#### Introduction and overview of MMCTP

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#### Layout



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- Current use at McGill
- McGill RT Format

#### 3 Research Software

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#### Why use MC?

What is the effect of more accurate MC dose distributions on patient clinical outcome?

- Need to investigate the correlation of MC calculated dose distributions with clinical outcome (tumor control and normal tissue toxicity)
  - 2003 De Jaeger *et al* demonstrated that the calculated incidence of radiation pneumonitis correlated better with observed incidence when using more accurate dose calculations
- 2006 Siebers *et al* investigated the use of more accurate dose calculations in IMRT optimization (DPEs and OCEs)

What do you gain from MC?

Clinical TPS are a black box

- Research MC codes are fully customizable
- Model detectors
- Model high Z materials in patient
- Model small fields
- Investigate beam vs patient transport differences

#### Monte Carlo Treatment Planning at McGill





#### CADPlan MCTP at McGill



#### Why build MMCTP

- Electron and photon Monte Carlo treatment planning system
- Establish large-scale retrospective studies
- Dose comparison tools
- Combine research projects under one platform

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#### MMCTP

#### "BEAMnrc without MMCTP is just uncivilized"

Ellis Mitrou



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#### What is MMCTP?

A full feature treatment planning system that uses various Monte Carlo calculation systems as dose calculation engines which run remotely on one or many multiprocessor computers.



http://www.medphys.mcgill.ca/~mmctp/MMCTP/Welcome.html

Alexander et al, Phys. Med. Biol., 2007



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## **MMCTP** Features

- Import DICOM RT, RTOG, CADPlan CART
- Export RTOG, DICOM RT Dose
- S Visualization options 2D, 3D, images, structures, doses
- Edit treatment planning settings
- Monte Carlo links to BEAMnrc, DOSXYZnrc, Cutout, XVMC
- Ose analysis tools
- Ø Available for MAC, Linux, Windows platforms

- Dose calculations are fully automated by logic scripts which determine when and where to submit MC simulations
  - Local computer or departmental cluster
  - Cloud or remote clusters (West grid, CLUMEQ)

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- Beam commissioning tools for profiles, PDDs, output tables

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# MMCTP Projects at McGill

- Modeling lung toxicity outcome with MC
- Ø Medulloblastoma MC recalculation
- Tomotherapy DQA and patient recalculation
- IMRT QA verification
- Energy and intensity modulated electron therapy (MERT,IMET)
- Olinical MMCTP station
  - MapCHECK<sup>TM</sup>
  - Head and neck IMRT recalculations
  - SBRT lung MC recalculation

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#### Head and Neck MC recalculation



AAA (dashed lines) and MC (solid lines). The PTV60, shown in light blue color, is surrounded by regions of high and low density materials. The MC 63 Gy isodose line deviates from the AAA isodose line in the top right conner. The deviation is believed to be caused by the high density mandible bone adjacent to the PTV60 structure.

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#### Lung SBRT MC recalculation



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# Lung SBRT MC recalculation on patient 15 with bullous lung disease





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## McGill RT Format



- Minimize redundant information
- Minimize number of files edited for save changes
- Use binary or text files when appropriate

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# Image files

- Image files are named sequentially with the file type as \*.img
- Files are binary, containing a slice Z position and the pixel data
- Image properties text file *RT.dir* stores header info

#### Table: RT.dir file

Patient Name	:= (string)
Patient ID	:= (integer)
IMAGE MODALITY	:= (string)
IMAGE TYPE	:= (string)
SCAN TYPE	:= (string)
CT offset	:= (integer)
Grid unit width	:= (double)
Grid unit height	:= (double)
Number representation	:= (string)
Bytes per pixel	:= (integer)
Size of Dimensions 1	:= (integer)
Size of Dimensions 2	:= (integer)
X offset	:= (integer)
Y offset	:= (integer)
CT-AIR	:= (integer)
CT-WATER	:= (integer)
Slice thickness	:= (cm)
Date	:= (string)
WINDOW LEVEL	:= (integer)
WINDOW WIDTH	:= (integer)

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#### McGill RT coordinate system



- 1<sup>st</sup> (X, Y) coordinate defines the center of the top right corner pixel
- Z increase into the page
- 1<sup>st</sup> coordinate Z is the lowest slice

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## Structure files

- Sequence of three-dimensional coordinates
- Files are text, type \*.*struct*
- Coordinates are grouped together in Z planes, which coincide with planes on the CT image

STRUCTURE NAME NUMBER REPRESENTATION STRUCTURE FORMAT STRUCTURE COLOUR RGB NUMBER OF SCANS

"NUMBER OF LEVELS" 3 "SCAN NUMBER" 1 "# OF SEGMENTS" 0 "SCAN NUMBER" 2 "# OF POINTS" 4 0, 1.459, 20.86305 -0.56, 1.387, 20.86305 -0.776, 1.387, 20.86305 -0.848, 1.315, 20.86305 "SCAN NUMBER" 3 "# OF SEGMENTS" 0

#### Table: Structure file

:= PTV2 := CHARACTER := SCAN-BASED := 252/18/41 := 3

(total number of scans) (=1 for first scan, etc) (number of segments in this level/scan)

(number of points in first segment) (X, Y, Z coordinates of each point)



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# Beam files

- Defines an external beam
- Files are text, type \*.Beam
- Two subfiles
  - MLC
  - Collimator

BEAM # TREATMENT UNIT BEAM MODALITY BEAM ENERGY BEAM APPLICATOR BEAM DESCRIPTION RX DOSE PER TX (GY) MUs NUMBER OF TX FRACTION GROUP ID BEAM TYPE COLLIMATOR TYPE APERTURE TYPE WEDGE ANGLE WEDGE ORIENTATION WEDGE DYNAMIC COLLIMATOR ANGLE GANTRY ANGLE COUCH ANGLE NOMINAL ISOCENTER DIST NUMBER REPRESENTATION PLAN ID OF ORIGIN

#### Table: Beam file

- := (beam number in plan)
- := (treatment unit name ex CL21EXA)
- := (electron, photon)
- := (energy ex 18 MV)
- := (for electrons only)
- := (text description of beam)
- := (fraction dose)
- := (number of MUs)
- := (number of fractions)
- := (id to group beams of common fraction
- := (static or arc)
- := (symmetric or asymmetric)
- := (block or MLC)
- := (wedge angle in degrees ex 15)
- := (wedge orientation ex in,out,left,right)
- := (dynamic wedge boolean ex true/false)
- := (angle in degrees)
- := (angle in degrees)
- := (angle in degrees)
- := (isocentre distance in cm)

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- := (character)
- ψЩψ := (plan ID of beam for grouping beams)

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## MLC files

- Defines an MLC pattern for one beam
- Files are text, type \*.MLC
- index(i) index(i-1)is the probability of field *i*

#### Table: MLC file

:= STATIC := 1 := := Varian 120M := 0 := 0
:= 1
:= 100.
:=
:=
:=
:= 0.
:= 0.
:= 0.
:= 0.
:= 0.
:= 1.089
:= 1.785
:= 2.164
:= 2.384
:= 2.487

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#### Beam angles



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## Beam jaws



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#### Dose Distribution files

- Sequence of 2D dose planes *x*, *y*
- Files are binary, type \*.Dose
- 100 byte header, 4 byte single per dose point
- Additional sequence of uncertainty values

#### Table: Dose Distribution file

Coordinate X 1st point := (cm) Coordinate Y 1st point := (cm) Coordinate Z 1st point := (cm)Size of Dimension 1 := (# horizontal points) Size of Dimension 2 := (# vertical points) Size of Dimension 3 := (# planes) Horizontal Grid := (cm > 0)Vertical Grid := (cm > 0)Depth Grid := (cm > 0)Dmax := (Gv)Dmin := (Gy)Dose Units := (string)

# MC control files

- These file keep a record of what has been done and what needs to be done
- per plan, 1 BEAMnrc file and a DOSXYZnrc and XVMC file per patient model
- Each file store the location of the simulation, progress, CPU time, normalization values etc...
- All input files are stored within the plan file this keeps a local record of the simulation
- Input files are named by the patient ID # followed by plan # and beam #, ex: 34392\_p01b3.egsinp

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## **MMCTP** License

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- Use this Software at your own risk
- The Software is in constant development and there is a non-zero probability that the user will encounter errors.

## **MMCTP** Sites

• Home page http:

//www.medphys.mcgill.ca/~mmctp/MMCTP/Welcome.html

- User google group page http://groups.google.com/group/mmctp
- MMCTP distribution page http: //www.medphys.mcgill.ca/~mmctp/MMCTP/MMCTPDis