

# EPID-based *in vivo* transit dosimetry in external beam radiotherapy: prediction of portal dose images using artificial neural networks

Côme Mével Dutertre<sup>ab</sup>, Alexandre Hakimi<sup>a</sup>, Eric Fadel<sup>a</sup>, François Smekens<sup>a</sup>, Xavier Franceries<sup>b</sup>, François Husson<sup>a</sup>

<sup>a</sup>DOSIsoft Physics R&D, Cachan, France

<sup>b</sup>INSERM Toulouse NeuroImaging Center, Toulouse, France

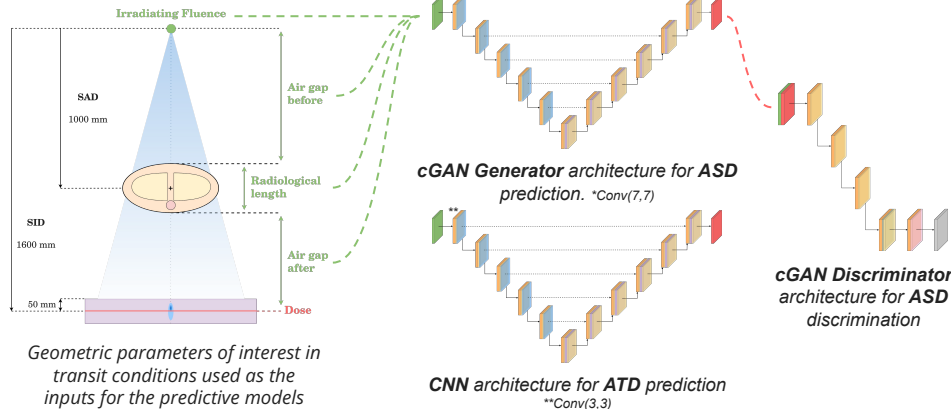
## Introduction

EPID-based *in vivo* transit dosimetry using the forward technique requires the accurate calculation of the predicted (reference) image to be compared in terms of absolute dose with the acquired image.

The **Collapsed Cone Convolution (CCC)** algorithm is a fast method for accurate dose predictions suitable to transit dosimetry, and allows a distinction between the dose generating phenomena. The **Detector Total Dose (DTD)** at the computation plane is the sum of the **Attenuator Transmitted Dose (ATD)** from direct photons beam transmission and the **Attenuator Scattered Dose (ASD)** from beam interaction in the attenuator.

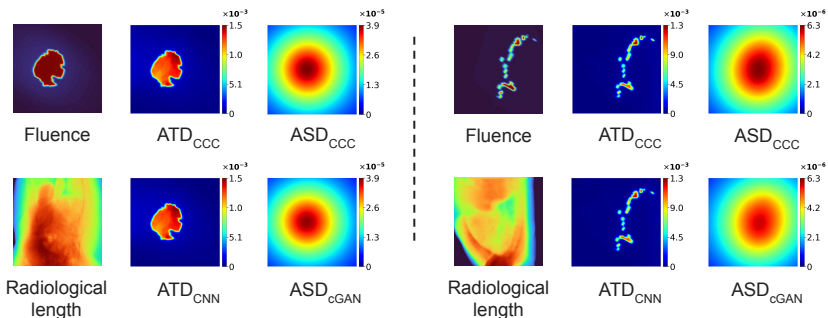
This work introduces a **Machine Learning** workflow based on CCC ground-truths, using a **Convolutional Neural Network (CNN)** for ATD calculation and a **conditional Generative Adversarial Network (cGAN)** for ASD computation.

## Machine Learning approach



- 17 unique CTs from various tumoral sites
    - 15 for training
    - 2 for testing
  - 16 unique VMAT RTPlans
    - 14 for training
    - 2 for testing
  - Irradiation segments
  - Gantry angles variations
  - Reference: CCC (ThinkQA DOSIsoft)
  - Elekta Synergy 6 MV
- 9263 training cases  
→ 2573 testing cases

## Results



	ATD <sub>CNN</sub> VS ATD <sub>CCC</sub>	ASD <sub>cGAN</sub> VS ASD <sub>CCC</sub>	ATD <sub>CNN</sub> VS DTD <sub>CCC</sub>	DTD <sub>CNN + cGAN</sub> VS DTD <sub>CCC</sub>
Average GPR (%)	95.89	54.50	91.98	95.90
Median GPR (%)	98.95	58.68	97.21	98.96
GPR > 95% (%)	78.43	14.57	61.17	78.35
GPR > 90% (%)	87.01	21.26	74.19	87.21

GPR (2%, 2mm, 10% threshold) analysis is performed on the test dataset in terms of separate dose components and total dose

## Conclusion

- The full ML model allows:
- CCC equivalent predictions in terms of total dose at the EPID level
  - Lower computation time for identical CPU/GPU (~100x less)
  - Generalization across various tumoral sites
  - Restitution of physics phenomena (transmission, scattering)