has N dimensions, the size of B is $size(A) \times [r_1, \dots, r_N]$.
- d. PCmerge was used to merge multiple point clouds. By the previous method, we get an array of point clouds for each layer, and then we need to merge and stack them to form a complete 3-dimensional point cloud map. This was expressed as: `ptCloudOut = pcmerge(ptCloudA, ptCloudB, ptCloudC..., ptCloudN, gridStep);` `ptCloud` refers to the point cloud data points, N = the number of coconut point cloud layers created, the merged point cloud was returned using the box grid filter, `gridStep` specifies the size of the filter's 3D box, and in addition, increases the size of `gridStep` when there were not enough resources to build a large fine-grained grid.

The merged point cloud was returned as a point cloud object, and the PCmerge function calculates the axial bounding box of the overlapping area between the two point clouds. The bounding box was divided into grid boxes of the size specified by `gridStep`. The points in each grid box are merged by averaging their positions, colors, and normals. Points outside the overlapping area are not affected according to the storage order of the image itself [17]. Because the coconut was placed in a fixed direction according to the orientation markings made during each scan, there was no need be concerned with the image alignment generated by CT. The first pixel of the previous image must correspond to the first pixel of the next image, and after repeated tests, the layer spacing of each image was set to 0.225 cm as the best. The point cloud

data are merged in this manner, thus constructing the coconut model. The 3D point cloud model of the coconut was constructed (Fig. 5a).

Model optimization

After the initial point cloud model was built, there were two main problems: irrelevant information and the existence of image noise, and the discrete discontinuity of points on the outer surface. These two factors would affect the effectiveness and accuracy of the model, so we carried out denoising and surface fitting to improve the quality of the overall model [18].

Denoising method

We used K-Nearest Neighbor (KNN) noise reduction method to remove the noise points and set the threshold to 0.13, determine the distance between the center point and the surrounding 50 points, take the average value and compare it with the threshold value, and categorize the points contained within the threshold distance as the valid information class; the remaining classes were deleted. Then for a new sample x , the k points in the training set that are nearest neighbors to x are found [19]. How is this nearest neighbor measured? We use L_p distance to measure it. The L_p distance of two vectors x_1, x_2 can be defined as $L_p(x_1, x_2) = \left(\sum_{i=1}^n |x_1^i - x_2^i|^p \right)^{\frac{1}{p}}$, next, find the class with the highest number of occurrences of the category among the k points and assign x to this class, expressed in a mathematical formula is

$$y = \arg \max c_j P_I(y_i = c_j), \quad i = 1, 2, \dots, N; j = 1, 2, \dots, K$$

, where i takes values from 1 to N , representing the number of samples in the training set, and j takes values from 1 to k , representing the number of categories in the training set. I is the indicator function, meaning that I is 1 when $y = c$, and I is 0 otherwise [20]. Thus obtain the effect after denoising (Fig. 5b).

Surface reconstruction and mesh quantization

The 3D model constructed by the point cloud method has the effect of displaying too many discrete points, which is not conducive to the integrity and attractiveness of the model. According to the relationship that defines the connection between two points (Fig. 4), we use the rolling ball method in order to construct the complete boundary information and calculate the associated volumes using integrals through the small grid thereby created.

- (1) For any point p , the radius of the rolling circle r , search all points within the distance $2r$ from the

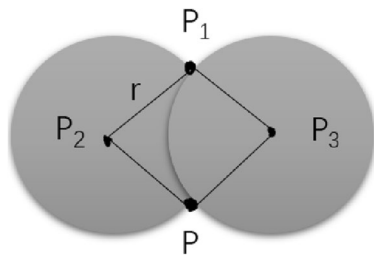


Fig. 4 Definition of the connection between two points

point p in the point cloud, and record it as the point set M .

- (2) Select any point $p_1(x_1, y_1)$ in M , and calculate the center coordinate based on the coordinates of these two points and r . Among them, $p_2(x_2, y_2)$ and $p_3(x_3, y_3)$ are the coordinates of the center of the circle when passing through the two points p and p_1 and the radius is r . The calculation formulas are:

$$x_2 = x + \frac{1}{2}(x_1 - x) - H \times (y_1 - y) \tag{2}$$

$$y_2 = y + \frac{1}{2}(y_1 - y) - H \times (x - x_1) \tag{3}$$

$$x_3 = x + \frac{1}{2}(x_1 - x) + H \times (y_1 - y) \tag{4}$$

$$y_3 = x + \frac{1}{2}(y_1 - y) + H \times (x - x_1) \tag{5}$$

$$H = \sqrt{\frac{\alpha^2}{S^2} - \frac{1}{4}} \tag{6}$$

$$S^2 = (x - x_1)^2 + (y - y_1)^2 \tag{7}$$

There will be some discrete independent points in the figure, because when the distance is greater than $2r$, they will be regarded as outliers and will not participate in the rolling process. The model effect of surface fitting can be obtained by this process (Fig. 5b)

Visualization

Because the point cloud features were composed of discrete points, the model lacked wholeness. We performed a closed meshing operation for surface fitting on the model. The alpha shape rolling ball method [21] was suitable for the feature requirements of the external coconut surface. To find the points that met the requirements, the radius of the rolling ball was set to 0.3 cm, while all external circles corresponding to the connection of points constituted the boundary information; the plot function was then used to draw them, and the color parameters were adjusted to remove the grid lines in order to obtain the model after fitting the discrete points.

At this point, the basic construction of the model was complete; it was then necessary to quantify it digitally [22]. The model can target any area to obtain its

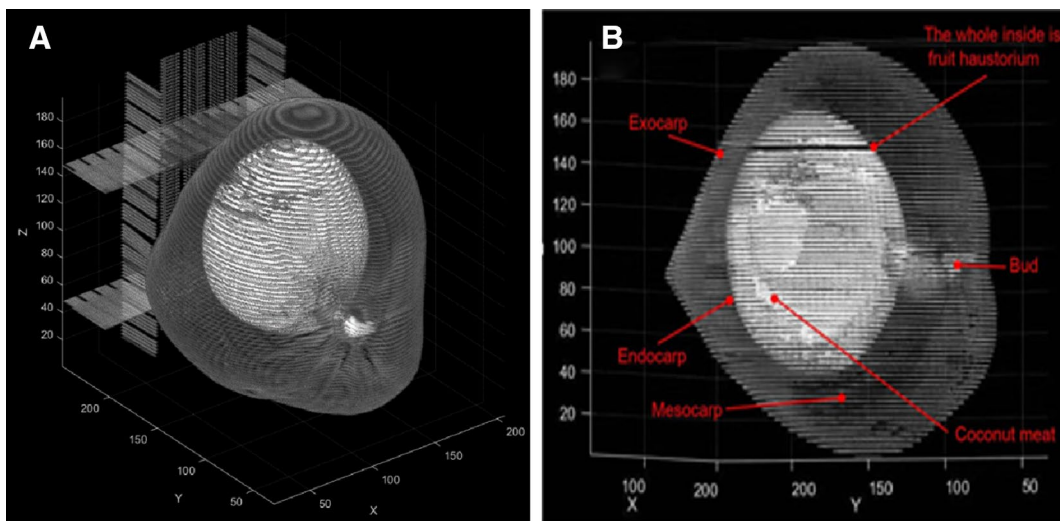


Fig. 5 3D model effect and denoising treatment. **A** a 3D point cloud image; **B** the model diagram after denoising and surface fitting

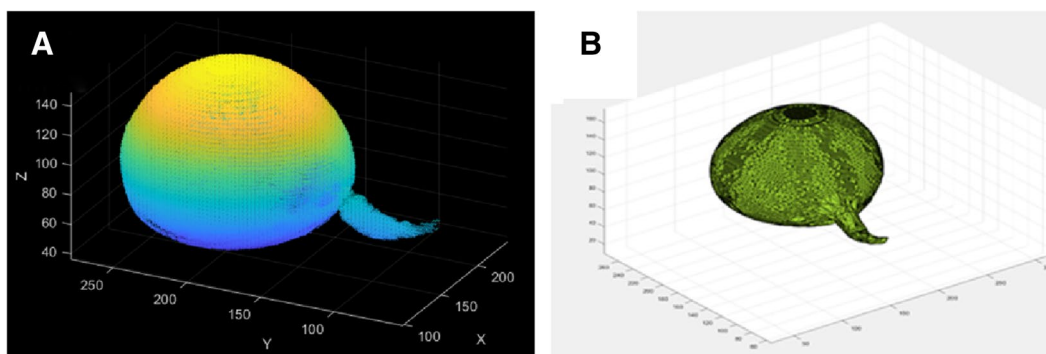


Fig. 6 A example of quantitative calculation of organ. **A** Extraction and coloring; **B** the haustorium after meshing

volume. The commonly used fruit haustorium was used as an example for the calculation. The haustorium fruit is extracted according to the RGB value [230,230,230]. Because the target in the color display is not conducive to observation and calculation, we subjected it to color processing (Fig. 6A). Then used alphashape to perform the grid operation. At this time, the haustorium was composed of small squares (Fig. 6B). Because the 3D coordinates of all points were known, the volume function could be used to obtain the volume value, the models of each coconut fruit in different stages are calculated according to this method [23], the volume value calculated by taking one of the models as an example is 602.9 cm^3 .

Model evaluation

After the above experimental work was completed, we then evaluated the accuracy of the constructed 3D model. We evaluated the model both internally and externally, conducted through in communication with experts at the coconut institute. For the internal factors, we used these four key sets of data as statistical indicators: the long axis of the haustorium fruit, the short axis of the haustorium fruit, the long diameter of coconut water, the short diameter of coconut water. For the external factors, we took the total volume and the triangular mesh number of the reconstructed coconut as the evaluation index. This can take the following form.

Definition 2 The absolute divergence of coconut is a tuple $\langle H_l, H_s, W_l, W_s, V_{ol}, N_{ob} \rangle$, where H_l and H_s represent the statistical difference of long and short axis of haustorium respectively, W_l and W_s represent the statistical difference of long and short diameter of coconut water

respectively. And when the error is less than 0.2 cm will be judged to be accurate. V_{ol} (volume) represents the volume of the model built, and N_{ob} (number of boundary triangle grid) represents the number of triangular meshes on the boundary, which is judged to be accurate as the volume is closer to the reference value and the number of triangular meshes is higher.

System prototype

The intelligent coconut analysis system (Fig. 7) is designed according to our working method and the practical application. The Browser-Server architecture is adopted in this system, the development mode is front-end and back-end separated, and the front-end part uses Vue. JS, the development tool is VS Code, and adds the Echarts plug in library. The back-end part uses the Java language, version 1.8, uses the Springboot framework.

When entering the system, followed by the “import the images” function (for uploading coconut CT images), the 3D point cloud (for converting images into the 3D point cloud), contour drawing (for connecting the discrete points of a point cloud map to make it more intuitive), color rendering (applying color according to one’s preferences can help improve the look or highlight important areas), and visual presentation (for outputting the final image after the above steps). Each function block is relatively independent and can be used according to demand.

The reconstructed 3D point cloud display structure is clear at a glance. Through the algorithm, the system can extract the required internal structure and obtain the relevant data. The system provides commonly used structural categories, such as coconut meat, coconut water, germ, coconut haustorium, coconut shell, and so on. After selection, the corresponding length and diameter,

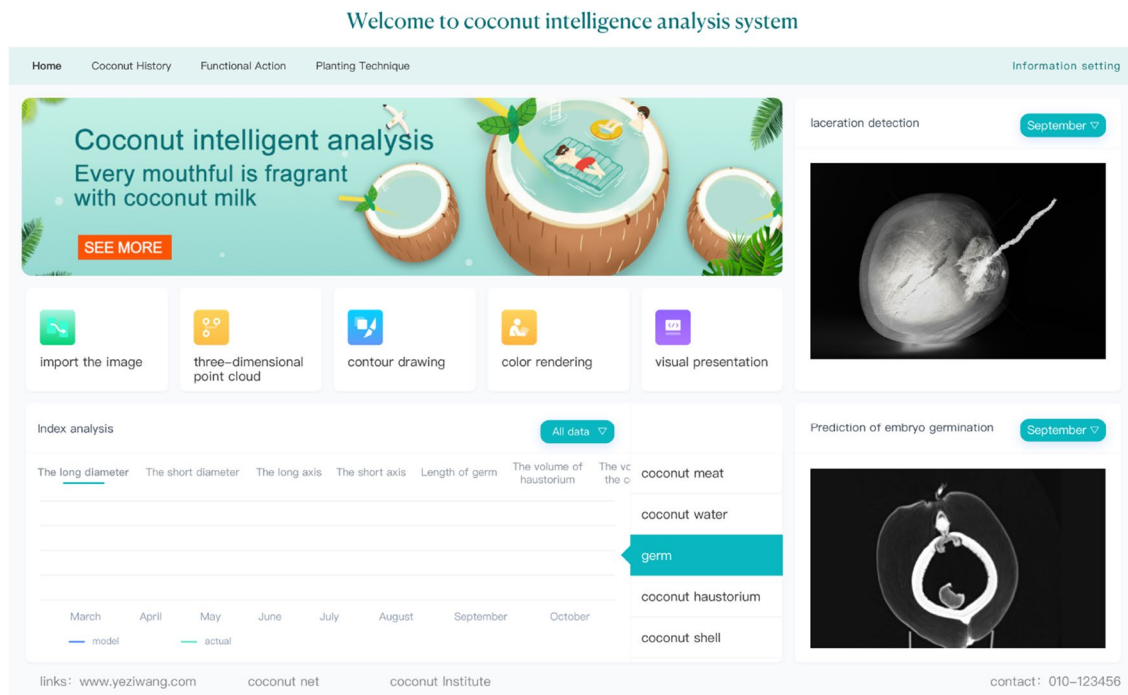


Fig. 7 Prototype demonstration of the system

long and short axis, volume and other index data can be obtained.

Result and discussion

In order to prove that the intelligent coconut system can effectively output 3D point cloud models of different coconuts and verify the high accuracy of the 3D model, we performed two experiments by way of illustration. We took 200 coconut fruits and observed them for weeks. Using CT scans, we used CT scanning to obtain a total of 37,950 non-destructive internal growth change maps of various types of coconuts to establish a coconut data set called “CCID”.

We next began our evaluation of internal factors, where the raw parameter values of the actual coconut fruit were measured by the CT machine, and the processed 3D model data thereby generated were measured using the MATLAB platform. Under the same experimental conditions and samples, we recorded the data of both the actual coconut fruit and its 3D model, and the weekly averages of these samples were analyzed and compared as a whole. The horizontal and vertical axes represent the measurement period and the values of each parameter, respectively (Fig. 8A and B). Figure 8A shows the values of these four factors of the coconut itself measured by the CT machine at different periods. And Fig. 8B shows the values associated with

the corresponding 3D model of the coconut at different periods. We observed the coconuts for twelve weeks. These two graphs not only show the growth changes of the internal structure of the coconut, but clearly indicate the small error between the model and the body in each time period. We then calculated their means and variances, further showing how the coconut fruit compared to the corresponding models in terms of important influences over multiple time periods (Table 1). These statistical differences are within the desired interval and strongly indicate the accuracy of our constructed model in terms of internal factors.

After the evaluation of the internal factors, we addressed the external factors [24]. In our work, two methods for meshing are used to express information, namely Alphashape and Convex Hull [25]. The results were compared to demonstrate which was more suitable.

The mesh reconstruction of the surface is performed using the rolling ball method (Fig. 9B). And we use the volume function in matlab to get its volume and use the numregions function to output the number of triangles in the surface shape, these two indicators are important factors to quantify whether the model is accurate from the outer surface. Correspondingly, we use the Convex Hull to perform surface reconstruction based on the 3D point cloud, the result (Fig. 9C) were obtained from the sample we selected. Then we allowed the program iterate through all the point sets, and based on the number

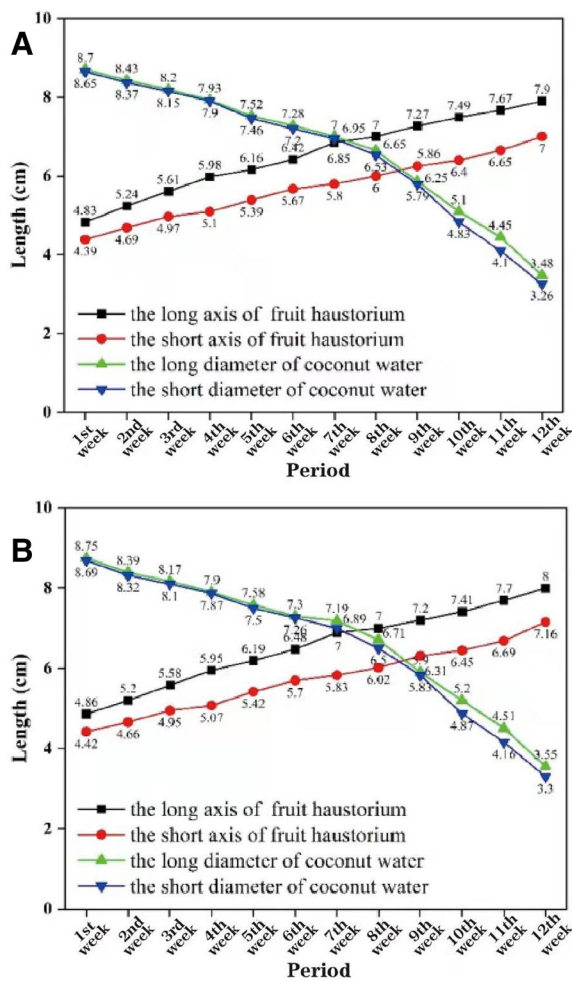


Fig. 8 Machine measurements, modeling data (same growth cycle). **A** Important parameter values of coconut fruit in different periods; **B** Corresponding parameter values in the 3D model

of vertices we can determine whether they were triangles or not, and thus to calculate how many triangles were included, as well as to the volume value as well.

Regarding the comparison between the volume value and the number of boundary grids, we used the data obtained when scanning coconuts on the CT machine as a reference value to compare the final results obtained

by the two methods. For the reconstruction of the outer surface, the number of regional grids to a certain extent reflects whether the construction is accurate. Through data comparison, it was found that the volume of the final model formed by the alpha shape method was closer to the reference value, and the number of triangular meshes contained in it was also more than that of the convex hull method. We also introduced the relative error rate. This concept was used to better show the degree of deviation of the results caused by the two methods. The volume value obtained by the alpha shape method was closer to the reference value, and the number of boundary grids was greater, indicating the number of captured points (Table 2). Moreover, the region description was more accurate and the overall error rate was smaller, which further proves the correctness of our selection of the alpha shape method to reconstruct the outer surface. The validity of our method is verified by the above comparison results from the internal structure and the external surface, and the accuracy of our constructed model is proved qualitatively and quantitatively.

Conclusion

A 3D quantitative imaging model for coconuts was developed, allowing non-destructive observation of the internal development of coconut fruit. The aim was to investigate the internal stratified physiological structure of coconut fruit while identifying its growth pattern through the computation of its relevant tissue and structure data, thereby contributing to coconut cultivation and improvement. The results reveal that the 3D quantitative imaging approach may be utilized to efficiently capture the internal development of coconut fruit. Simultaneously, the 3D point cloud model for coconuts established by this system has high precision, provides a powerful auxiliary function for enhancing coconut cultivation technology, and has a high practical application value in the coconut research sector.

In the future, we hope to utilize the deep learning network to detect cracking inside coconuts and forecast embryo germination. With continuous improvements in

Table 1 Summary of average and variance comparison results (cm)

Category	Attribute	Long axis of haustorium	Short axis of haustorium	Long diameter of coconut water	Short diameter of coconut water
Coconut fruit	Average	6.535	5.6925	6.7167	6.5992
	Variance	0.90172	5.60087	2.54586	2.82079
3D model	Average	6.5383	5.7233	6.7625	6.6167
	Variance	0.91813	0.65974	2.46632	2.75259

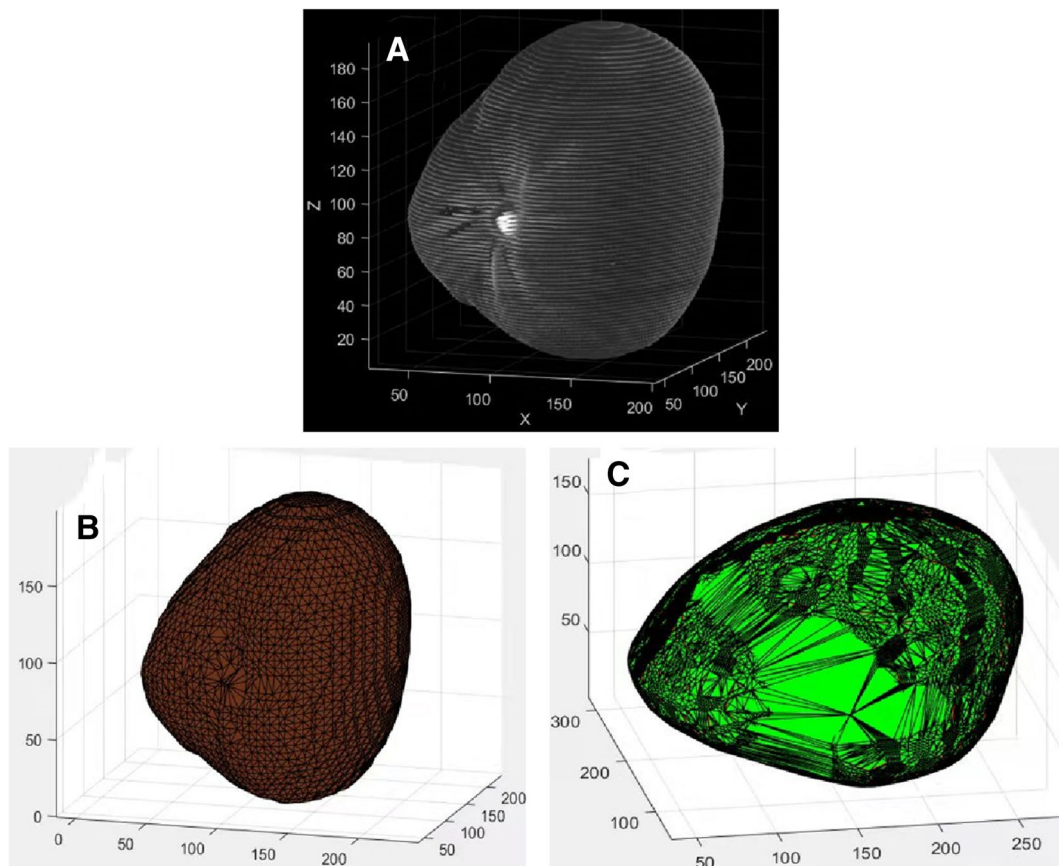


Fig. 9 Surface fitting. **A** 3D point cloud of the sample coconut; **B** Alphashape grid; **C** Convex Hull boundary

Table 2 Comparison of the results of outer surface reconstruction between the two methods

Method	Volume(cm ³)	Number of boundary triangle grid	Relative error of volume (%)
Alphashape	549.03	21038	4.10
Convex Hull	571.58	10934	8.38

Description: The volume reference value of this sample 527.39cm³, the relative error calculation formula is (actual value – reference value)/reference value

technology, imaging instruments and computing power, it is expected that CT can be fully implemented on the sorting line to categorize coconut based on maturity and internal quality. It is anticipated that technological and research advancements will allow for the development of reconstruction techniques and image processing algorithms that will achieve fast and affordable inline CT sorting systems for quality evaluation. We also intend to continuously add new functions to the system. The intelligent system developed herein provides a platform

to empower decision-making for researchers within the coconut industry.

Acknowledgements

The units participating in the cooperation are Coconut Research Institute-Chinese Academy of Tropical Agricultural Sciences, Hainan University, Haikou Municipal People's Hospital and Xi'an University of Technology. We would like to express our sincere gratitude to the above organizations for providing us with the necessary space, materials and equipment for the experiment.

Author contributions

YZ and QL completed the construction of the point cloud model, Fig. 1, 2, 3, 4 and wrote the first draft of this paper, J C developed the work plan of CT scan and experimental Figs. 5, 6 and Table 1, CS and HC provided coconut samples and completed coconut model validation experiments, and completed Figs. 8, 9 and Table 2, SL completed the software development of coconut intelligent analysis system (Fig. 7), ZX and MH guide the whole work and complete the paper optimization. All authors read and approved the final manuscript.

Funding

This work was supported in part by the Major Science and Technology Project of Haikou (Grant #: 2020-009), in part by the Key R & D Project of Hainan province (Grant #:ZDYF2021SHFZ243), in part by the National Natural Science Foundation of China (Grant #: 62062030, 82260362), in part by the National Key Research and Development Program of China (Grant #: 2021ZD0111000, 2018YFB1404400).

Availability of data and materials

You can contact the first author or corresponding author for coconut CT images and experimental data. Their e-mail addresses are: yuzhang2015@hainanu.edu.cn, hnsuncx@qq.com.

Declarations**Ethics approval and consent to participate**

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Author details

¹School of Computer Science and Technology, Hainan University, Haikou, China. ²Radiology department, Central South University Xiangya School of Medicine Affiliated Haikou Hospital, Haikou, China. ³Coconut Research Institute, Chinese Academy of Tropical Agricultural Sciences, Wenchang, China. ⁴School of Computer Science and Engineering, Xi'an University of Technology, Xi'an, China. ⁵School of Information and Communication Engineering, Hainan University, Haikou, China.

Received: 21 July 2022 Accepted: 27 February 2023

Published online: 09 March 2023

References

- Arumugam T, Hatta MAM. Improving coconut using modern breeding technologies: challenges and opportunities. *Plants*. 2022;11:3414.
- Landis EN, Keane DT. X-ray microtomography. *Mater Charact*. 2010;2010(61):1305–16.
- Nicolai BM, Defraeye T, Ketelaere B, Herremans E, Hertog MLATM, Saeys W, et al. Nondestructive measurement of fruit and vegetable quality. *Annu Rev Food Sci Technol*. 2014;5:285–312. <https://doi.org/10.1146/annurev-food-030713-092410>.
- Cantre D, East A, Verboven P, Araya XT, Herremans E, Nicolai BM, Pranamornkith T, et al. Microstructural characterisation of commercial kiwifruit cultivars using X-ray micro computed tomography. *Postharvest Biol Technol*. 2014;92:79–86.
- Ting VJL, Silcock P, Bremer PJ, Biasioli F. X-ray micro-computer tomographic method to visualize the microstructure of different apple cultivars. *J Food Sci*. 2013;2013(78):E1735–42.
- Magwaza LS, Opara LU. Investigating non-destructive quantification and characterization of pomegranate fruit internal structure using X-ray computed tomography. *Postharvest Biol Technol*. 2014;95:1–6.
- Muziri T, Theron KI, Cantre D, Wang Z, Verboven P, Nicolai BM, Crouch EM. Microstructure analysis and detection of meakiness in 'Forelle' pear (*Pyrus communis* L.) by means of X-ray computed tomography. *Postharvest Biol Technol*. 2016;120:145–56.
- Herremans E, Verboven P, Bongaers E, Estrade P, Verlinden BE, Wevers M, et al. Characterisation of 'Braeburn' browning disorder by means of X-ray micro-CT. *Postharvest Biol Technol*. 2013;75:114–24.
- Chigwaya K, Schoeman L, Fourie WJ, Crouch I, Viljoen D, Crouch EM. 'Fuji' apple internal browning explored via X-ray computed tomography (CT). *Acta Hort*. 2018;1201:309–16.
- Herremans E, Verboven P, Defraeye T, Rogge S, Ho QT, Hertog MLA, et al. CT for quantitative food microstructure engineering: the apple case. *Nucl Instruments Methods Phys Res Sect B Beam Interact with Mater Atoms*. 2014;324:88–94.
- Kritzinger I, Lötze E, Jooste M. Stone hardening and broken stones in Japanese plums (*Prunus salicina* Lindl.) evaluated by means of computed tomography scans. *Sci Hort*. 2017;2017(221):1–9.
- Karmoker P, Obatake W, Tanaka F, Tanaka F. Visualization of porosity and thermal conductivity distributions of Japanese apricot and pear during storage using X-ray computed tomography. *Eng Agric Environ Food*. 2019;12:505–10.
- Janssen S, Verboven P, Nugraha B, Wang Z, Boone M, Josipovic I, Nicolai BM. 3D pore structure analysis of intact 'Braeburn' apples using X-ray micro-CT. *Postharvest Biol Technol*. 2020;159: 111014.
- Herrero-Huerta M, Meline V, Iyer-Pascuzzi AS, et al. 4D structural root architecture modeling from digital twins by X-ray computed tomography. *Plant Methods*. 2021;17:123. <https://doi.org/10.1186/s13007-021-00819-1>.
- Jiaxin Wang, Fuqun Zhao. Research on point cloud data preprocessing. *Modern Inf Technol*. 2020;4(2):129–30.
- Hui L, Boxiong W, Huaiyi R, Xiuzhi L. Research on two-way point cloud denoising method based on 3D reconstruction data. *J Electron Meas Instrum*. 2013;27(1):1–7.
- Huijun Y, Ruiping W, Zengying W, Xin W. Three-dimensional phenotype reconstruction of crop fruits based on multi-view images. *J Nanjing Normal Univ*. 2021;44(02).
- Li Q, Gao X, Fei X. Tree canopy 3D model construction using alpha-shape algorithm. *Mapp Bull*. 2018;(12).
- Zhang Q, Chen H, Zhu S. Application of density clustering algorithm in denoising of continuously distributed point clouds. *Geospatial Inf*. 2011;9(06):101.
- Zhang L, Wang M. Point Cloud Data Segmentation Based on Discrete Expansion of K-Neighborhood. *GUIDE Softw*. 2009;(12).
- Edelsbrunner H, Mücke EP. Three-dimensional alpha shapes. *ACM T Graphic*. 1994;13:43–72.
- Paturkar A, Sen Gupta G, Bailey D. Plant trait measurement in 3D for growth monitoring. *Plant Methods*. 2022;18:59. <https://doi.org/10.1186/s13007-022-00889-9>.
- Bin L, Junbo W, Wang L, Bochao M, Mingxia X. A comparative analysis of two point cloud volume calculation methods. *Int J Remote Sens*. 2019. <https://doi.org/10.1080/01431161.2018.1541111>.
- Ziamtsov, Navlakha, Saket, Su, Hao. An algorithmic toolbox for plant phenotyping with 3D point clouds, UC San Diego, 2021.
- Yoel S, Moti S, Avishai H. Shaping the way from the unknown to the known: the role of convex hull shape in numerical comparisons. *Cognition*. 2021;217: 104893.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

