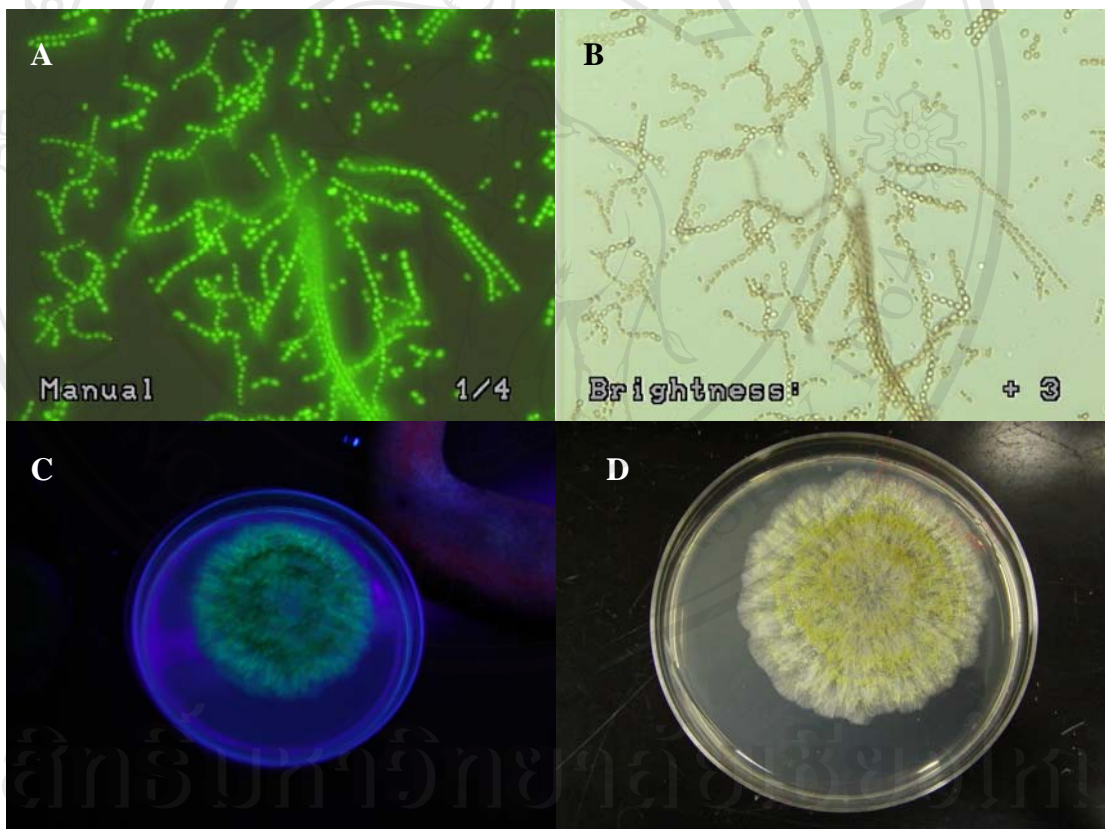


APPENDIX

Appendix A Conidia of GFP *Aspergillus flavus* observed under (A) an UV-illuminated microscope and (B) a white light microscope. *A. flavus* colony observed when illuminated with (C) UV light and (D) normal light.



Appendix B A minirhizotron camera system.



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Appendix C Installation and operation of QuaCos (Quantification of Color System)

Installation

The QuaCos directory includes files named **Quacos.cab**, **setup.exe**, and **setup.lst**. To install, use either Windows Explorer double click on **setup.exe** or click **Start** then **Run**. Select **Browse** to find path to **setup.exe** and then click **Ok**. We recommend installing QuaCos under the program director, which is set as the default.

Operation

1. Open QuaCos, for example, by selecting **Start**, the **Programs**, and finally **QuaCos**.

Please select Disclaimer, and then read and accept the terms of using QuaCos.

2. Select **Ok**, which will lead to the display of a window with four tabbed cards at the bottom.

3. Under the **Input Option** tab select the path where the images to be analyzed are stored.

4. Click the **Output Option** tab to select the color values to be stored and the path for storing data.

To analyze an entire image:

5. Select the **Full Image Analysis** tab.

6. Select the size of pixel groups to be averaged. Note that the smaller the size of pixel groups, the longer the analysis requires. It necessary, you may also rotate the data to match the orientation of the image.

7. To analyze a single image select the **Single** button. To analyze all images in a folder using the same setting select the **Batch** button.

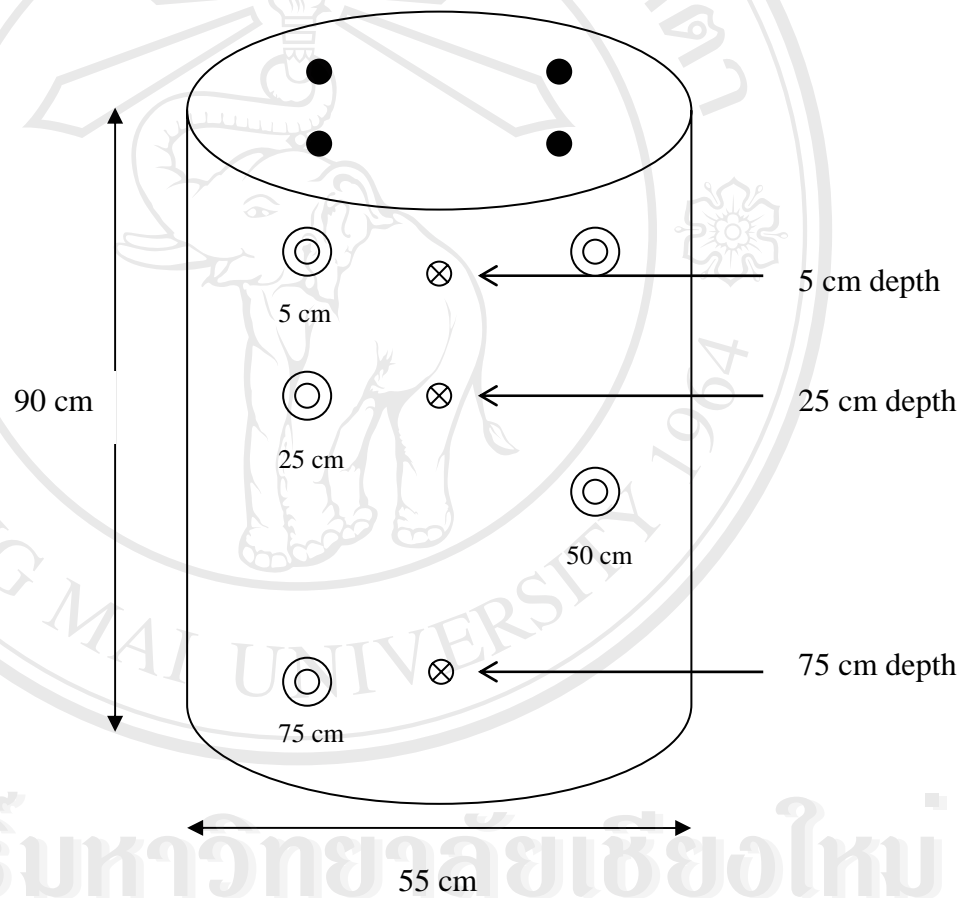
8. To analyze a portion of an image: Select the **Partial Analysis** tab.

9. Select area to be analyzed from options provided, that is, 10×10 , 25×25 , 50×50 , 75×75 , 100×100 , or 500×500 pixels. The area selected must be smaller than the image.
10. Position the selector over that portion of the image to be analyzed.
11. Select the **Analyze** button to analyze a single area. Repeat the process to analyze other areas of the image. Alternatively, select **Batch** to analyze the same area of all images in the folder.

Displaying Data

Data are stored in comma delimited ASCII format. They are named as image_name_color.map. To open these files with Excel (© Microsoft), first open Excel, select **File**, **Open** and browse to locate data. Select **Delimited** then **Next**, **Comma** then **Next**, and **Finish**. To display the data, we find it convenient to create a graph using **Surface** chart type with the **Contour-color** option.

Appendix D Diagram of 214-L container fitted with minirhizotron tubes at 5, 25, 50, and 75 cm. Soil moisture blocks and thermocouples installed at 5, 25, and 75 cm depths.

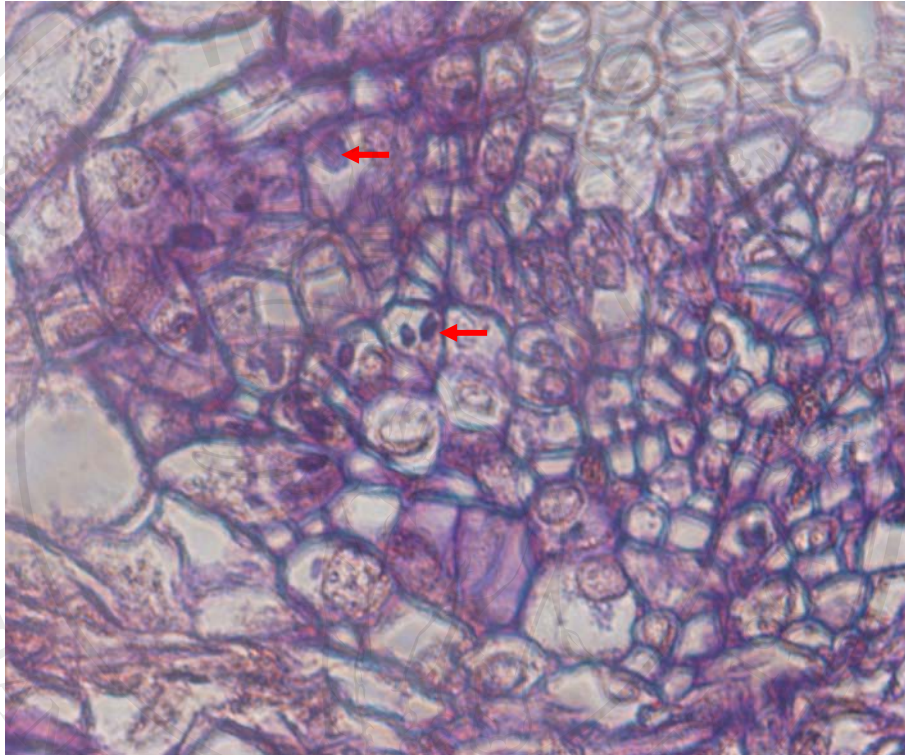


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Appendix E Wilting symptom of plants received T4 compared to T2 water treatment.



Appendix F Crossed section of seed coat of 511CC genotype. The red arrows showed tannin compound in epidermis cell.



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Appendix G Fixative and stain used for preparation of slide.**1. Formalin-acetic acid-alcohol (FAA) mixture**

Ethyl alcohol (50%)	90.0 ml
Glacial acetic acid	5.0 ml
Formalin	5.0 ml

2. Cotton blue lactophenol

Anhydrous lactophenol	67.0 ml
Distilled water	20.0 ml
Cotton blue	0.1 g

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