

Artificial Intelligence (AI)-Driven Estimation of Total Kidney Volume (TKV) in ADPKD

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Results

Background

ADPKD characteristics:

- Fluid-filled cysts
- Bilateral renal enlargement
- Eventual renal failure

Regulatory guidance:

US FDA supports using imaging-based Total Kidney Volume (TKV) assessment as a qualified prognostic biomarker in clinical trials

Al-driven methods:

- Enable automated kidney segmentation
- Provide high repeatability

The study objective:

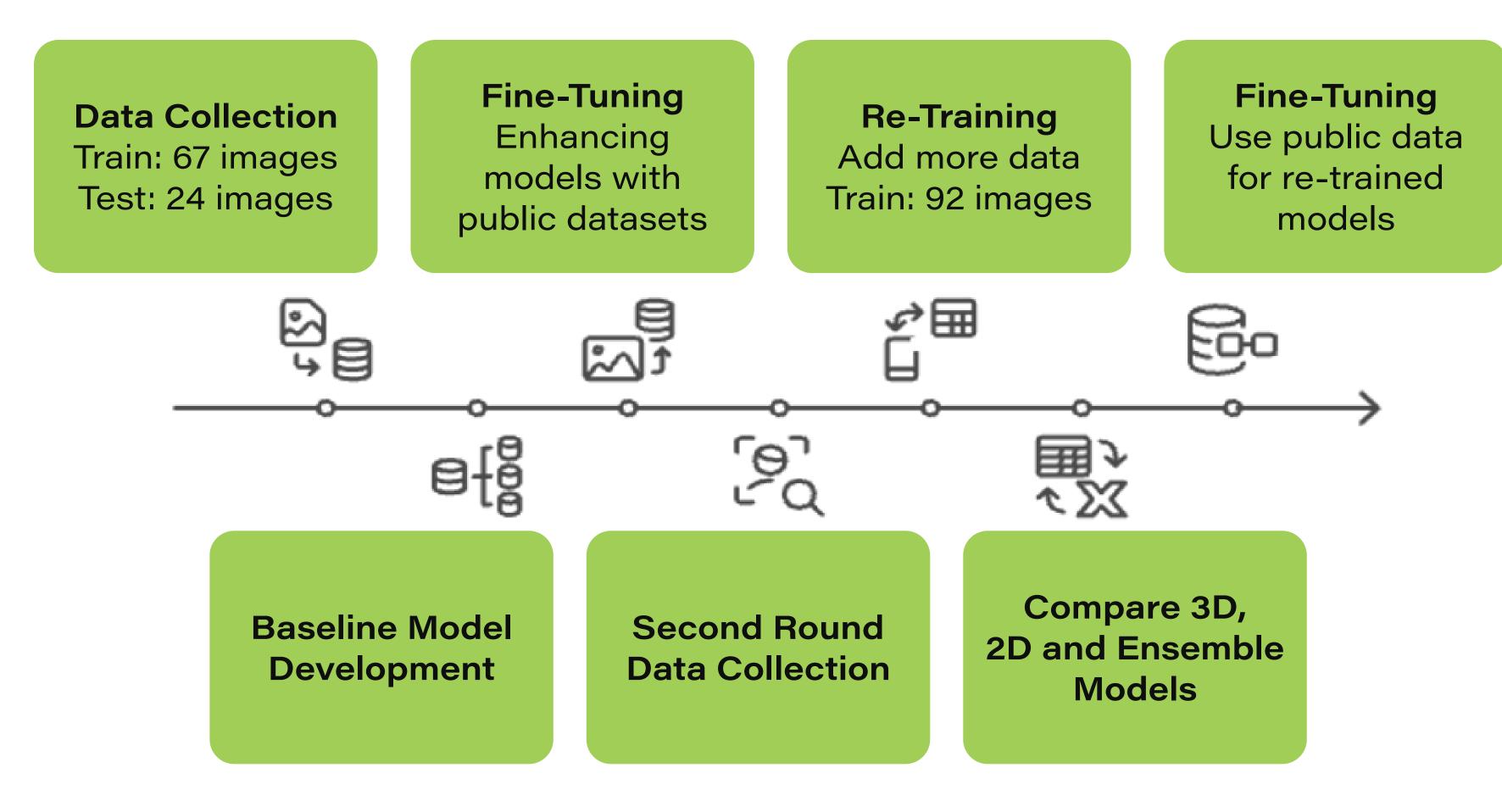
- Develop an Al-driven methodology to estimate TKV on MRI
- Ensure reproducibility

Method

A retrospective cohort of MRI scans from ADPKD patients enrolled in the CRISP study was analyzed (T1w and T2w images). One reader (15 yrs experience) annotated the T2w images.

Data Preparation: 1) registering T1w images rigidly to corresponding T2w images, 2) splitting the dataset into training, validation and test sets.

Model Development:



Public Datasets: MRI (AMOS, CHAOS, Totalsegmentator, CKD) and CT (AbdomenCT-1K).

Evaluation Metrics: The performance of all trained models was evaluated on the held-out test set using Dice Similarity Coefficient (DSC), Volume Error (VE), Hausdorff Distance (HD) and Intersection over union (IoU).

Best Initial Training (67 images, 24 test cases) Expanded Training (92 images, hold-out test set)

Total Kidney Performance (Initial Test Set):

- ► DSC: 0.843
- Hausdorff: 39.1 mm
- ► IoU: 0.745

0.90

Volume Error: 155,135 mm³

3D Model (T2_3D):

DSC: 0.897, IoU: 0.822, Volume Error: 133,531 mm³

2D Model (best performance) (T2_2D):

► DSC: 0.917, IoU: 0.852, Volume Error: 106,522 mm³

Ensemble Model (T2_Ensemble):

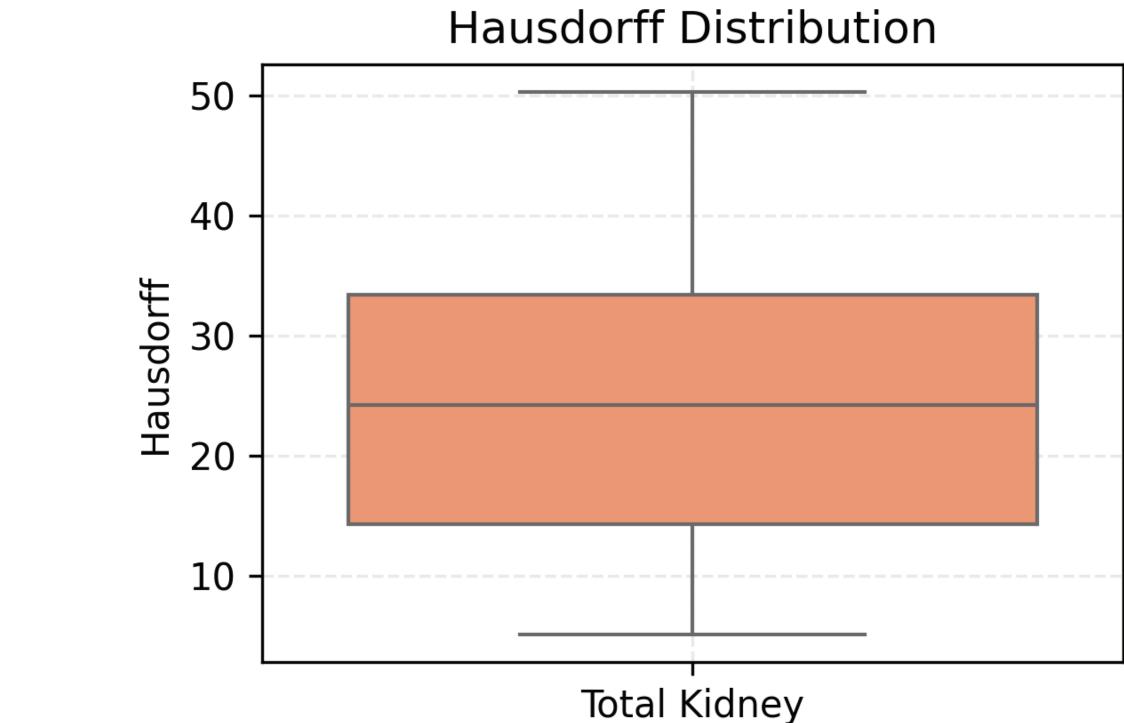
- ► DSC: 0.920, IoU: 0.856, Volume Error: 113,495 mm³
- ► Hausdorff: 21.1 mm (improved boundary accuracy)

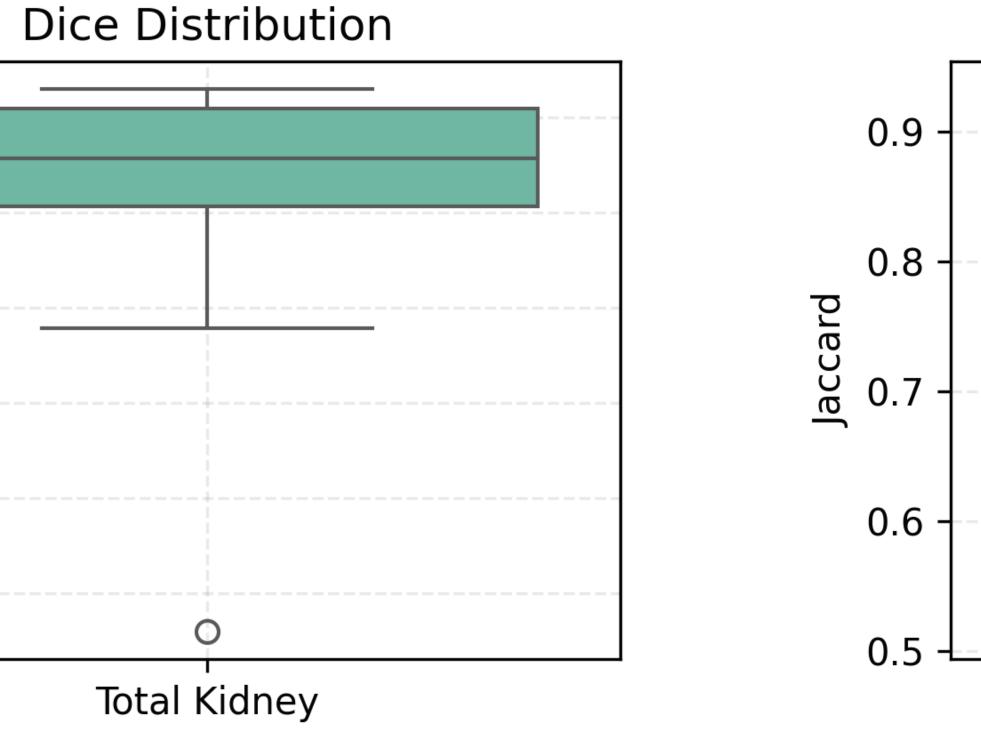
Fine-Tuning (T2_FT):

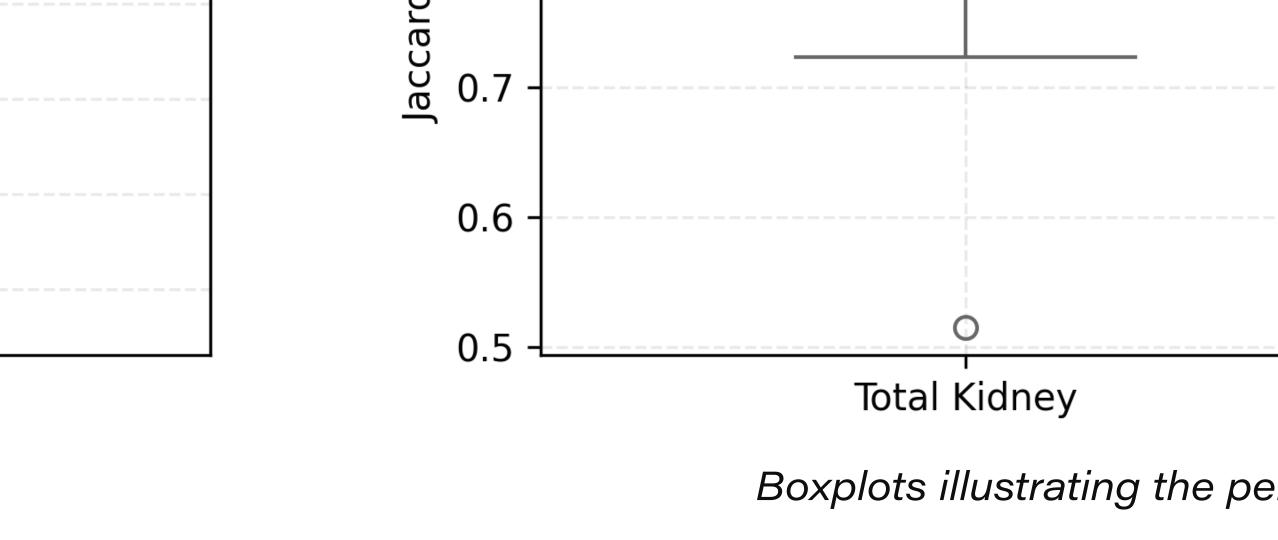
- ► DSC: 0.84, IoU: 0.75, Volume Error: 131,158 mm³
- Performance did not surpass the 2D/Ensemble models, but remained consistent across cases.

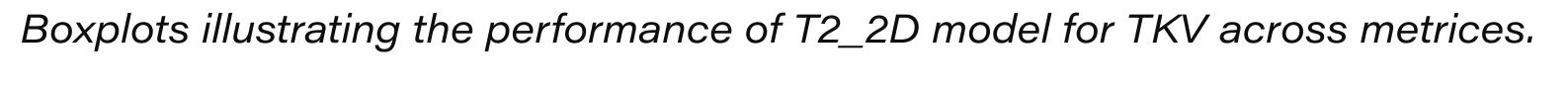
Jaccard Distribution VolumeError Distribution 200000 100000 -

Total Kidney



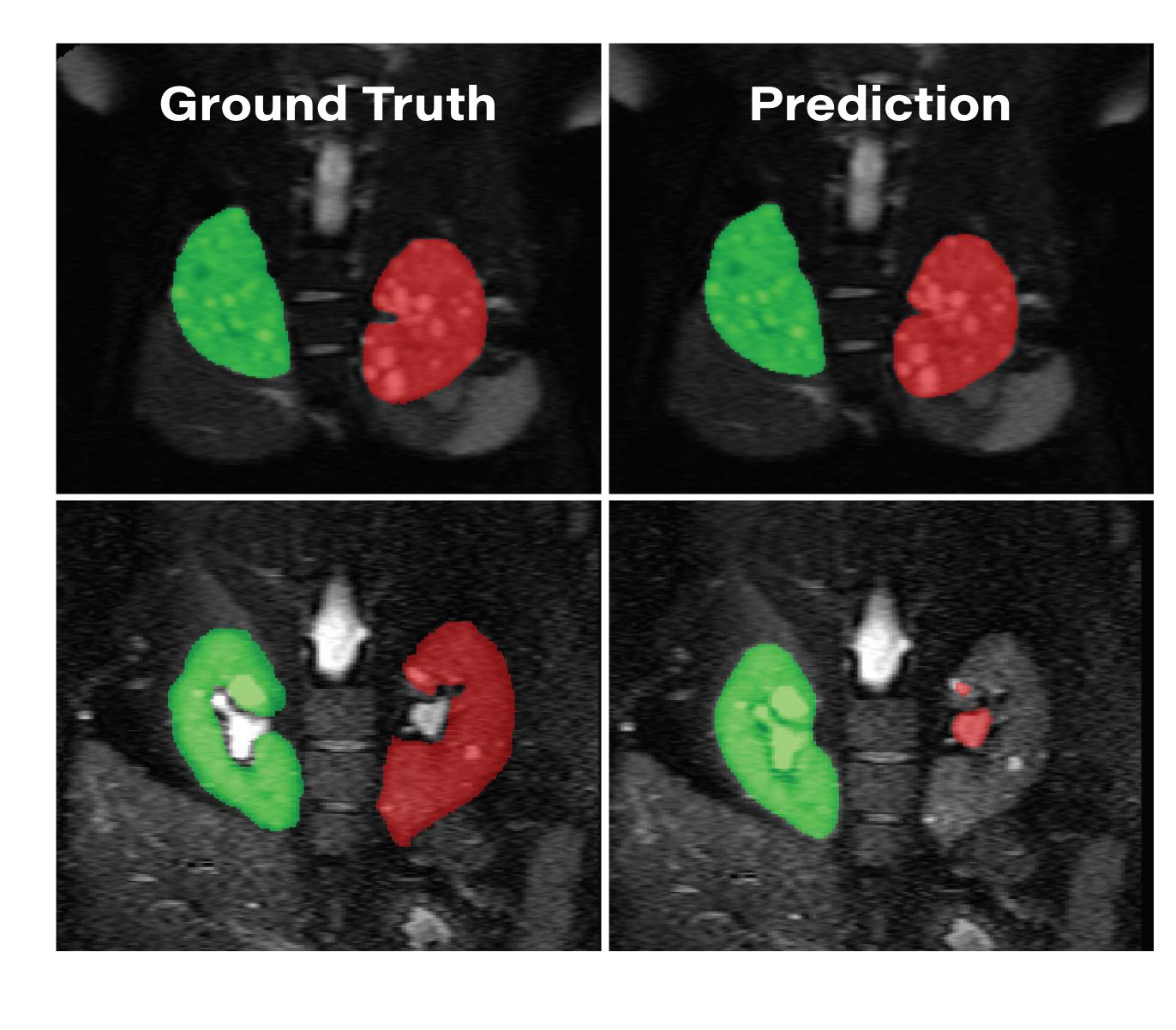






DSC vs Volume Error (Mean per Model) T2_3D_baseline 150000 5 140000 T2_FT •T2_3D ⊕ 130000 -> 120000 T2_Ensemble 110000 T2 2D

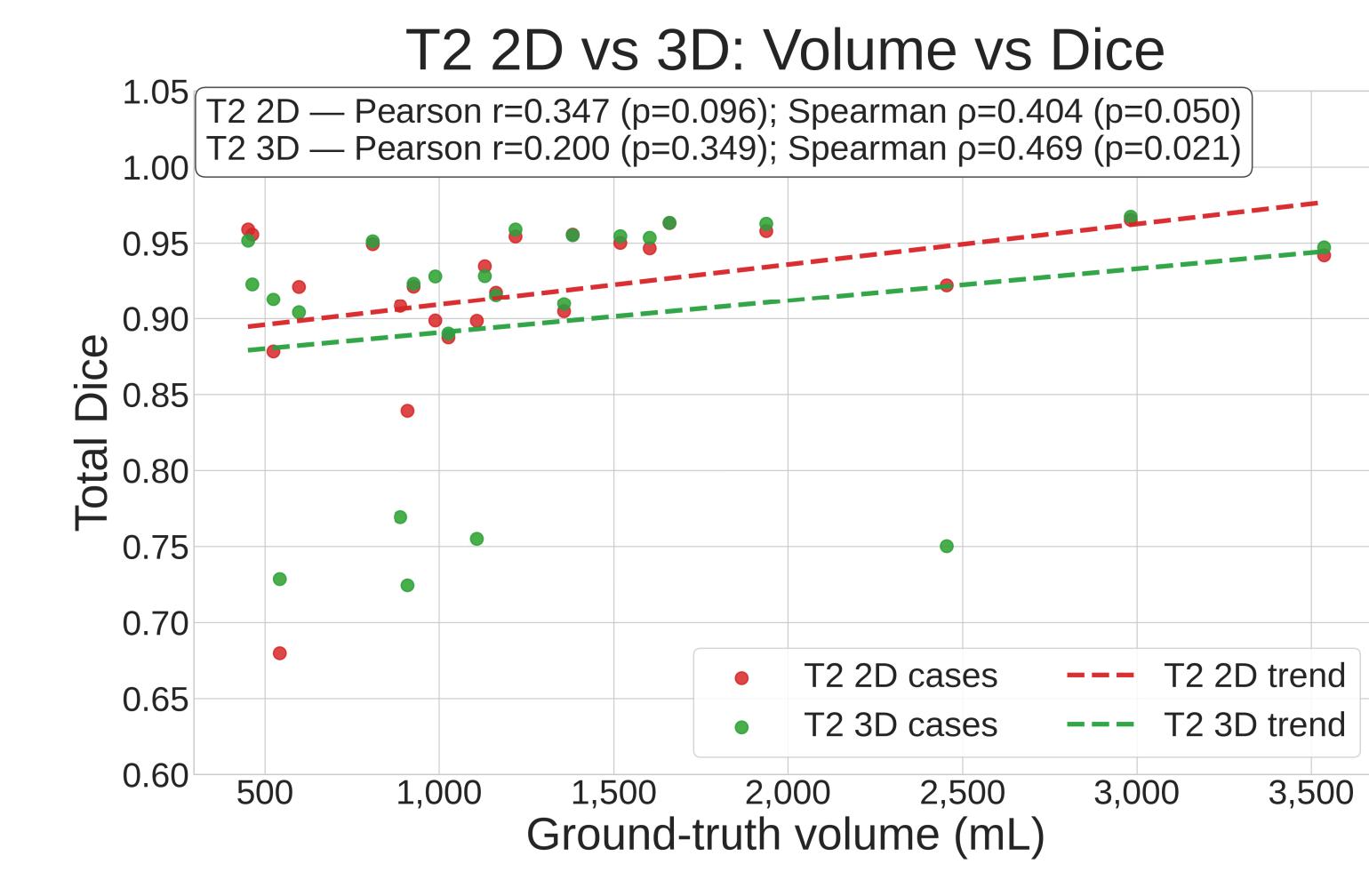
Comparison of models performance showing Dice Similarity Coefficient (DSC) versus Volume Error across different models.

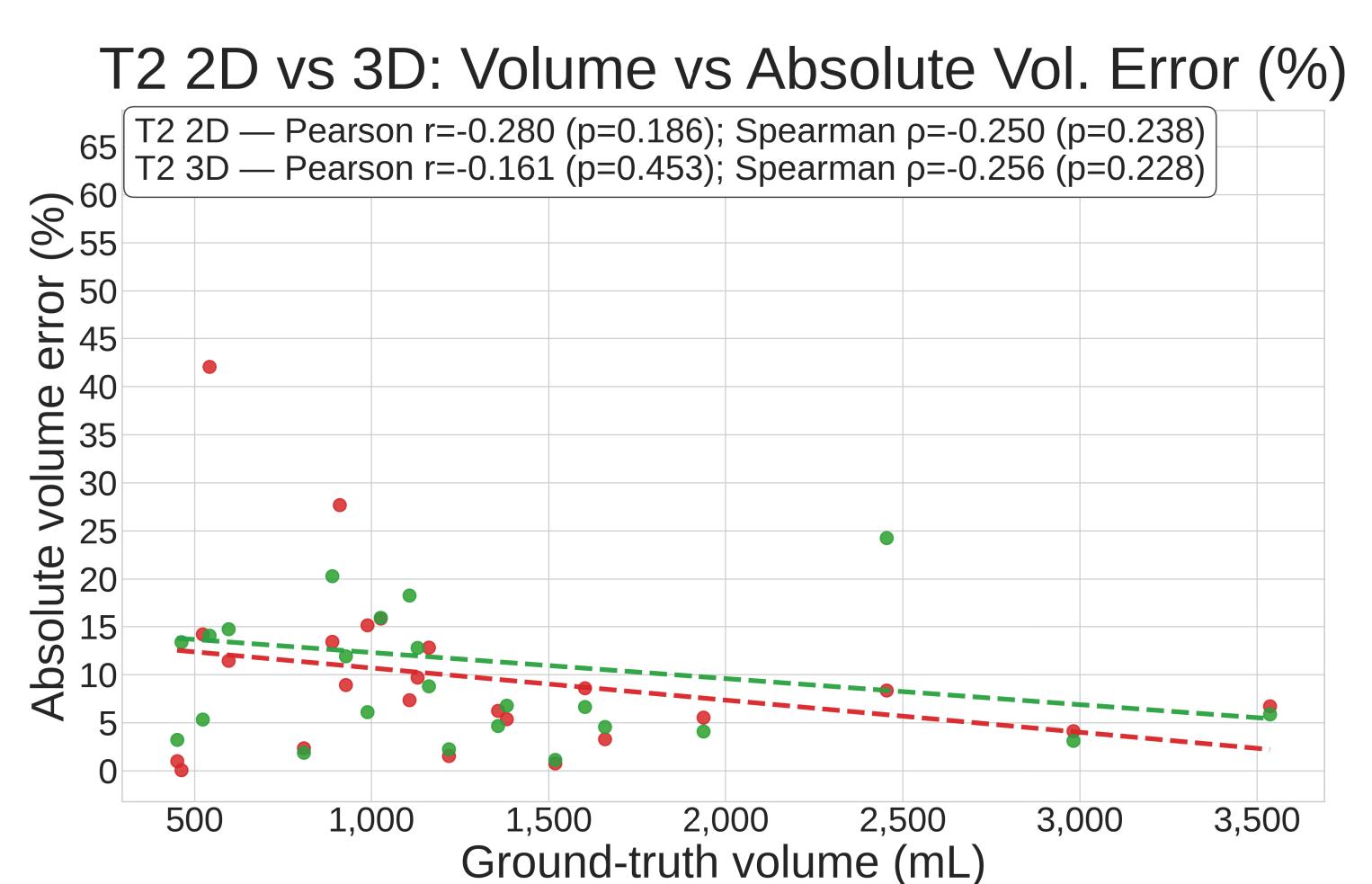


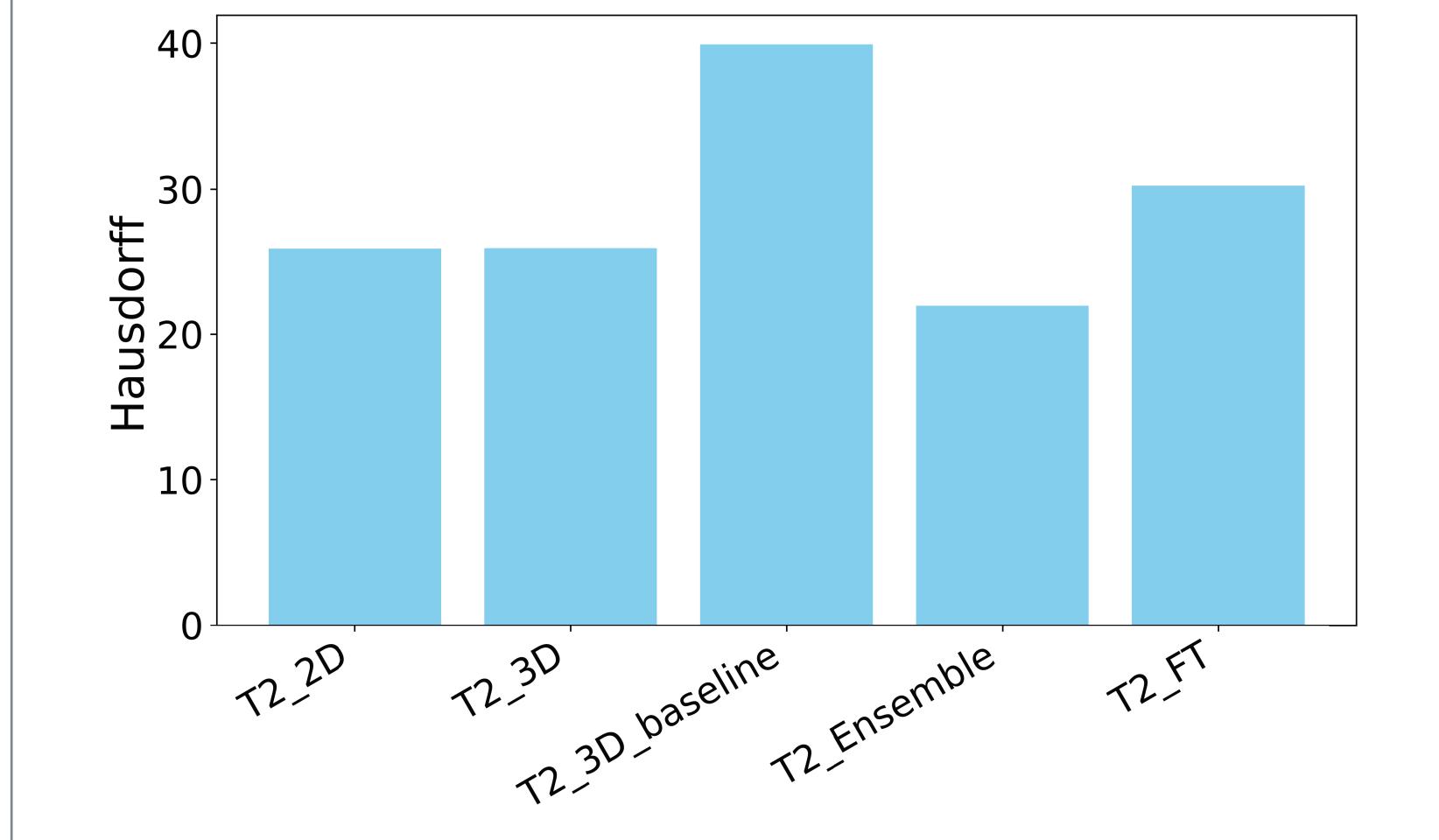
Overlay examples of the best prediction (DSC 0.96) and less accurate prediction (DSC 0.68).

Conclusions

- ► Al-driven models achieved high accuracy in estimating Total Kidney Volume (TKV) in ADPKD patients, with DSC approaching 0.92 and strong volumetric agreement.
- Expanding the dataset from 67 to 92 training images substantially improved performance, particularly with ensemble modeling, reducing both segmentation error and Hausdorff distance.
- Although fine-tuning did not improve accuracy, the results suggest better generalizability across held-out test sets, an important property for clinical translation.
- These findings support the feasibility of deploying AI-based tools for reliable, automated TKV estimation in ADPKD clinical and research settings.
- ► The models do better with larger kidney volumes, evident from the Pearson and Spearman correlation coefficients.







Hausdorff by Model (Mean)

Total Kidney

Comparison of the Hausdorff Distance metric across all evaluated segmentation models to assess boundary accuracy.