

Workshop on Innovative Delivery Systems in Particle Therapy

23-25 February 2017 Molecular Biotechnology Center

Online dose calculation using GPUs

Simona Giordanengo Istituto Nazionale di Fisica Nucleare Torino, February 25, 2017





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Online dose calculation using GPUs integrated in the CNAO DDS

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ONLINE → DURING treatment so integrated with the Dose Delivery Systems (DDSs)

When can be computed the dose?

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For synchrotron online means during the inter-spill time → Synchronous with spills

ONLINE → DURING treatment so integrated with the Dose Delivery Systems (DDSs)

When can be computed the dose?



For synchrotron online means during the inter-spill time → Synchronous with spills For cyclotron ...? Could be in parallel with the delivery but not synchronous

Is this feasible? Yes, using GPUs

GPU-based

- pencil beam algorithms (Da Silva 2015) and
- Proton MC packages (Yepes 2009, Jia 2012, Renaud 2015, Wan Chan Tseung 2015,) have recently been developed.



... in place of CPU

CPU MULTIPLE CORES



GPU THOUSANDS OF CORES

The computation time was reduced (up to a few minutes for MC-based dose simulation) via both **hardware accelerations** and the use of **simplified physics models**

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PRECISION vs computation TIME is the relevant TRADE-OFF

Fast MC dose calculations shows great promise for accurate inverse plan optimization as well as for pre and post treatment independent dose calculation for dose verification in place of measurements

BUT are NOT yet SUITABLE for ONLINE applications.

Sub-second forward dose calculation: Is it feasible? Yes for Pencil-Beam algorithms

Sub-second pencil beam dose calculation on GPU for adaptive proton therapy

Joakim Da Silva et al. University of Cambridge, UK **Phys. Med. Biol. 60 (2015)** 4777



- Tesla K40 GPU (2880 cores @ 845 MHz)
- Skull base test case
 - 2 oblique beam directions
 - 38 + 45 energy layers, total 6776 spots
 - 2x2x2 mm³ dose resolution, 1x1x1 mm³ ray resolution
- Total calculation time 0.22 seconds
- Individual energy layers: 2.2-6.4 ms

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Physics in Medicine & Drawy PMB 2015

VALIDATED vs Syngo and Fluka



- 2%/2 mm γ-index compared to Fluka
- Passing rate: 96.7% (voxels receiving >10% of max dose)
- Corresponding value for Syngo: 96.8%

The INFN and University of Torino experience on DDS and TPS

It started in 1993 developing beam monitors and MC dose simulations for hadrontherapy

Leading to two important products

The CNAO DDS in collaboration with CNAO Foundation, which is now a CE Medical Device



The PlanKIT TPS in collaboration with IBA



Giordanengo et al. Med. Phys. 42 (2015)

Russo et al. Med. Phys. 61 (2016)

The Dose Engine Kernel (DEK) TPS

The INFN-IBA Beam Model allows computing the 3D effect of **an ion field** incident on a **water-like material**. This model is implemented in a Dose Engine Kernel (DEK) to estimate the outcome of a therapeutic ion irradiation delivered with the Pencil Beam Scanning (PBS) technique.

Biological Dose RBE **Dose-averaged LET** 1.4 1.2 Protons 125 100 **Carbon** ions

In the following DEK doses will refer to CPU-based doses by DEK TPS

S. Giordanengo - INFN Torino - Online dose calculation using GPUs

PlanKIT OUTCOMES:

- Physical Dose
- RBE
- Biological Dose
- LET
- Dose-averaged LET
- **α**
- [
- Survival

For different ions!

GPU-based DEK Forward Planning



C++ DEK for single core CPU

CPU single core

Not optimize to be fast but to be versatile in order to easily be adapted to different ions, facilities and computational requirements

GPU-based DEK Forward Planning

In the Framework of the INFN RIDOS project (2014-2015) A Fast Forward Planning implementation has been done porting on CUDA the INFN-IBA TPS libraries (SW)



Why GPUs? (I) Thousands narrow beams over thousands voxels

1000 - 50000512 x 512 x 250 voxels pencil beams for field 1 thread for each voxel 155 150 145 (MeV/u) Pencil beam thread for each PB 130 125 120 30 Energy 20 30 20 -10 -10 -20 -30 Y (mm) -30 \times (mm)

Thread = Sequence of Single Instruction on Multiple Data (SIMD) operations

Algorithms for PBS dose: (I) Raytracing

 $Beam \rightarrow ray$

Raytracing \rightarrow

Find intercepts + sorting intercepts + path length evaluation



Voxels hit by a ray (pencil beam direction)

Raytracing : Find intercepts with CT voxels

GPU code: check the intercept between the ray and each plan along the 3 voxels direction (stream compaction algorithm) Max N intercepts: $(N_x+N_y+N_z) \times N_{ray}$ N = CT voxels ~ (512+512+256)

 N_{rav} (Min ÷ Max) = 1 ÷ 5000000

N intercepts range : $10^3 \div 5 \times 10^9$



Raytracing(II-III): Sorting the intercepts and measuring the ray path length (L_i)



Bitonic Sort : the intercepts are sorted in function of the distance from the CT entrance



Why GPUs? (II) Interpolations exploit the use of GPU-Texture memory



Why GPUs? (II) Interpolations exploit the use of GPU-Texture memory



Large cut-off \rightarrow beam halo is considered \rightarrow better accuracy Large cut-off \rightarrow more voxels \rightarrow high computation time for standard C++ codes

2nd Texture Interpolation: dose evaluation from pre-computed LUTs



GPU-DEK time performances

Entire Per spill				Computing Grid = 170x170x125; N of Rays = 1248; N of slice = 39; CPU: XeonE5-1620v2-3.7-4C-16GB GPU: Nvidia TESLA-K20c		
Radial cut- off (mm)	Number of voxels	RIDOS F-	FP time (s)	PlanKIT FP time (s)	Gain	
20	44678	1.35	0.035	157	116	
40	178702	2.33	0.059	636	273	
50	279243	2.76	0.071	997	361	
80	714861	4.68	0.120	2534	541	

GPU-DEK vs CPU-DEK for clinical patient

Dose (GPU) [Gy] Differences [Gy] Dose (DEK) [Gy] 1.8 1.8 10-4 -1.8 3 20 1.6 1.6 1.6 2 ^{1.4} 1.4 1.4 **Protons** 1 1.2 1.2 0 1.0 **Physical Dose** 1.0 -1 100 0.8 0.8 120 -2 0.6 0.6 0.4 -3 0.4 0.2 0.2 -4 20 40 60 80 100 120 20 40 60 80 100 120 40 60 80 100 120 20 2 10-4 4.5 4.5 4.0 4.0 **Carbon Ions** 3.5 3.5 3.0 3.0 **Biological Dose** 2.5 2.5 <u>∽</u> 30 1 2.0 2.0 1.5 1.5 Com Grid = 170x170x127; N of Rays = 10761; 1.0 1.0 N of slice = 51;0.5 0.5 N voxels= 7292; radial cut-off=10 mm. n 0 100 40 60 80 40 60 80 100 60 voxel × voxel x voxel x

GPU-DEK vs Syngo TPS for cube in water



Protons – SOBP21cm Cube 3x3x3 cm³

0.4 **GPU-DEK vs SYNGO CNAO** patient – dose to water 0.3 proton – Small H&N tumor TPS Lateral profiles value 0.2 DEK RIDOS SYNGO 0.8 0.1 0.6 -0.0 -200 -100 ò z(mm) TPS value DEK RIDOS SYNGO TPS DEK RIDOS SYNGO 0.2 -0.0 200 -100 100 -200 Ó z(mm) 0.00 -100 -200 Ó 100 200

z(mm)

GPU-DEK vs PinPoint measurements



What we can do with so fast dose calculation?

Upgrade the new generation of DDS



New dose Planning and Delivery System

able to check ON-LINE the uncertainties in the delivered DOSE DISTRIBUTIONS due to measured patient and beam deviations from planned conditions.



...Goal of the INFN RIDOS project

Grant for Young Researcher 2014/2015

RIDOS HW and interfaces



- *Text* file with nominal patient and treatment data (CT, nominal beams characteristics)
- Spill data binary file (Spill_N.ridos 1-100 kB/spill)

RIDOS HW and interfaces



RIDOS HW and interfaces

RIDOS GUI in Matlab

RIDOS "4D"

10 time [s]

0 -

Ó

5

[90,100)

Dose accumulation with Deformable Image Registration (DIR)

Doses accumulated spill by spill in different respiratory phases

Preliminary results of a GPU-based Dose accumulation in different respiratory phase

NVIDIA Quadro K2000

CUDA Cores: 384 Memoria Globale: 2 GB

NVIDIA Tesla K20

Cuda Cores: 2496 Memoria Globale: 5 GB

Tempi interpolazione	CPU	GPU K2000	GPU K20	Gain
DVF interp.	0,15 s	0,011 s	0,006 s	~14 / ~25
Dose interp.	5,1 s	0,18 s	0.08 s	~30 / ~63

Future and perspectives: Adaptive radiotherapy with PBS (?)

Adaptive radiotherapy consists in performing dose feedbacks

INTER-fraction dose feedback will be soon available for proton beams with commercial TPS and also with Fast MC

INTRA-fraction dose feedback is possible ...if you have access to DDS HW and FIRMWARE and you have a sub-seconds dose calculation

BUT: *on-line patient imaging* is mandatory for significant clinical improvement

Conclusions

GPUs allow online accurate PB dose calculation

A FAST GPU-based Dose Engine Kernel for different ions has been integrated in the CNAO DDS and is under test.

The online doses can be calculated on different CTs exploiting the use of 4D-CT to improve treatments of moving targets.

Vincenzo Monaco Anna Vignati Andrea Attili Felix Mas Milian Marco Donetti Mario Ciocca Silvia Molinelli Roberto Cirio Flavio Marchetto Germano Russo Roberto Sacchi Lorenzo Manganaro Giorgio Magro Mohammad Varasteh Anvar

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