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Weakly supervised 3D ConvLSTMs for Monte-Carlo radiotherapy dose simulations

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Introduction

The **most precise** method to compute clinical dose volumes in radiotherapy is **Monte-Carlo** simulation, but it remains **prohibitively time consuming** if well converged results are required.

Goal: to infer high precision dose volumes from noisy Monte-Carlo dose simulations [1].

Contributions and intuition

Intuition:

ConvLSTM [2] derives meaningful features from the sequential nature of the input to infer the next logical step.

Contributions:

- 1. Investigate 3D ConvLSTM-based architecture
 - → To infer high-precision dose volumes from a sequence of noisy Monte-Carlo generated dose distributions.

2. Noise-to-noise [3] training scheme on VMAT [4] cases

→ Reduces the computation time required to generate the training set.

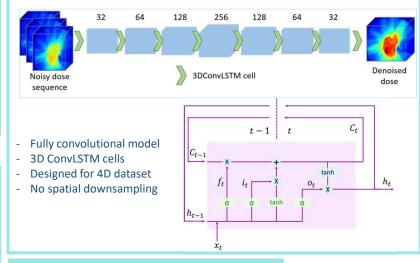
Dataset

- 50 VMAT patients
- Resolution: 2 mm³
- For 1 patient:

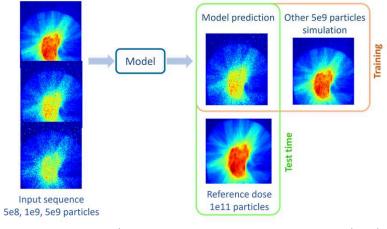
#Particles #Simulations Computation time

5e8	5	5 hours		
1e9	5	10 hours		
5e9	5	50 hours		
1e11	1	4000 hours		
Reference				

Model: Stacked 3D ConvLSTMs



Training scheme : Noise-to-noise



- Noise-to-noise training does not require expensive-to-compute ground-truth.
- Train / validation / test sets: 40 / 5 / 5 patients.
- Patch-based training: 64 x 64 x 64 pixels
- Hyperparameters for AdamW: learning rate=0.001, weight decay=0.01.

- Loss function - weighted sum of SSIM and L1:

 $Loss = SSIM(y_{model}, y_{5e9}) + 20 * L1(y_{model}, y_{5e9})$

Results

- Evaluation with gamma index and clinical metric Gamma Passing Rate [5]
- Dose-to-agreement / tolerance on dose values = 3mm / 3% within 30% 100% of maximum dose.
- Benchmark with BiONet [6] adapted to handle 3D data.
- Gamma index < 1 = 🗸

Denoised dose	Re	eference dose	Gamma inde	x 2 1
Model 3D BiONet	#Weights 178 M	SSIM (%) 91.1 ± 2.2	L1 0.102±0.022	GPR (%) 50.5 ± 4.8
(seq. input) 3D BiONet (5e9 input)	178 M	90.4 ± 3.4	0.072±0.011	71.9 ± 5.1
Stacked 3DConvLSTMs	5 M	86.4 ± 5.2	0.104±0.022	83.8 ± 3.4

Future work

In future studies, we will:

- 1. Assess sensitivity of the model to input noise
- 2. Perform ablation studies to size up the potential of 3D ConvLSTMs

Reference

[1] T. Bai, et al. Deep dose plugin towards real-time monte carlo dose calculation through a deep learning based denoising algorithm. Machine Learning: Science and Technology, 2020.

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[5] D. Low, et al. A technique for the quantitative evaluation of dose distributions. Medical Physics, 1998.

[6]T. Xiang, et al. Bio-net: Learning recurrent bi-directional connections for encoder-decoder architecture. MICCAI 2020.