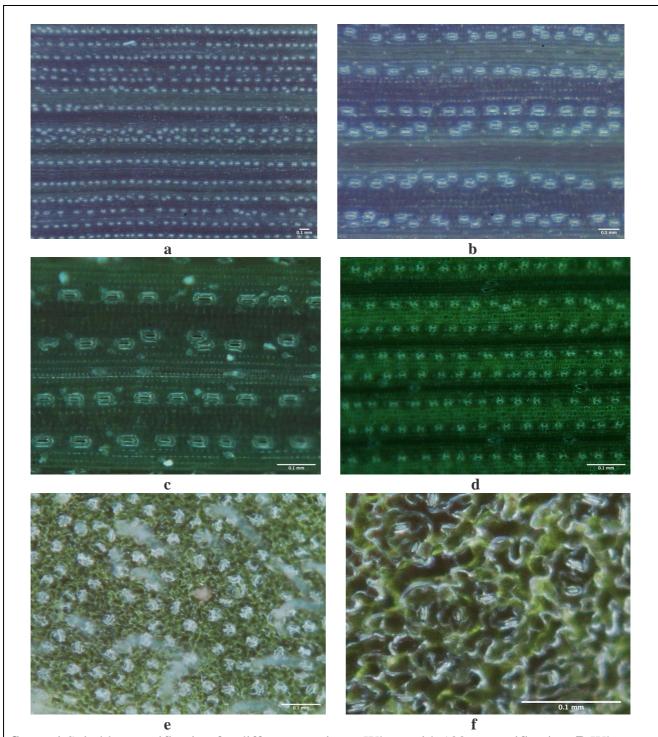
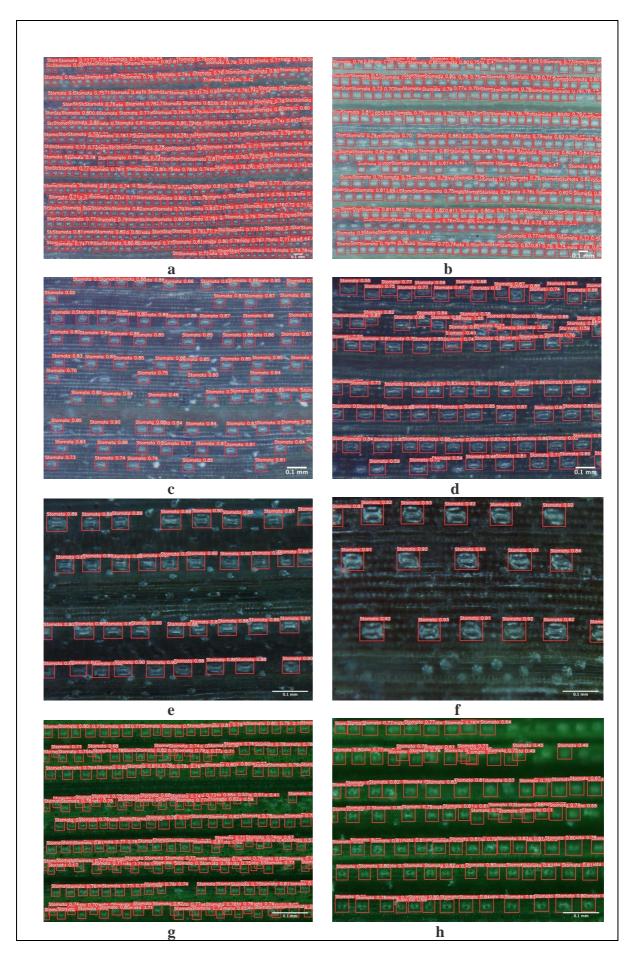
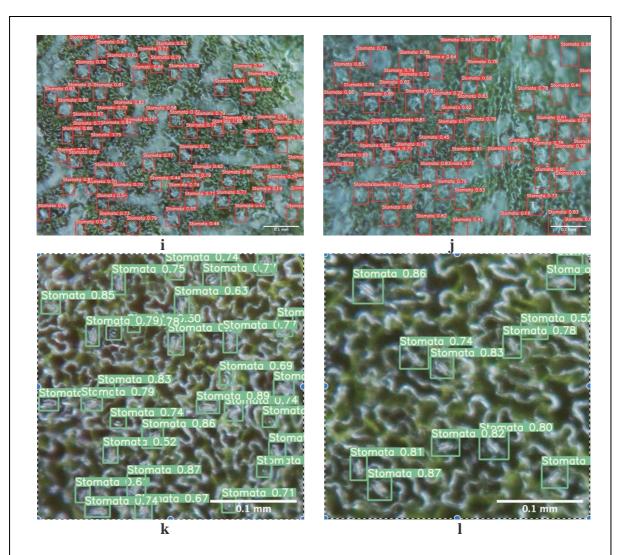
# Supplementary figures



**Supp. 1** Suitable magnification for different species. **a** Wheat with  $100 \times$  magnification. **B** Wheat with  $200 \times$  magnification. **c** Wheat with  $400 \times$  magnification. **d** Rice with  $400 \times$  magnification. **e** Tomato with  $400 \times$  magnification. **f** Arabidopsis with  $400 \times$  magnification. Scale = 0.1 mm





**Supp. 2** Stomata detection models identify the differences in stomata size within the same data set. **a** 100x wheat smallest stomata trait. **b** 100x wheat largest stomata trait. **c** 200x wheat smallest stomata trait. **d** 200x wheat largest stomata trait. **e** 400x wheat smallest stomata trait. **f** 400x wheat largest stomata trait. **g** 400x Rice smallest stomata trait. **h** 400x Rice largest stomata trait. **i** 400x Tomato smallest stomata trait. **j** Tomato largest stomata trait. **k** 400x Arabidopsis smallest stomata trait. **l** Arabidopsis biggest stomata trait. Scale bar = 0.1 mm

**Supp. 3** Link to Stomata detection model Google Colab

8 https://drive.google.com/drive/folders/1MX-

PBlCtKsBIjkJSN9PQQf3gfAVX4mNZ?usp=sharing

#### **USER GUIDE**

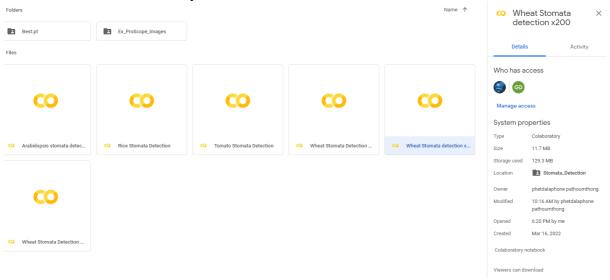
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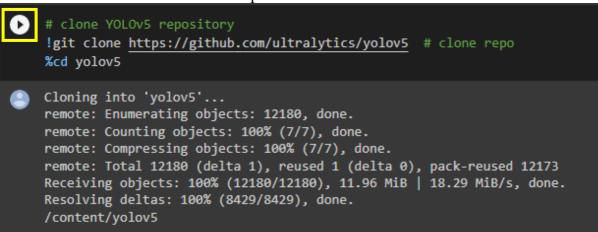
2223

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- 14 Rapid non-destructive method to phenotype stomatal traits
- 15 Phetdalaphone Pathoumthong<sup>1</sup>, Zhen Zhang<sup>2</sup>, Stuart Roy<sup>1</sup>, Abdeljalil El Habti<sup>1\*</sup>
- <sup>1</sup>School of Agriculture, Food and Wine, The University of Adelaide, Urrbrae 5064, Australia
- <sup>2</sup>Australian Institute for Machine Learning, The University of Adelaide, Adelaide 5000,
- 18 Australia
- \* Corresponding author: abdeljalil.elhabti@adelaide.edu.au
- 21 It is important to have good quality images that allow the model to recognise stomata.
- 24 Click on the link <a href="https://drive.google.com/drive/folders/1MX-">https://drive.google.com/drive/folders/1MX-</a>
- 25 PBlCtKsBIjkJSN9PQQf3gfAVX4mNZ?usp=sharing. If you can't open it using Chrome,
- open in Microsoft Edge
- 27 Copy Collab folder "Wheat Stomata detection x200" on your Google Drive so you don't
- 28 have to look for the link every time



#### 31 Click on the start button of the first script



Click on Run anyway

# Warning: This notebook was not authored by Google.

This notebook was authored by **phetdalaphone1989@gmail.com**. It may request access to your data stored with Google, or read data and credentials from other sessions. Please review the source code before executing this notebook. Please contact the creator of this notebook at phetdalaphone1989@gmail.com with any additional questions.

Cancel

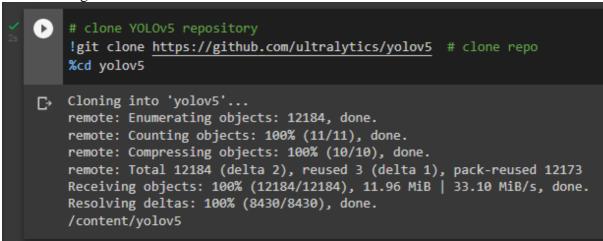
Run anyway

34 35

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You should get



37 Click on the second script

```
# install dependencies as necessary
!pip install -qr requirements.txt # install dependencies (ignore errors)
import torch

from IPython.display import Image, clear_output # to display images
from utils.google utils import gdrive_download # to download models/datasets

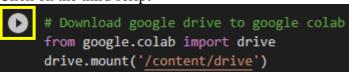
# clear_output()
print('Setup complete. Using torch %s %s' % (torch.__version__, torch.cuda.get_
```

38 39 You should get

```
# install dependencies as necessary
    !pip install -qr requirements.txt # install dependencies (ignore errors)
    import torch
    from IPython.display import Image, clear_output # to display images
    from utils.google_utils import gdrive_download # to download models/datasets
    print('Setup complete. Using torch %s %s' % (torch. version , torch.cuda.get device
                                         | 596 kB 29.0 MB/s
₽
    ModuleNotFoundError
                                              Traceback (most recent call last)
    <ipython-input-2-d9ae8b9a3cb3> in <module>()
          5 from IPython.display import Image, clear_output # to display images
    ----> 6 from utils.google_utils import gdrive_download # to download models/datasets
          8 # clear output()
    ModuleNotFoundError: No module named 'utils.google_utils'
   NOTE: If your import is failing due to a missing package, you can
    OPEN EXAMPLES
                   SEARCH STACK OVERFLOW
```

Ignore the error message

### 42 Click on the third script



43 44

Connect to Google Drive

### Permit this notebook to access your Google Drive files?

This notebook is requesting access to your Google Drive files. Granting access to Google Drive will permit code executed in the notebook to modify files in your Google Drive. Make sure to review notebook code prior to allowing this access.

No thanks

Connect to Google Drive

45

- 46 Choose the account that will contain images
- 47 Sign in
- 48 Allow

## Make sure you trust Google Drive for desktop

You may be sharing sensitive info with this site or app. You can always see or remove access in your **Google Account**.

Learn how Google helps you share data safely.

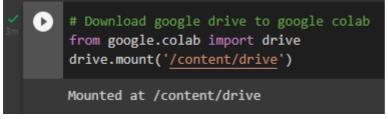
See Google Drive for desktop's Privacy Policy and Terms of Service.



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You should get



Before clicking on the fourth script

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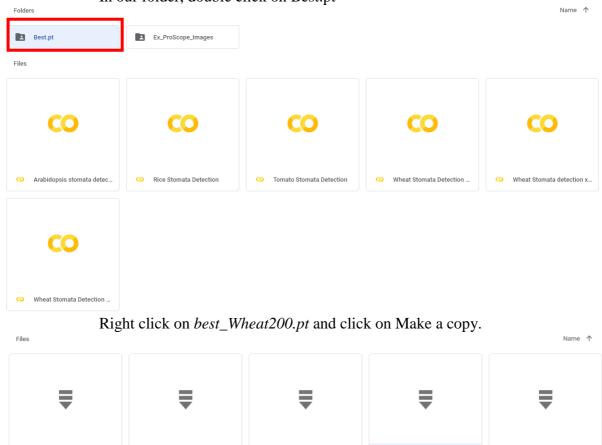
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Copy machine learning model to your Google Drive.

In our folder, double click on Best.pt



best\_x100.pt

Name ↑

■ best\_Arabidopsis.pt

 □ best\_Rice.pt

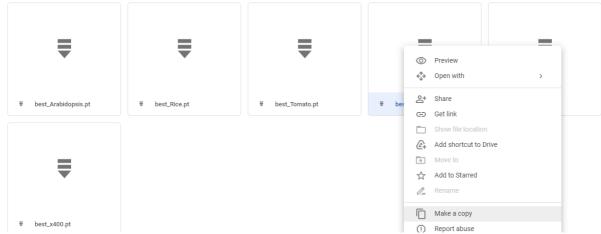
 □ best\_Tomato.pt

 □ best\_wheat200.pt

 □ best\_wheat200.pt

 □ best\_x400.pt

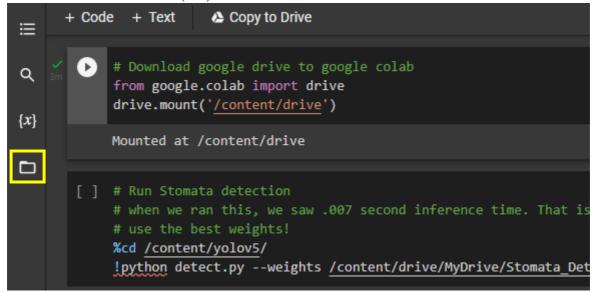
Files



This folder will now appear in your Google Drive. No need to paste the file in your Google Drive.

 Create new folder in your Google Drive and transfer images you would like to analyse

 Click on folder tab (left)



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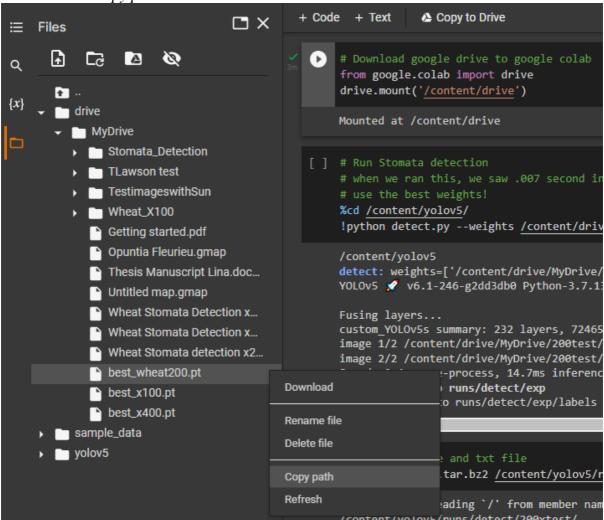
84

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Specify model location

Click on drive – MyDrive – click on three dots next to best\_wheat200.pt –

copy path



Replace

"/content/drive/MyDrive/Stomata\_Detection/Best.pt/best\_wheat200.pt" by pasting the copied path

```
# Run Stomata
 cd /content/yolov5/
```

Specify image set location

Click on drive - MyDrive - click on three dots next to the image folder - copypath

Paste after source. Add \ when there is a space, add / at the end before save txt

```
# 052 (content/yolov5/
!python detect.py --weights /content/drive/MyDrive/Stomata_Detection/Best.pt/best_wheat200.pt --img 416 --conf 0.4 --source /content/drive/MyDrive/200test/ --save-txt
[ [Errno 2] No such file or directory: '/content/yolov5/'
    python3: can't open file 'detect.py': [Errno 2] No such file or directory
```

For example, if the file path is

- "/content/drive/MyDrive/TLawson test", it will become
- "/content/drive/MyDrive/TLawson\ test/"

#### 86 Click on the start button

```
# Run Stomata detection

# when we ran this, we saw .007 second inference time. That is 140 FPS on a TESLA P100!

# use the best weights!

%cd /content/yolov5/
!python detect.py --weights /content/drive/MyDrive/Stomata_Detection/Best.pt/best_wheat200.p

[Errno 2] No such file or directory: '/content/yolov5/'

/content

python3: can't open file 'detect.py': [Errno 2] No such file or directory
```

You should get

```
# Run Stomata detection
# when we ran this, we saw .007 second inference time. That is 140 FPS on a TESLA P100!
# use the best weights!
%cd /content/yolov5/
!python detect.py --weights /content/drive/MyDrive/best_wheat200.pt --img 416 --conf 0.4 --source /content/

/content/yolov5
detect: weights=['/content/drive/MyDrive/best_wheat200.pt'], source=/content/drive/MyDrive/TLawson test/, d
Y0L0v5 / v6.1-246-g2dd3db0 Python-3.7.13 torch-1.11.0+cu113 CPU

Fusing layers...
custom_Y0L0v5s summary: 232 layers, 7246518 parameters, 0 gradients
image 1/1 /content/drive/MyDrive/TLawson test/wheat stomata.jpg: 416x326 75 Stomatas, Done. (0.228s)
Speed: 0.9ms pre-process, 228.3ms inference, 11.6ms NMS per image at shape (1, 3, 416, 416)
Results saved to runs/detect/exp/labels
```

The number of stomata for each image is highlighted in blue. Copy the results and paste it in an Excel file.

94 Click on the fifth script



95 96

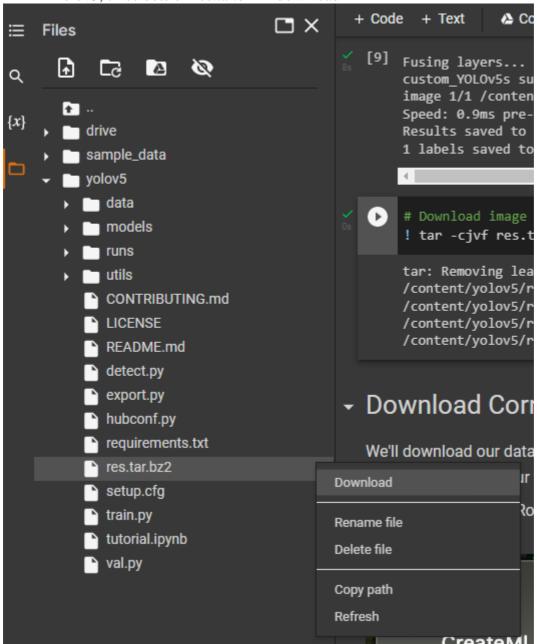
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Go to folder at the left

Yolov5 – runs – detect – rename exp

Yolov5, three dots on res.tar.bz2 – download



99 100

You will find this folder in your Downloaded folder.

This folder contains: stomata size in a txt file, one file per image; labelled images.