

1. Introduction & Motivation

- **Context.** Intercropping wheat (cereal) with pea (legume) improves soil health and reduces chemical inputs.
- **Problem.** Evaluating it needs per-species plant density across hundreds of micro-plots: manual counting does not scale.
- **Research question.** Can a point-based model localize and count each species separately from smartphone photos?
- **Contributions.**
 - Adapt PET to per-species wheat/pea counting in intercropping
 - Identify the optimal input resolution for small plant structures
 - Quantify edge-border filtering trade-offs (bias vs. localization)

2. Proposed Approach

- **Step 1: Data.** INRAE STICSMIX field trial; wheat labeled at leaf tips (orientation-invariant), pea at plant center. 116 wheat / 63 pea images, split at the plot level (train/val/test).
- **Step 2: Model.** PET [1], a VGG16-bn point-query transformer, finetuned from a crowd-counting checkpoint.
- **Step 3: Test-time levers.** Input resolution (1500-4000 px) and edge-border point filtering.

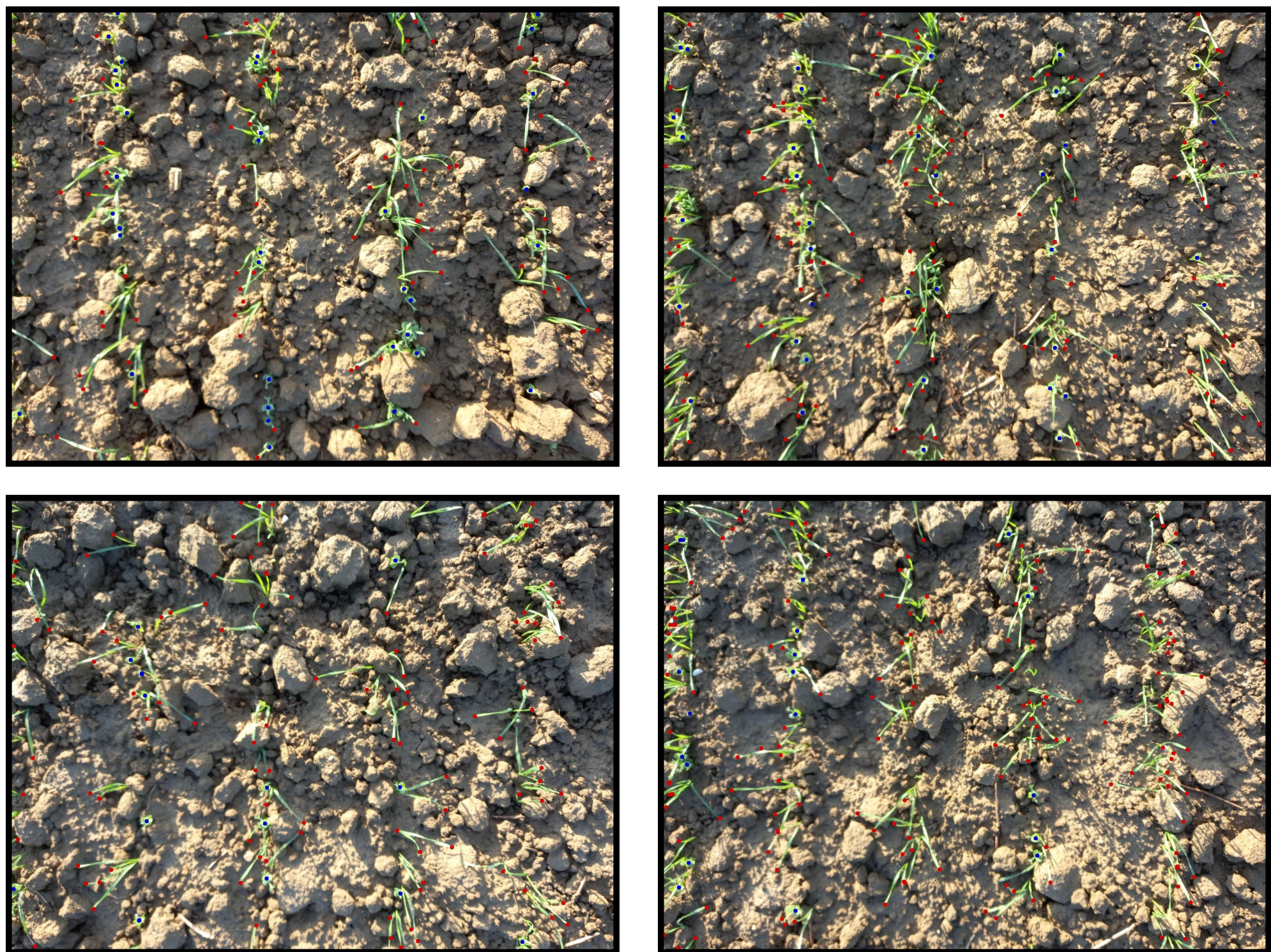


Fig. 1. Point annotations on two mixed wheat-pea plots, two photos each (red: wheat leaf tips, blue: pea plant centers; 1500 px, border filtering on).

3. Experimental Setup

- **Metrics** (greedy point matching, fixed distance tolerance):
 - MAPE: count error (%) ↓
 - R^2 : count variance explained ↑
 - Signed bias (%): + over-counting, - under-counting, closer to 0 is better
 - F1 (%): point-level localization ↑
- **Grid.** 5 resolutions × {with, without} edge-border filtering, per species.
- **Implementation.** VGG16-bn backbone; finetuned from a crowd-counting checkpoint; model selected on validation MAE.

4. Results

- **Best trade-off:** wheat 1500 px + border → **4.7% MAPE, 74% F1** · pea 2048 px, no border → **7.8% MAPE, 86% F1**.
- MAPE can mislead: the best-MAPE resolution is not always the best-localized one (FP/FN cancellation).
- Beyond 2048 px, both species degrade: pea detection **collapses at 4000 px** ($R^2 = -4.33$).

Species	Res.	Without border				With edge border			
		MAPE	R^2	Bias(%)	F1(%)	MAPE	R^2	Bias(%)	F1(%)
Wheat	2048	8.74	0.90	-1.65	62.3	12.18 _{-1.44}	0.81 _{-0.09}	11.30 _{-0.05}	65.9 _{+0.6}
	1500	9.03	0.90	-7.49	72.3	4.69 _{-1.34}	0.97 _{-0.07}	-0.48 _{-0.01}	74.2 _{+0.3}
	2500	9.73	0.87	0.28	62.1	10.50 _{-0.17}	0.91 _{-0.04}	9.53 _{-0.05}	64.4 _{+0.3}
	3000	12.54	0.79	4.25	53.4	18.11 _{-0.07}	0.75 _{-0.04}	16.76 _{-0.05}	55.9 _{+0.5}
	4000	13.77	0.63	-13.02	63.7	7.15 _{-0.02}	0.89 _{-0.01}	-4.14 _{-0.01}	66.7 _{+0.0}
Pea	2048	7.83	0.82	0.70	86.2	8.59 _{+0.19}	0.90 _{-0.05}	4.00 _{-0.01}	86.5 _{+0.3}
	1500	9.65	0.68	-7.07	82.6	9.66 _{+0.03}	0.80 _{-0.02}	-1.59 _{-0.04}	84.9 _{+0.3}
	2500	9.56	0.87	-0.13	84.5	9.82 _{+0.05}	0.93 _{-0.01}	6.88 _{-0.01}	85.6 _{+1.1}
	3000	13.15	0.78	1.31	76.4	15.33 _{-0.18}	0.81 _{-0.03}	7.37 _{+0.01}	78.1 _{+1.7}
	4000	67.94	-4.33	-67.70	33.5	64.11 _{-0.03}	-2.95 _{-1.06}	-64.11 _{-0.01}	36.2 _{+0.7}

Table 1. Test-set performance across input resolutions, with/without edge-border filtering. Subscripts: Δ vs. no-border (green = gain, red = loss). **Bold** = best per column within each block.

5. Discussion

- Border filtering drops truncated points from both GT and predictions ⇒ fewer boundary false negatives (F1↑), but fewer counted plants ⇒ over-counting bias.
- At native resolution, pea plants occupy far more pixels than PET's crowd-counting pretraining scale, a likely driver of the 4000 px collapse.
- F1 alongside MAPE is essential: MAPE alone can hide false-positive/negative cancellation.
- **Limits:** single trial, single growth stage.

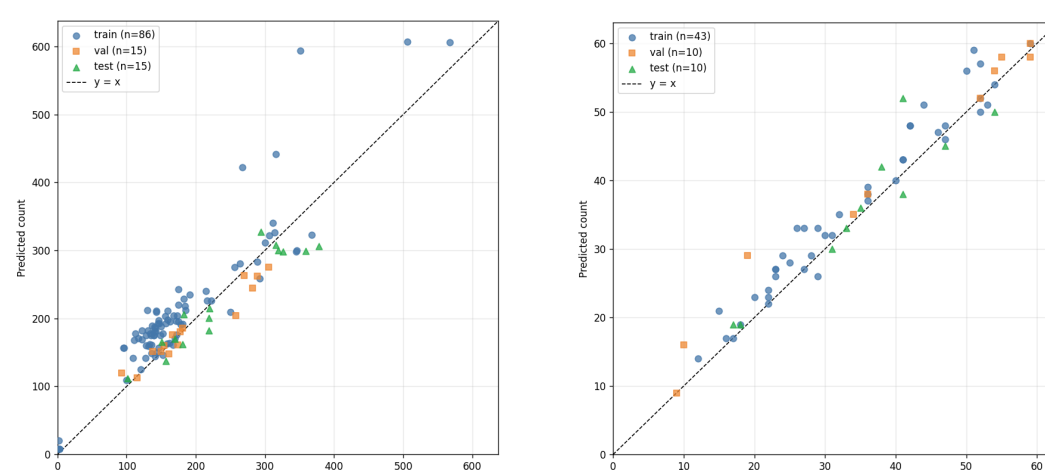


Fig. 2. Predicted vs. ground-truth counts, wheat (left) / pea (right), across train/val/test splits (dashed: $y=x$).

6. Conclusion & Future Work

- **Take-home.** PET reliably counts wheat and pea separately from smartphone photos, reaching single-digit MAPE and 74-86% F1 at the right input scale.
- **Field gap.** To our knowledge, no general baseline exists yet for per-species point-based counting in intercropped fields.
- **Next steps.**
 - Count both species jointly in a single pass, instead of per-species models
 - Cross-reference counts with post-harvest yield data

References & Acknowledgments

[1] Liu, C., Lu, H., Cao, Z., Liu, T.: Point-query quadtree for crowd counting, localization, and more. ICCV (2023).

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paper / code