

PHenOSPEx

Smart Plant Analysis

Machine Learning Engineer - Take Home Assignment

Plant Counting and Localization Under Edge Constraints

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1. Problem Definition

The objective of this work is to design a pipeline that **counts individual plants and estimates their centroid locations** from **2D and 3D sensor data**. Given an input, the system must output the **total number of plants**, and the **coordinates** of each plant's center.

This task is formulated as a **dense localization problem** rather than a **bounding-box detection problem**. Instead of predicting object extents, the focus is on accurately identifying **plant centers**. This formulation naturally supports **heatmap-based regression approaches**, where the model learns a **continuous spatial representation of plant likelihood**.

In addition to accuracy, the solution must satisfy practical **deployment constraints** such as: **CPU-only inference, offline operation, low memory usage and fast runtime** and a **reproducible and lightweight training pipeline**.

For the assignment, two approaches are implemented: a **machine learning-based solution** as the primary method and a **classical computer-vision baseline** for comparison.

2. Dataset Description

This work utilizes the provided phenotyping dataset with varying **grid configurations and plant counts**, focusing exclusively on **2D top-down images**. The **2D modality** was selected to prioritize **computational efficiency, low inference latency, and CPU-only deployment**.

An annotated dataset was built and split into **50 training images, 12 validation images, and 12 test images**, preserving the **distribution of grid setups** across all splits. An additional **augmented test set of 36 images** is constructed for **robustness evaluation**.

2.1. Annotation Procedure and Assumptions

Plant centroid annotations are generated using a **semi-automated annotation pipeline using OpenCV**. Centroids are defined as **2D points (x, y)** , corresponding to the **approximate stem base** and assumed to align with the **center of the pot**. While more accurate **centroid ground truth** could be obtained from **3D point cloud data**, this approximation is considered **sufficient for the proposed approach**.

3. ML Method: Architecture, Training Process, and Rationale

The ML approach is formulated as a **heatmap regression** problem and implemented using a lightweight **U-Net** architecture. This formulation is well suited to **plant layouts** in which instances have **high visual similarity** and are primarily defined by their **center locations** rather than precise object boundaries. Given an input image, the model predicts a **single-channel heatmap** where each plant center is represented by a **2D Gaussian peak**. Plant centroids are obtained by detecting **local maxima** in the predicted heatmap, and the **total plant count** corresponds to the number of detected peaks.

Compared to **bounding-box detectors** (e.g., YOLO-style models), heatmap regression avoids **anchor design, bounding-box regression, and non-maximum suppression**, which are unnecessary for **point-level localization**. **Semantic segmentation** approaches, while expressive, introduce **higher computational cost** and **annotation complexity** without directly optimizing centroid accuracy. By directly supervising **spatial plant locations**, the proposed method yields a **simpler training objective, lower inference overhead, and greater stability**, making it well suited for **CPU-only deployment under edge constraints**.

3.1. Architecture Differences from Original U-Net

The proposed model includes the core **U-Net encoder-decoder structure with skip connections** but is **simplified**. This variant uses **fewer encoder-decoder stages** and **reduced channel widths**, resulting in a **significantly lower parameter count** and **improved CPU inference speed**. This **compact design** achieves a **favorable trade-off** between **localization accuracy** and **inference efficiency** under **strict latency and memory constraints**.

Unlike the **original U-Net**, which is typically trained for **semantic segmentation** using **pixel-wise classification losses**, the proposed model outputs a **single continuous heatmap** and is trained using **mean squared error loss** on **Gaussian targets**.

3.2. Training Procedure

Ground-truth plant centroids are converted into **isotropic Gaussian kernels** centered at each centroid, which are summed to form a single **ground-truth heatmap** per image. All images are resized to **512 × 512** in

the model prior to training.

The network is trained to regress heatmaps using **mean squared error (MSE)** loss and optimized with the **Adam** optimizer. To **stabilize convergence**, a **ReduceLROnPlateau** learning rate scheduler is employed, reducing the learning rate when validation loss plateaus. **Early stopping** is applied to mitigate overfitting and avoid unnecessarily long training runs. **Best model checkpoint selection** is based on validation loss.

Initial training is performed **without data augmentation** to establish a baseline. Subsequently, augmented training is done using **rotations, occlusions, and noise** to improve **robustness and generalization**, and evaluated on an augmented test set. Due to the **limited number of irregular grid samples**, an **oversampling strategy** is applied during training to increase their contribution and mitigate dataset imbalance.

4. Classical Computer Vision Baseline

A **classical computer-vision pipeline** is implemented to provide an alternative **baseline**. This baseline relies on **manually tuned thresholds** and **morphological parameters**. The pipeline begins with **region cropping** to remove **irrelevant background areas** outside the **plant grid**. **Bright glare** and **table surfaces** are then suppressed using **color-space thresholding**, followed by **binary segmentation of plant regions** via **grayscale conversion, denoising, and Otsu thresholding**. In a **post-processing stage**, **small noise components** are removed and **fragmented plant regions** are merged using **morphological operations**. Finally, **connected components** are identified, and **plant centroids** are computed from the resulting **components**.

5. Results

Performance is evaluated using:

- **F1 score:** Where **predicted centroids** are **greedily matched** to **ground-truth centroids** within a **fixed distance threshold**.
- **Localization error:** **Mean Euclidean distance (in pixels)** between **matched predicted and ground-truth centroids**. The **localization error** is reported only over **matched pairs** within a **5 px threshold for ML approach**, resulting in values **bounded by this threshold**.
- **Count error** is defined as the **mean absolute error (MAE)** between **predicted and ground-truth plant counts**, computed **per image**.

5.1. Effect of Hyperparameters on the ML Model

A set of training runs was conducted to assess key hyperparameters, including the **Gaussian heatmap sigma (σ)**, the **oversampling weight (OSW)** for irregular grid samples, and the **sampling multiplier (SM)**.

The **Gaussian sigma** has the strongest influence on performance, with **larger values outperforming smaller ones**. Also, applying OSW during training helps mitigate dataset imbalance and improves overall results.

Further experimentation could yield better results; however, based on the current test set results, the configuration (**$\sigma = 3.0$, OSW = 25, SM = 2**) is selected. While this setting shows a slight reduction in localization accuracy compared to the **OSW = 12** configuration, it achieves a **lower counting error (MAE) and higher F1**, where the minor localization trade-off is acceptable given the improved true-positive detection rate.

Table 1: Test-set results of key ML experiments.

σ	OSW	SM	F1	Loc. Err (0–5 px)	Count Err (MAE)
3.0	1	1	0.973	1.62	1.25
3.0	25	2	0.971	1.61	0.83
3.0	12	2	0.968	1.53	2.08
1.0	12	2	0.750	2.38	14.33

5.2. Robustness and Augmented Test Evaluation

Robustness is evaluated by testing the final model on both the **test set** and the **augmented test set**, comparing training conducted **with** and **without data augmentation**. Models trained **without data augmentation** generalize poorly to the augmented test set. In contrast, models trained **with data augmentation** maintain strong performance across both test conditions. These results demonstrate that data augmentation is critical for achieving robustness and reliable generalization.

Table 2: Test and augmented test performance with and without data augmentation.

Aug.	Test			Test-Aug		
	F1	Loc. Err (0–5 px)	Count Err (MAE)	F1	Loc. Err (0–5 px)	Count Err (MAE)
No	0.971	1.71	0.83	0.404	4.07	25.81
Yes	0.984	1.58	0.58	0.957	1.88	1.64

5.3. Comparison with Classical Baseline

Comparison of the **best ML model** with a **classical CV baseline** on both the **test** and **augmented test set** was done. The **classical method** performs **poorly** under a **5 px matching threshold**, which is expected given the **mismatch between pot-based ground truth and leaf-based segmentation**. However, relaxing the threshold to **50 px** still results in **substantially inferior performance** compared to the ML model.

Table 3: Best ML model vs. classical baseline under different matching thresholds.

Method	Test		Test-Aug	
	F1 Score	Loc. Err (px)	F1 Score	Loc. Err (px)
U-Net-Tiny (best)	0.983	1.57	0.959	1.81
Classical CV (5 px)	0.043	3.08	0.013	3.24
Classical CV (50 px)	0.624	21.31	0.188	23.49

5.4. Runtime Comparison

For training, **GPU acceleration** was used to reduce training time. On CPU, a single epoch required approximately **3:30 min**, whereas in GPU it is **1:30 min**. Peak resource usage during GPU training was **4.7 GB of VRAM** and approximately **1.2 GB of system RAM**.

For inference, on **CPU**, the **classical pipeline** runs in **0.15 s per image**, while the **ML model** requires **0.60 s** with **PyTorch** and **0.25 s** after **ONNX** export, remaining only slightly slower while offering substantially improved accuracy and robustness.

6. Deployment Considerations

For **deployment**, the trained model is exported to **ONNX** and validated. The exported model size is **7.35 MB**, making it suitable for **resource limited environments**. **Runtime evaluation** using **ONNX Runtime on CPU** shows an **average inference time** of **0.23 s per image**, which is adequate for **near real-time runs**. **Memory usage** was measured using **resident set size (RSS)**. Loading the ONNX model increased memory usage from about **500 MB** to **540 MB**. During inference, **peak memory usage** reached approximately **900 MB**, mainly due to the **runtime environment**, **image data**, and **intermediate tensors**, rather than the model itself.

Overall, the **small model size**, **reasonable CPU runtime**, and **framework-independent ONNX format** make the proposed method suitable for **practical deployment**. Further **memory and runtime reductions** could be possible using a **lighter runtime** (e.g., C++ ONNX Runtime).

7. Limitations and Future Improvements

The **proposed approach** relies on **simplifying assumptions**, notably that **plant centroids** are approximated by the **pot center**, which ignores both **depth information** and the true **geometric centroid** of the plant.

The model is trained on a **limited dataset** with mostly **regular grid layouts** and **visually similar plants**, which can restrict **generalization** to other **plant species**, **growth stages**, and **grid configurations**, even though **data augmentation** improves **robustness**.

Another limitation arises from the **dependence on pot visibility**. Because annotations are defined at the **pot center**, the model may primarily learn **pot-specific visual cues**. Consequently, performance can **decline** when pots are **occluded**, even if the plant remains visible, leading to **false negatives**. Conversely, **false positives** may occur when a pot is present without a plant. These effects are observed in samples containing pots without plants, where rarely false predictions are produced.

Future work can include using **3D data** to obtain **accurate centroid annotations**. Expanding to **larger and more diverse datasets** may further improve **generalization** under **accuracy**, **runtime**, and **resource** constraints. Additionally, introducing **negative annotations** for empty pots, applying **post-processing filters**, or adopting a **plant segmentation model** could mitigate pot-only detections.