

On TotalSegmentator's Performance on Low-Dose CT Images

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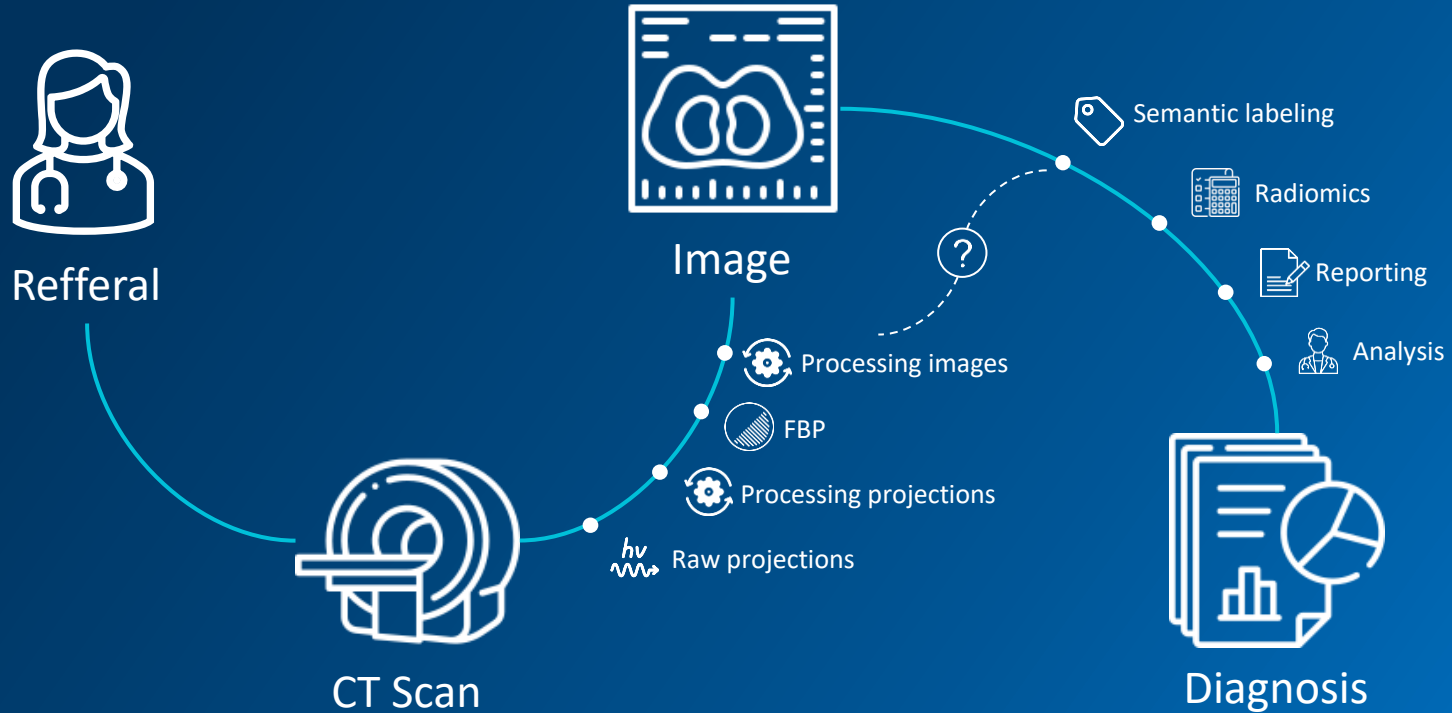
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Presenter:

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How low-dose protocols affect segmentation?

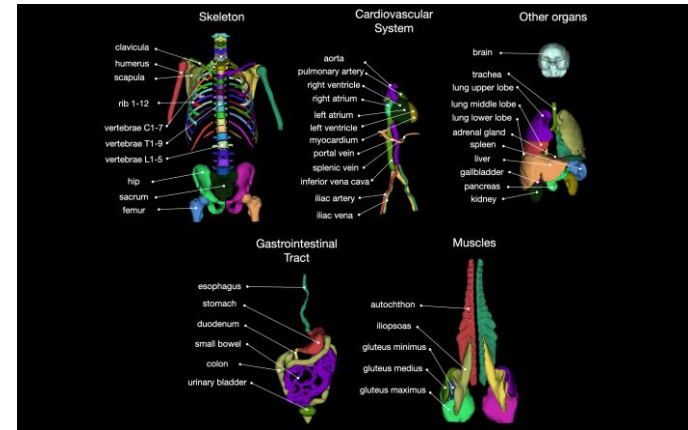
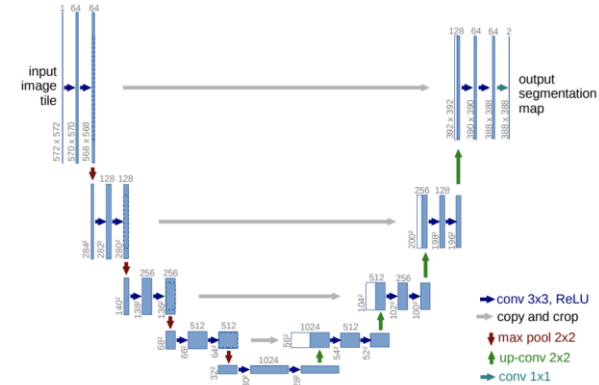


Segmentation Milestones

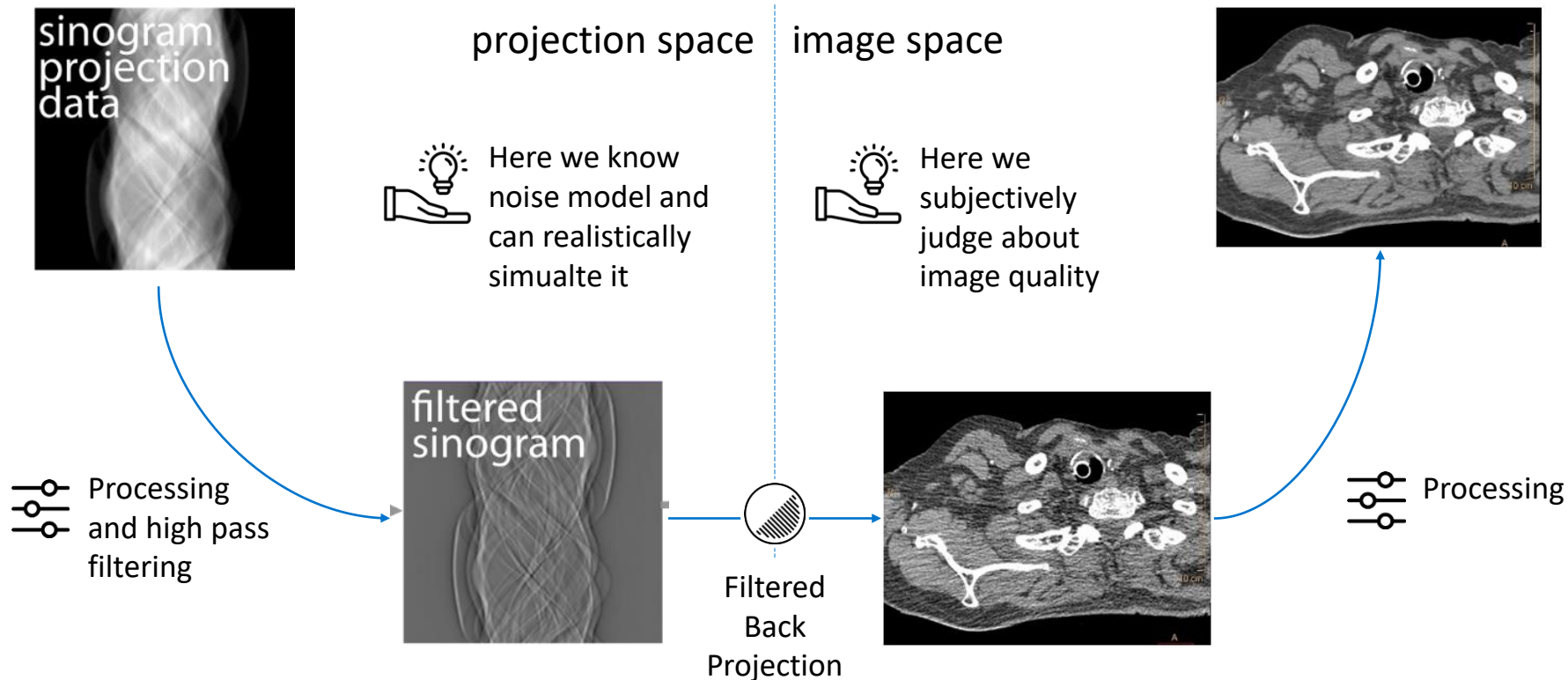
2015 ● U-Net: Convolutional Networks for Biomedical Image Segmentation
 A novel encoder-decoder architecture with skip-connections. Originally applied for segmentation of neuronal structures in electron microscopic stacks.

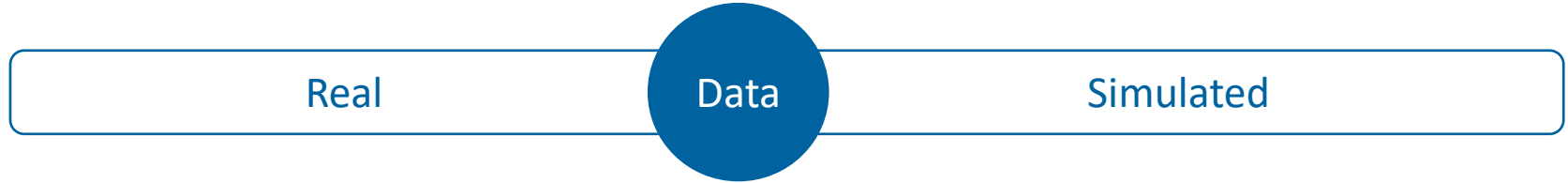
2018 ● nnU-Net: a self-configuring method for deep learning-based biomedical image segmentation
 "nonew-Net". The method automatically configures itself, including pre-processing, network architecture, training and post-processing for any new task

2022 ● TotalSegmentator: robust segmentation of 104 anatomical structures in CT images



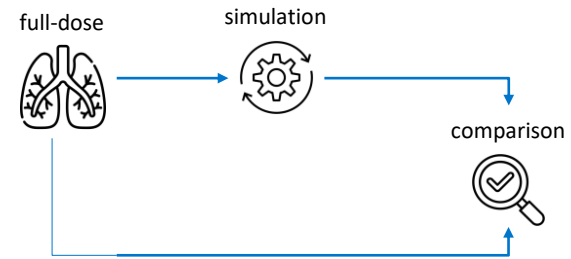
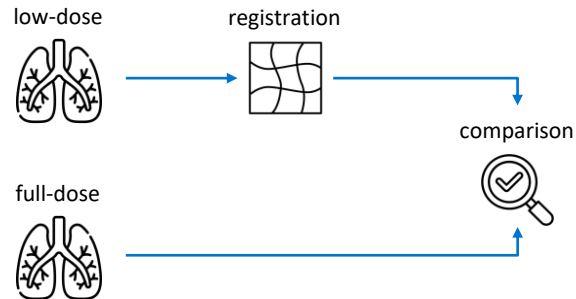
CT Image Formation



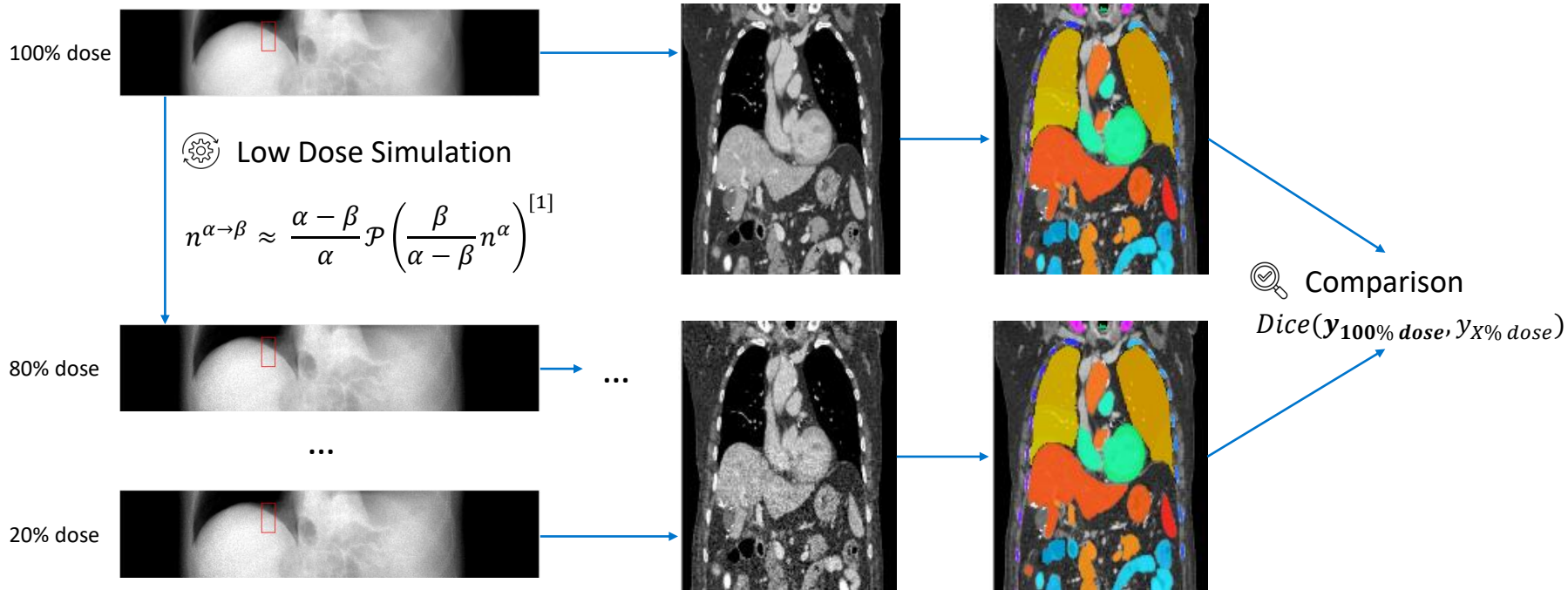


- Low-dose data is taken from real acquisitions
- Leads to unpaired low- and full-dose images
- Low quantity

- Low-dose data is simulated
- Perfectly registered
- **Accurate simulations require raw data and information about a CT system**



Method



[1] Žabić, Stanislav, et al. "A low dose simulation tool for CT systems with energy integrating detectors." *Medical physics* 40.3 (2013): 031102.

Experiment

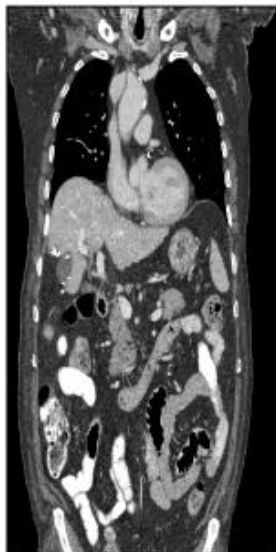
- 42 patients, 99 CT scans
- Raw projections of the abdominal area in helical geometry with 8 cm collimation and pitch factor 1.38
- Reconstructed using 420mm field of view with pixel numbers along X and Y set to 512
- 5 dose levels: 20%, 40%, 60%, 80%, 100%
- TotalSegmentator model consisting of 5 models for different body parts

Table 1. Dice scores calculated between segmentations of full- and low-dose CT images for different dose levels. Since the full-dose segmentation was used as a reference, the scores for the 100% dose level equal 1.0.

Model Name	Dose Level				
	20%	40%	60%	80%	100%
totsegm-all	0.983 ± 0.008	0.990 ± 0.005	0.993 ± 0.003	0.996 ± 0.002	1.000 ± 0.000
totsegm-organs	0.987 ± 0.011	0.992 ± 0.007	0.995 ± 0.004	0.997 ± 0.003	1.000 ± 0.000
totsegm-vertebrae	0.988 ± 0.002	0.993 ± 0.002	0.995 ± 0.001	0.997 ± 0.001	1.000 ± 0.000
totsegm-cardiac	0.982 ± 0.010	0.989 ± 0.006	0.993 ± 0.004	0.996 ± 0.002	1.000 ± 0.000
totsegm-muscles	0.986 ± 0.004	0.992 ± 0.003	0.995 ± 0.002	0.997 ± 0.001	1.000 ± 0.000
totsegm-ribs	0.974 ± 0.006	0.984 ± 0.004	0.990 ± 0.002	0.994 ± 0.002	1.000 ± 0.000

- The median and the median absolute deviation aggregate the results across multiple CT images as some scores are falling into the extreme of the value range
- The absolute decline does not exceed 3%, even at the 20% dose level.

Results



(a) A CT image at 100% dose level.



(b) Segmentation results at 100% dose level



(c) A CT image at 20% dose level.



(d) Segmentation results at 20% dose level



(e) Differences between the segmentation results

Summary & Conclusion

- We analyzed the robustness of the TotalSegmentator model for anatomical segmentation with respect to lower doses in CT imaging.
- By employing raw CT projections, we could accurately simulate low-dose CT data that were intrinsically registered with the original CT images.
- The results reveal surprising robustness of the model even at 20% dose level
- The study is limited due to absence of the ground-truth segmentations
- Future work may include analysis of segmentation models for pathologies

Questions



Please, send your questions to the
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Thank you!