

Installing MMCTP

Andrew Alexander

Medical Physics Unit
McGill University

June 6, 2011



Layout

1 Install files

2 Configurations

- MMCTP Preferences
- MMCTP Shells
- CT to Density



Install MMCTP

- Future updates from http:
`//www.medphys.mcgill.ca/~mmctp/MMCTP/MMCTPDIs`
- Latest version on USB key
- Move MMCTP folder to local drive



Files

	Mac	Windows	Linux
Package	mpkg	msi	no
App Dir	Applications	Program Files	/usr/lib/
MMCTP Setting	user:Library: Application Support:	user\AppData\Roaming\	/home/user
Quesa	Library: Frameworks: Quesa.framework	\Windows\ Quesa.dll	no

- App Directory : MMCTP program file, Configuration folder, DICOM library text file
- MMCTP settings directory : Stores MMCTP user files
- Quesa (Open Source, 3D graphics library) : Required for 3D viewer



Windows and Linux

- ① Windows users move plink.exe and pscp.exe from application configurations folder to C:\
- ② Linux users move linux MMCTP folder to /usr/lib



User files and folders

- ① Specify a MMCTP home directory on your computer (USB MMCTPData folder), this directory houses:
 - McGill-RT directory
 - BEAMnrc directory (input files and dosxyznrc_material.txt file)
 - Create a Commissioning directory
- ② Move STT files to MMCTP settings directory



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MMCTP Settings and files

These files are unique to each user and are edited within MMCTP

- ① Login file
- ② Preference file
- ③ MLC file
- ④ Linac file
- ⑤ MC control file



MMCTP Preferences

- Specify folder locations
 - ① McGill RT folder (stores imported McGill RT patients)
 - ② DICOM folder (dynamic)
 - ③ BEAMnrc folder (stores BEAMnrc and DOSXYZnrc input files, and preferences)
 - ④ Commissioning folder (stores beam data, profiles, PDDs, output tables)
 - ⑤ McGill Cutout (for electron cutouts)
- Export options
- DVH calculation options



MMCTP Preferences

MMCTP Configurations

Beam Configurations MLC CT to Density Monte Carlo Settings Shell Refresh Shell Run Shell Download Shell Login **Preferences** Dose Stats

McGill RT Folder	/Users/andrew/Documents/MMCTP\Programming/McGillRT	<input type="button" value="Change"/>
BEAMnrc Input Folder	/Users/andrew/Dropbox/MMCTP/BEAMnrc	<input type="button" value="Change"/>
DICOM Folder	/Users/andrew/Library/Application\ Support/MMCTP/MMCTP\Settings	<input type="button" value="Change"/>
RTOC Folder	/Users/andrew/Dropbox/MMCTP/Main	<input type="button" value="Change"/>
CADPLAN Folder	/Users/andrew/Dropbox/MMCTP/Main	<input type="button" value="Change"/>
Commissioning Data folder	/Users/andrew/Dropbox/MMCTP/Commission\Data	<input type="button" value="Change"/>
VMC Folder	/Users/andrew/Dropbox/MMCTP/VMC	<input type="button" value="Change"/>
McGill Cutout folder	/Users/andrew/Dropbox/MMCTP/Cutout	<input type="button" value="Change"/>

Export Dose Plane

Interpolate Export in DICOM format
 Do not Interpolate Export in text format

DVH Calculation

Use Graphics Use Graphics and Is Within Routine Use Only Is Within Routine



MMCTP Shells

MMCTP Configurations

Beam Configurations MLC CT to Density Monte Carlo Settings Shell Refresh Shell Run Shell Download **Shell Login** Preferences Dose Stats

Shell **MPU-New** Delete Shell

Properties

Title: MPU-New Machine IP: 192.168.0.202 Mac Linux PC
User: andrew Password: ********* Online

Line Feed Prompt: andrew\$ Bash: **Connection Test**

List Files:

Command: ls -l Column number for file size: 5 Column number for date: 6 0
Column number for file name: 8 Column number for time: 7

Tail:

Command tail After file name

Cluster:

File transfer protocol

FTP FTP line feed prompt: ftp> SCP **Connection Test**

Batch: keg Queue: Number of Queues: 0 Name:

MC:

Max # of running jobs 4 BEAMnrc release

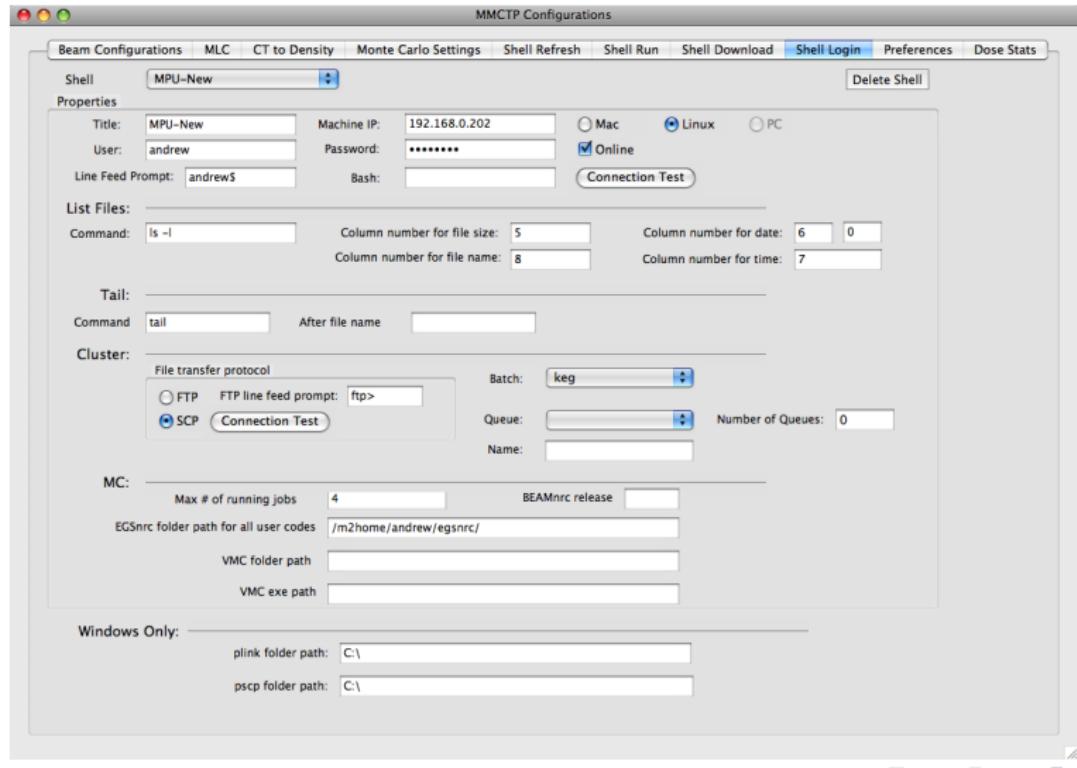
ECSnrc folder path for all user codes /m2home/andrew/egsnrc/

VMC folder path

VMC exe path

Windows Only:

plink folder path: C:\ pscp folder path: C:\



CT to Density Curves

MMCTP Configurations

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Model Eclipse-het Delete Model

CT Model

Name	Eclipse-het
Pegs file	700icru
Number of Materials	3

Material	HU Low	HU High	Density Low	Density High
H2O700ICRU	-1000.	0.	0.001	1.
H2O700ICRU	0.	6000.	1.	3.9
H2O700ICRU	6000.	10000.	3.9	3.9

