

DESIGN AND SIMULATION OF A PASSIVE-SCATTERING NOZZLE IN PROTON  
BEAM RADIOTHERAPY

A Thesis

by

FADA GUAN

Submitted to the Office of Graduate Studies of  
Texas A&M University  
in partial fulfillment of the requirements for the degree of

MASTER OF SCIENCE

December 2009

Major Subject: Health Physics

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Approved by:

Chair of Committee,	John W. Poston, Sr.
Committee Members,	Leslie A. Braby
	Michael A. Walker
Head of Department,	Raymond J. Juzaitis

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## ABSTRACT

Design and Simulation of a Passive-Scattering Nozzle in Proton Beam Radiotherapy.

(December 2009)

Fada Guan, B.E., Tsinghua University, China

Chair of Advisory Committee: Dr. John W. Poston, Sr.

Proton beam radiotherapy is an emerging treatment tool for cancer. Its basic principle is to use a high-energy proton beam to deposit energy in a tumor to kill the cancer cells while sparing the surrounding healthy tissues. The therapeutic proton beam can be either a broad beam or a narrow beam. In this research, we mainly focused on the design and simulation of the broad beam produced by a passive double-scattering system in a treatment nozzle.

The NEU codes package is a specialized design tool for a passive double-scattering system in proton beam radiotherapy. MCNPX is a general-purpose Monte Carlo radiation transport code. In this research, we used the NEU codes package to design a passive double-scattering system, and we used MCNPX to simulate the transport of protons in the nozzle and a water phantom. We used “mcnp\_pstudy” script to create successive input files for different steps in a range modulation wheel for SOBPs to overcome the difficulty that MCNPX cannot be used to simulate dynamic geometries. We used “merge\_metal” script, “gridconv” code, and “VB script embedded in Excel” to process the simulation results. We also invoked the plot command of MCNPX to draw

the fluence and dose distributions in the water phantom in the form of two-dimensional curves or color contour plots.

The simulation results, such as SOBP and transverse dose distribution from MCNPX are basically consistent with the expected results fulfilling the design aim. We concluded that NEU is a powerful design tool for a double-scattering system in a proton therapy nozzle, and MCNPX can be applied successfully in the field of proton beam radiotherapy.

## DEDICATION

To my dear wife - Liuyang

No words can express my love and appreciation to her.

## ACKNOWLEDGEMENTS

I would like to thank my committee chair, Dr. John W. Poston, Sr. for his patient guidance in my study and enthusiastic help in my life. I also would like to thank my committee members Dr. Leslie A. Braby and Dr. Michael A. Walker for their support and help throughout my study and research.

I want to thank Dr. Reece and Dr. Ford for teaching me about nuclear instrumentation and internal dosimetry and radiation biology. Thanks go to faculty and staff in the Department of Nuclear Engineering at Texas A&M University.

I want to thank the author of the NEU code, Dr. Bernard Gottschalk at Harvard University, and all the members in MCNP and MCNPX development group at Los Alamos National Laboratory for developing useful design and simulation tools. I thank Daryl Hawkins for his maintenance of the computation codes on the grove server. I thank the researchers at M.D. Anderson Cancer Center for their support in this research. I thank Dr. Harald Paganetti from Massachusetts General Hospital for his excellent presentation in the topic of proton therapy at Texas A&M University. Without their cooperation and information, this research could not have been completed.

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I also want to thank all my teachers, classmates, colleagues and friends.

Finally, thanks go to my family members and parents-in-law for their love and encouragement.

## NOMENCLATURE

2D	Two-dimensional
3D	Three-dimensional
ASCII	American Standard Code for Information Interchange
CATANA	Centro di AdroTerapia e Applicazioni Nucleari Avanzate, Italy
CSDA	Continuous Slowing-Down Approximation
EGS	Electron and Gamma Shower
FLUKA	FLUctuating KAscades
FWHM	Full-Width at Half Maximum
GEANT4	GEometry ANd Tracks, version 4
GSI	Helmholtzzentrum für Schwerionenforschung GmbH, Germany
HCL	Harvard Cyclotron Laboratory (1949-2002)
IBA	Ion Beam Applications, S.A., Belgium
LANL	Los Alamos National Laboratory
LLUMC	Loma Linda University Medical Center
MCNP	Monte Carlo N-Particles
MCNPX	Monte Carlo N-Particles eXtension
MCNP(X)	MCNP or MCNPX
MGH	Massachusetts General Hospital
NEU	Nozzle with Everything Upstream
NIST	National Institute of Standards and Technology, Gaithersburg, MD

NPS	Number of Particles to be run in MCNP(X)
PHITS	Particle and Heavy Ion Transport code System
PSI	Paul Scherrer Institute, Villigen, Switzerland
rms	root mean square
SOBP	Spread-Out Bragg Peak
SSD	Source Surface Distance
VB	Visual Basic



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## 1. INTRODUCTION

### 1.1 Overview

Currently, cancer is one of the most threatening diseases for humans and other beings. Scientists worldwide are trying to find more effective methods to conquer this disease. The current cancer therapy methods mainly include surgery, chemotherapy, radiotherapy or a combination of these (Bayle and Levin 2008). A little over 50% of all cancer patients will require radiotherapy at some time during their illness. The basic principle is to use ionizing radiation (internal or/and external) to deposit energy in a tumor to kill the cancer cells. In this thesis, we will introduce an emerging radiation treatment tool - proton beam radiotherapy, which is called “the state of the art” technique in radiation therapy (Paganetti and Bortfeld 2005).

The purpose of radiation therapy is to maximize the dose delivered to the tumor region while minimizing the dose to the surrounding healthy tissues or organs (Weber 2006). The optimized result from radiation therapy is that the profile of the spatial distribution of absorbed dose is exactly the same as the profile of the tumor volume. Although it is not so easy to obtain this optimized result, some techniques can be used to achieve it. Conformal techniques have been and are being developed to achieve this aim (Paganetti and Bortfeld 2005). “Conformal” means the shape or profile of the high dose region is the same as the tumor region, while low-dose or non-dose regions cover the

---

This thesis follows the style of Health Physics.

surrounding healthy tissues or organs. The depth-dose distributions for high-energy protons make it possible for this radiation to offer a high degree of conformity to the tumor volume. This is one important reason proton therapy techniques are increasingly popular in clinical applications.

## 1.2 Scope of This Thesis

In this thesis, we will provide a detailed design and simulation procedure of a passive-scattering nozzle in proton beam radiotherapy.

Section 2 will provide an overview of proton beam therapy, including the history, basic principles, facilities, and some considerations about this treatment method.

Section 3 will introduce the design and simulation codes and several other scripts used in this research.

Section 4 will provide a detailed design procedure of a double-scattering system. The design results will be used in Section 5.

Section 5 will provide a detailed Monte Carlo simulation procedure of a passive-scattering nozzle to verify the design results from Section 4 and obtain the dose and fluence distributions. The geometric models, particle tracks, and simulation results will be shown graphically.

Section 6 will summarize this research and point out the uncompleted tasks and future work.

## 2. OVERVIEW OF PROTON BEAM RADIOTHERAPY

### 2.1 History and Principle of Proton Beam Radiotherapy

In 1946, Robert R. Wilson (Wilson 1946) first proposed the use of protons for radiation therapy, at Harvard Cyclotron Laboratory (HCL). In 1954, the first patient was treated at Lawrence Berkeley Laboratory. By December, 2007, thirty-three proton therapy centers had been established worldwide, and twenty-six of them are still active. Seven future proton therapy facilities are currently under construction or are fully funded. From 1954 to 2007, 53,439 patients had received proton therapy (ICRU 2007).

Proton therapy is a tool to treat cancer. It uses the ionizing ability of protons to damage the DNA of the cancer cells. Like other forms of radiotherapy, proton therapy works by aiming energetic ionizing particles onto the target tumor.

The physical rationale of proton therapy is based on the well-defined penetration range of protons. In the depth-dose curve, a dose peak (called the “Bragg peak”) occurs at the end of range, shown in Figure 2-1. If a tumor is located at the Bragg peak, it can obtain the highest dose from the radiation. Since a tumor is not a point but rather a volume, a single Bragg Peak cannot cover the whole tumor. If several Bragg peaks from protons with different ranges are superimposed properly to form a wide dose peak, the target tumor can be covered longitudinally. This wide dose peak is called a “Spread-Out Bragg Peak” (SOBP), shown in Figure 2-2. The incident proton beam can form an SOBP

by sequentially penetrating absorbers with different thickness, e.g., via a range modulator or ridge filters (Chu et al. 1993).

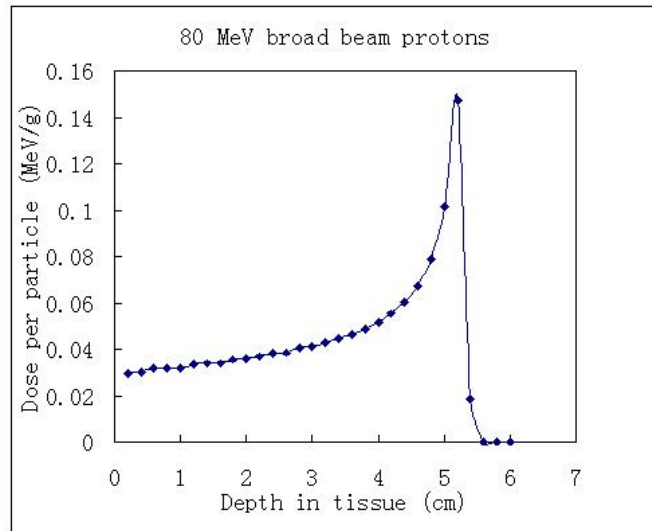


Figure 2-1. Depth-dose distribution of a broad proton beam in soft tissue

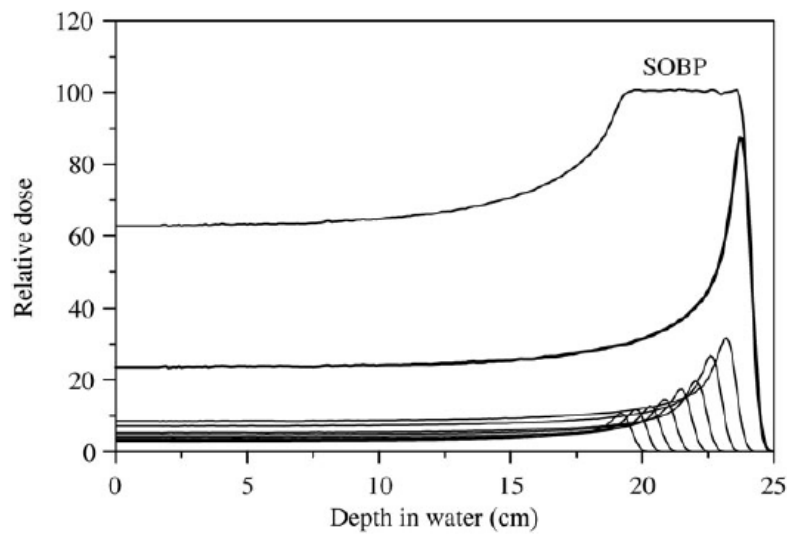


Figure 2-2. The formation of a spread-out Bragg peak  
(ICRU 2007)

The clinical rationale of proton therapy is the feasibility of delivering higher dose to the tumor, leading to an increased tumor control probability (Paganetti and Bortfeld 2005). The physician can choose the energy and intensity of proton beam according to the position and profile of the tumor. Compared with traditional photon therapy, proton therapy has many advantages, e.g., good high-dose conformity to tumor and lower dose to healthy tissue, lower tumor reoccurrence rate, and fewer adverse side effects, etc. (Olsen et al. 2007). Proton therapy is a preferred treatment method for pediatric cancers because it will not affect the growth of the young patients. A comparison of dose distributions between photons and protons is shown in Figure 2-3.

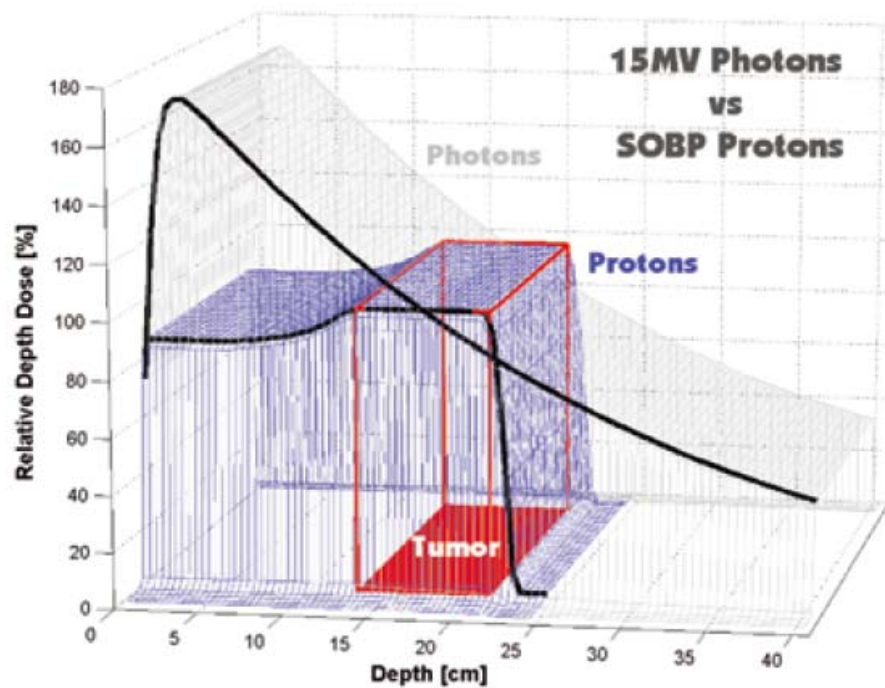


Figure 2-3. A comparison of depth-dose distributions between photons and protons  
(Weber 2006)

## 2.2 Proton Therapy Facility

### 2.2.1 Layout of a Typical Proton Therapy Facility

A typical proton-therapy facility includes three main components: an accelerator with energy selection system, a beam transport system, and a treatment delivery system. An illustration of a typical proton-therapy facility is shown in Figure 2-4, and the layout of proton-therapy facility at Massachusetts General Hospital (MGH) is shown in Figure 2-5.

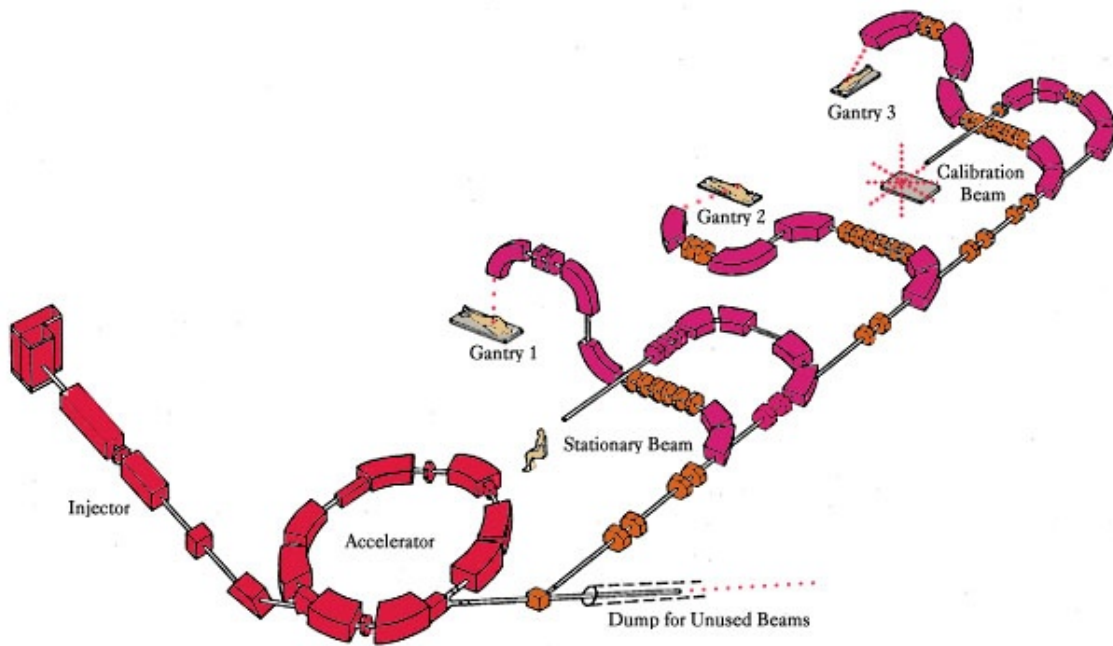


Figure 2-4. A typical layout of proton therapy facility  
(The National Association for Proton Therapy 2009)

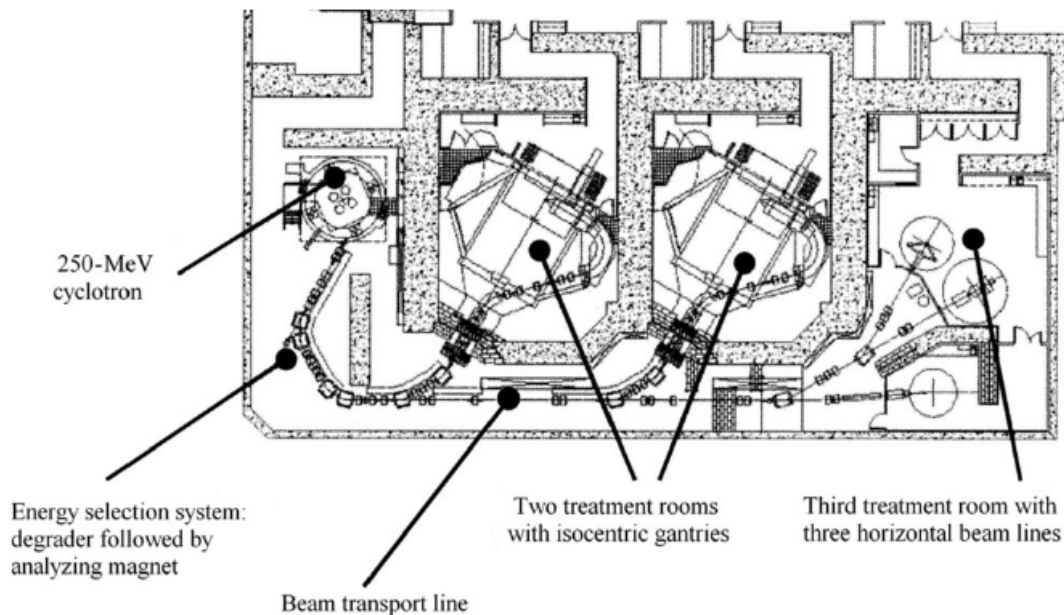


Figure 2-5. Proton therapy facility layout at MGH

(ICRU 2007)

### 2.2.2 Proton Accelerators and Beam Transport System

Accelerators are used to produce the treatment proton beams with the energy high enough to reach the deepest position of the tumors. According to the range (Berger et al. 2005) chart shown in Figure 2-6, the energy of 215 MeV is required for incident protons to achieve the depth of  $30 \text{ g cm}^{-2}$  in the body. The protons can lose some energy in the beam-modifying components, so the emerging energy of protons from an accelerator should be a little higher, i.e., 225 to 250 MeV. In addition, the beam intensity must be high enough to deliver the required therapeutic doses to the tumor within a reasonable time. If the required dose rate delivered uniformly to a one liter target tumor is 2 Gy per minute, the typical beam intensities should be in the range of  $1.8 \times 10^{11}$  to



$3.6 \times 10^{11}$  particles per minute. The exact energy and intensity requirements are dependent on the beam delivery mode (either scattering or scanning) actually used in therapy (ICRU 2007).

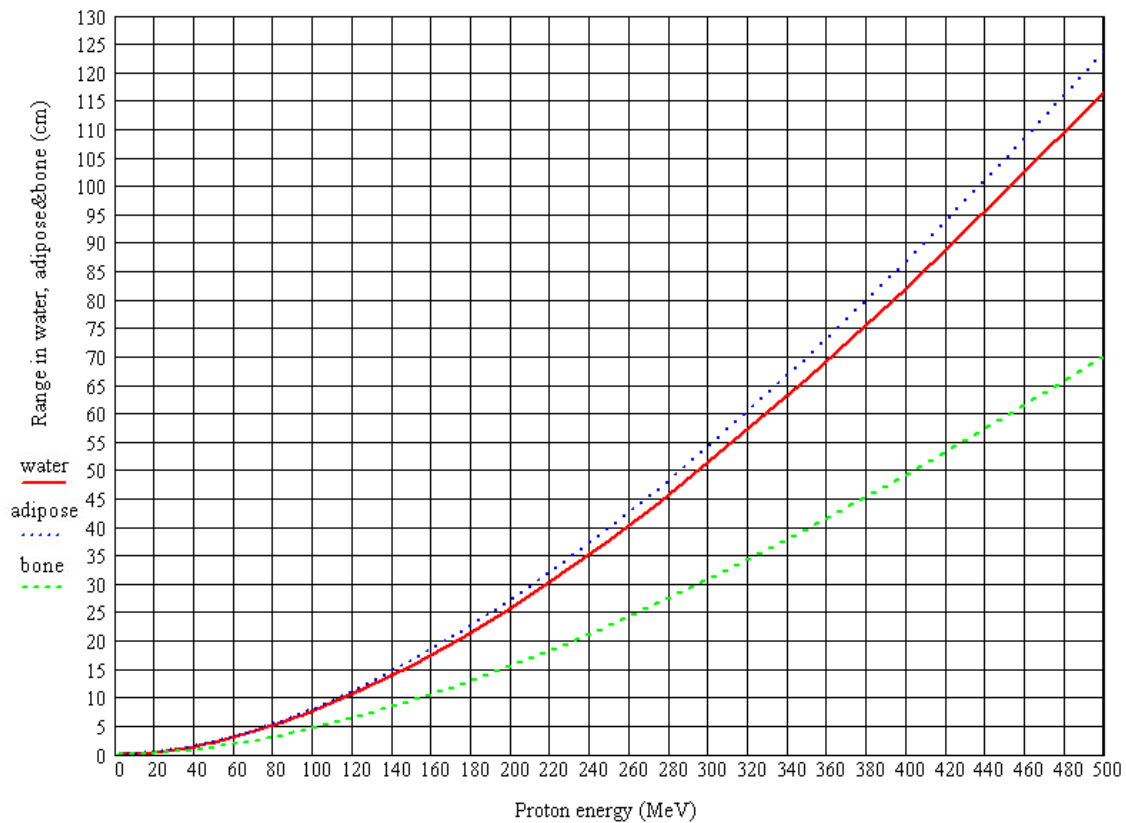


Figure 2-6. CSDA ranges of protons in water, adipose and bone

The types of accelerators that can be used to produce energetic protons include linear accelerators, classical cyclotrons, isochronous cyclotrons, synchrocyclotrons or synchrotrons. At present, only cyclotrons and synchrotrons are used in dedicated hospital-based, proton-therapy facilities. The mechanisms of different accelerators

related to proton therapy can be found in ICRU Publication 78 (2007). One cyclotron designed by Ion Beam Applications, S.A. (IBA) is shown in Figure 2-7.



Figure 2-7. An IBA isochronous cyclotron  
(Medical Physics Web 2009)

The beam transport system is actually the connection part between the accelerators and treatment rooms. It is used to transfer the beam from the outlet of the accelerator to the entrance of the treatment rooms. A series of magnets are used to bend, steer and focus the beam in the beam transport line.

### 2.2.3 Beam Delivery System

A beam delivery system can comprise several subsystems and may include some or all of the following: a gantry, a beam nozzle (equipped with a snout), a volume-

tracking and beam-gating device, and a patient-positioning and immobilization system.

The beam delivery system is enclosed in a treatment room to separate the accelerator and beam line. This protects the patients and allows personnel to move freely between treatment rooms while the beam is in use within adjacent restricted areas (ICRU 2007).

According to the alignment of the treatment head (also called “nozzle”) in the treatment rooms, the beam delivery systems are categorized into two types: a gantry system or a stationary beam delivery system, previously shown in Figure 2-4 and Figure 2-5. A fixed horizontal beam (stationary beam) can only be used to treat patients in a seated or near-seated position, e.g., a patient with an ocular tumor or tumor of the skull. Figure 2-8 shows a fixed horizontal beam delivery device at Centro di AdroTerapia e Applicazioni Nucleari Avanzate(CATANA), Italy. To irradiate a patient from any desired angle, the gantry system is introduced. A gantry equipped with a treatment head can rotate around the patient lying on a movable table. The nozzle delivers the beam to the patient at the desired angle. Two typical gantries are shown in Figure 2-9 and Figure 2-10. One is the IBA gantry at MGH, and the other is the gantry used at Paul Scherrer Institute (PSI) in Switzerland. Please notice that the bending paths for protons and isocenters are different in these two gantries.



Figure 2-8. A horizontal beam at CATANA, Italy  
(Proton Therapy Room 2008)

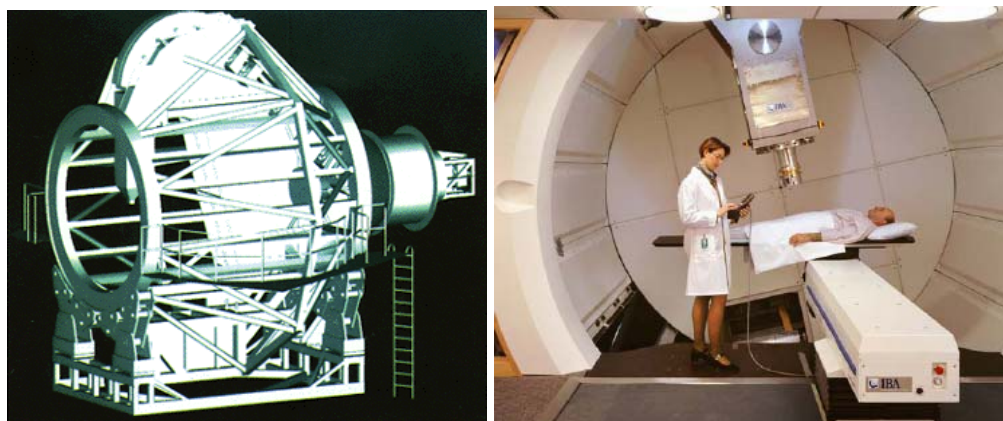


Figure 2-9. IBA gantry system at MGH  
(Lawrence Berkeley National Laboratory 2009; Paganetti and Bortfeld 2005)

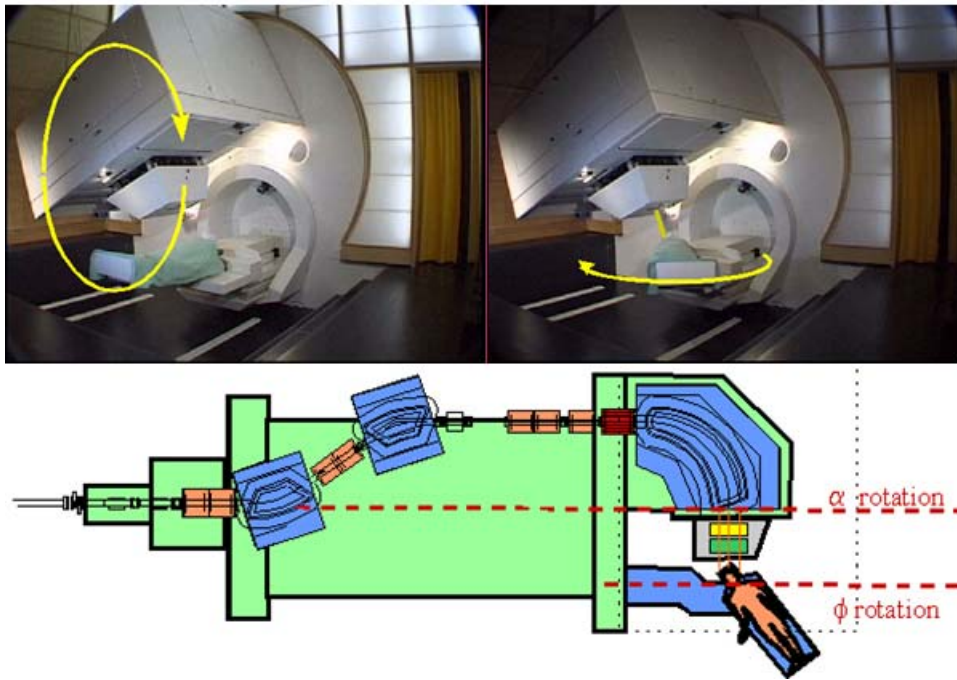


Figure 2-10. PSI gantry in Switzerland  
(PSI Proton Therapy Facility 2009)

To spread out the beam laterally, the beam-delivery techniques are categorized into two types: dynamic scanning or passive scattering (ICRU 2007). The dynamic scanning technique is a time-dependent method to deliver the desired dose to the target tumor in a scanning mode. In a passive scattering nozzle, a narrow beam is scattered by a scattering system to form a broad beam laterally.

In some dynamic scanning modes, the direction and path of a narrow beam (pencil beam) is controlled by magnets in order to scan the target tumor in a zigzag route laterally and layer by layer longitudinally. This working mode is also called “active scanning”, which looks like a painting procedure. Pencil beam scanning is an Intensity

Modulated Proton Therapy (IMPT) technique. A scanning beam is composed of a number of finite pencil beams. The schematic of an active scanning beam is shown in Figure 2-11. A broad beam (with a Gaussian lateral profile) can also be used as a scanning beam. This method is usually called “uniform scanning”, which is only used for the delivery of uniform-intensity dose distributions. It needs wobbling magnets to move the beam; besides, it also needs a scatterer, patient collimators and a compensator.

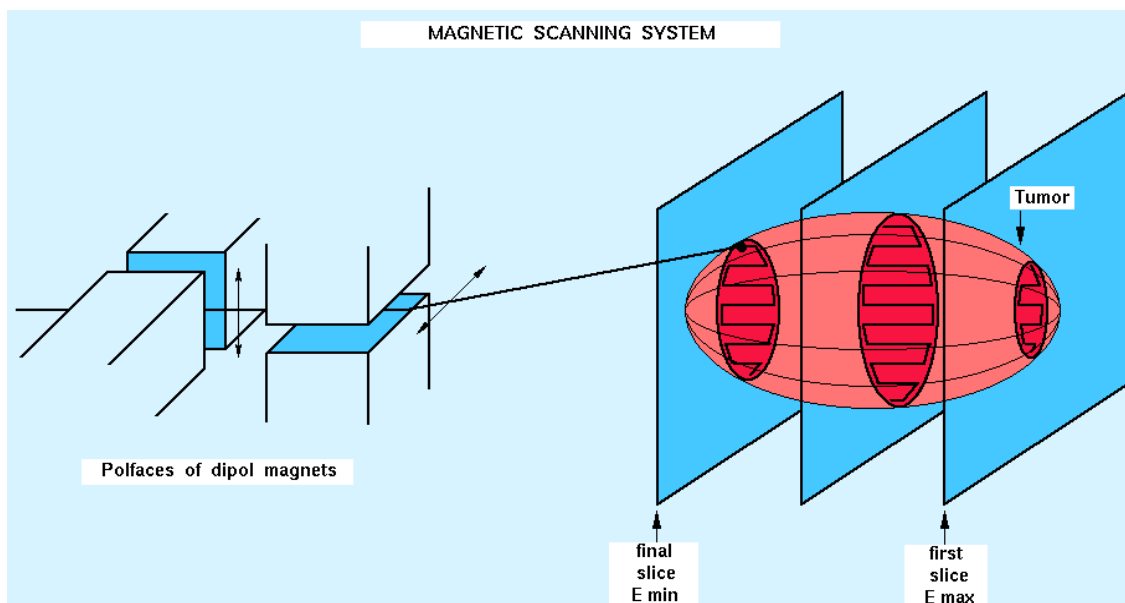


Figure 2-11. A schematic figure of a scanning beam

(GSI Heavy Ion Research Center 2008)

The design and simulation of a passive-scattering nozzle is the topic of this research. We will discuss it in detail. A passive-scattering nozzle used at Loma Linda University Medical Center (LLUMC) is shown in Figure 2-12 (ICRU 2007).

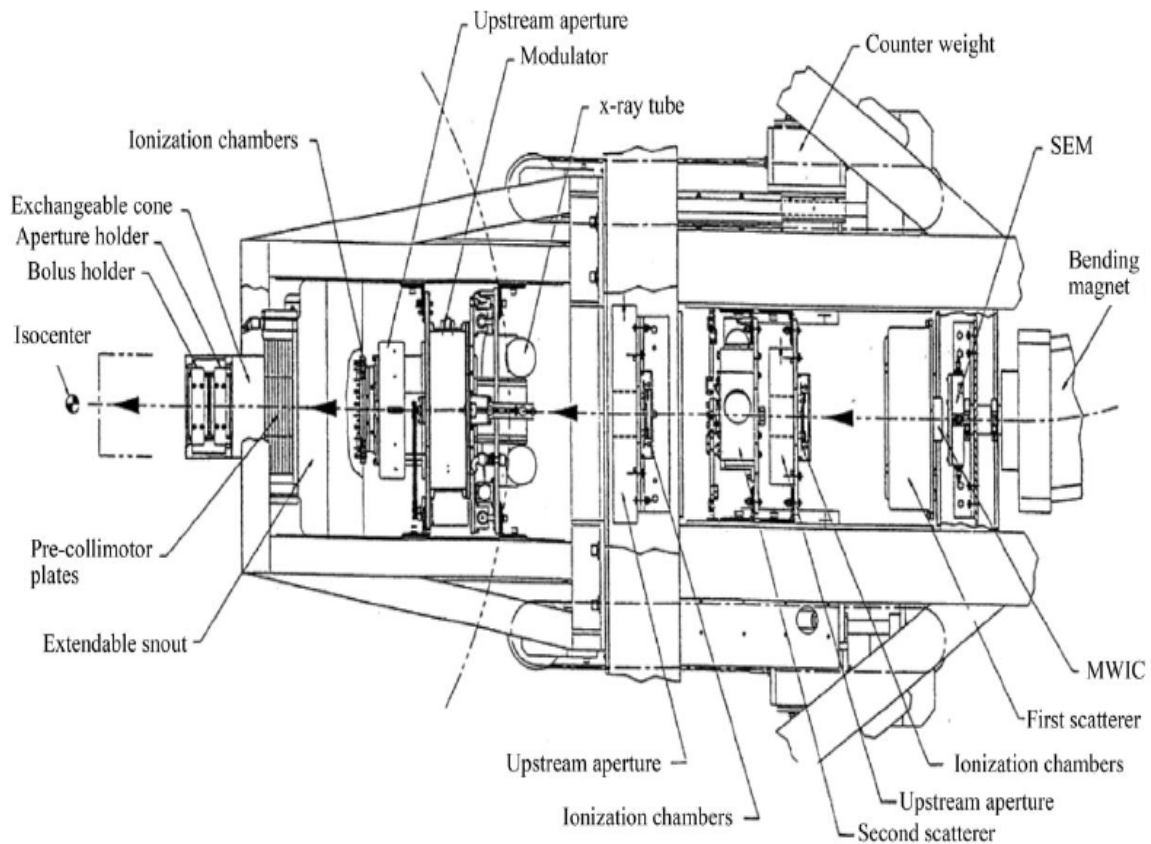


Figure 2-12. Schematic diagram of a passive-scattering nozzle at LLUMC  
(ICRU 2007)

The main components of a passive-scattering nozzle include some or all of the following: a vacuum chamber, a first scatterer, a range modulation wheel or ridge filters, a second scatterer, several collimators, a final aperture, a patient-specific range compensator, beam profile monitors, ion chambers, and a set of range shifters.

The function of scattering system is to broaden the narrow beam to form a flat transverse dose distribution to cover the tumor laterally. The function of a range modulation wheel or ridge filters is to modulate the beam to form a SOBP depth-dose

distribution to cover the tumor longitudinally. The combination of a scattering system and a range modulator (wheel or filters) make it possible to form a high-dose region to cover the tumor region fully.

The other devices, including collimators, a final aperture and a patient-specific compensator, and range shifters are used to spread and limit the beam profile to make the high dose distribution region “conformal” to the tumor region. All efforts are made to achieve one purpose: maximizing the dose in the target tumor and sparing the surrounding healthy tissues. However, even though there is a very sharp distal drop in depth-dose distribution, the healthy tissues proximal to the target are still exposed to very high doses, which can be seen from the SOBP curve discussed later. Scattering theory will be discussed in Section 4 and a simplified passive-scattering nozzle will be modeled in Section 5.

Figure 2-13 shows several different range modulation wheels. The common materials used in range modulation wheel can be plastic or aluminum alloy, depending on the therapeutic requirements. These common plastic materials include poly methyl methacrylate (PMMA), acrylonitrile butadiene styrene (ABS) resin and Lexan. Figure 2-14 shows a final aperture (brass) and a patient-specific range compensator (acrylic). Figure 2-15 shows the schematic of the formation of the “conformal” dose distribution in a passive-scattering treatment method.



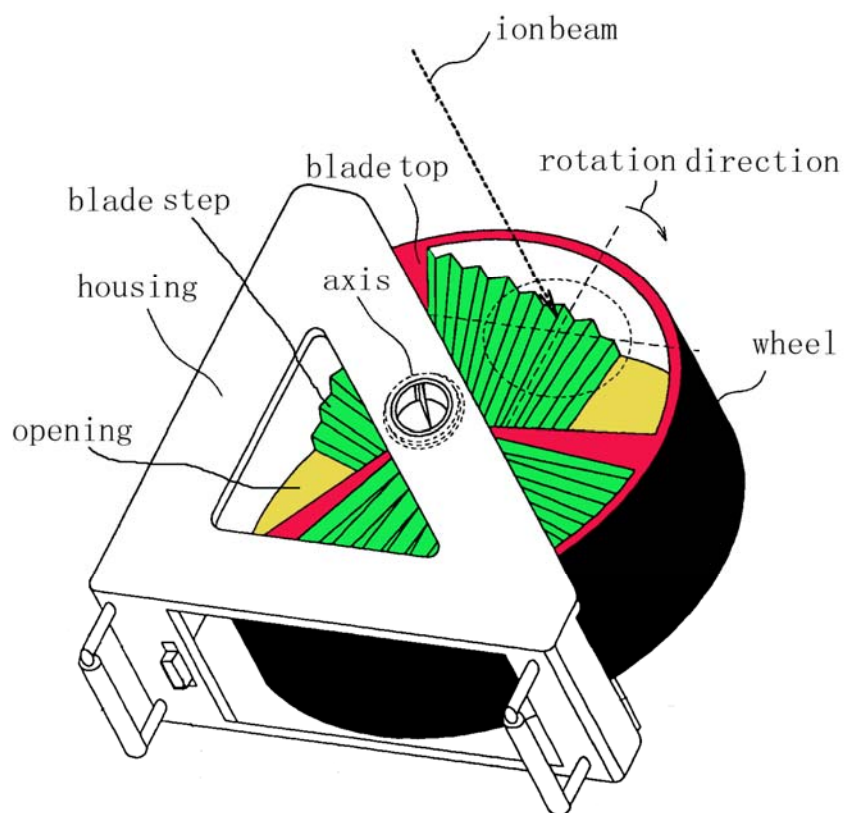
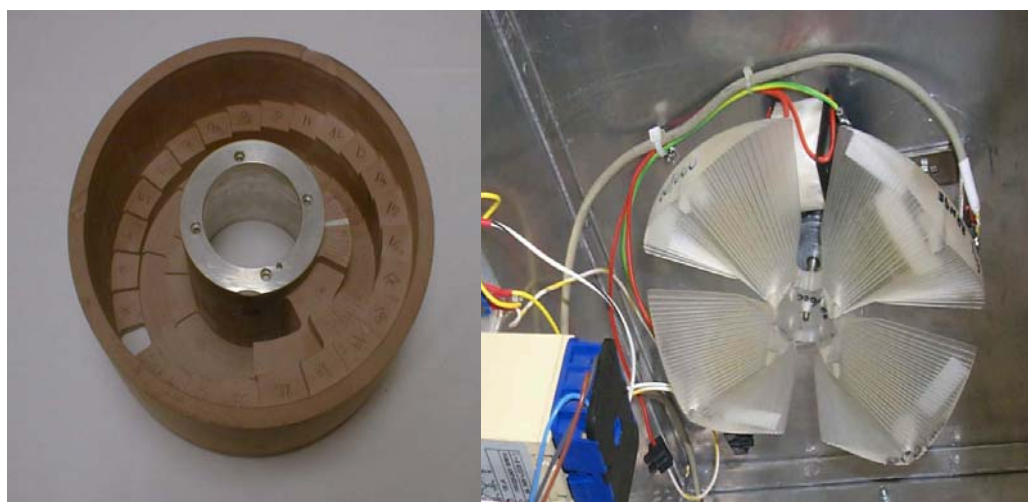


Figure 2-13. Three types of range modulation wheels

(Paganetti and Bortfeld 2005; Proton Therapy Room 2008; Free Patents Online 2009)

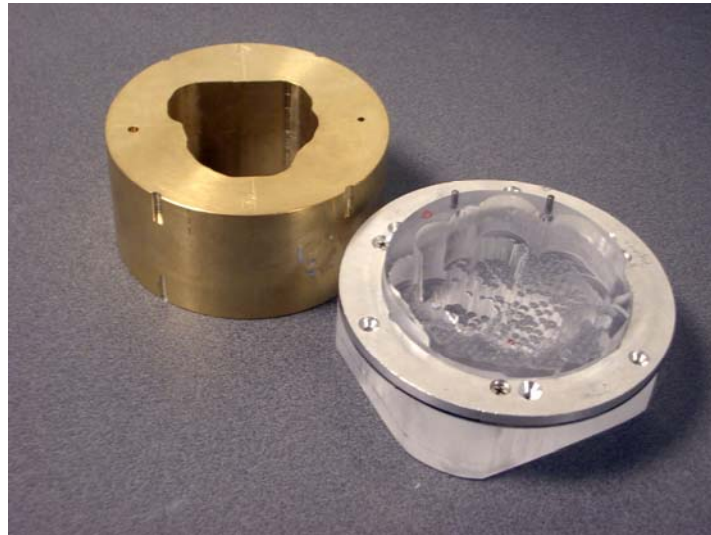


Figure 2-14. A final aperture and a patient-specific range compensator

(Gottschalk 2009)

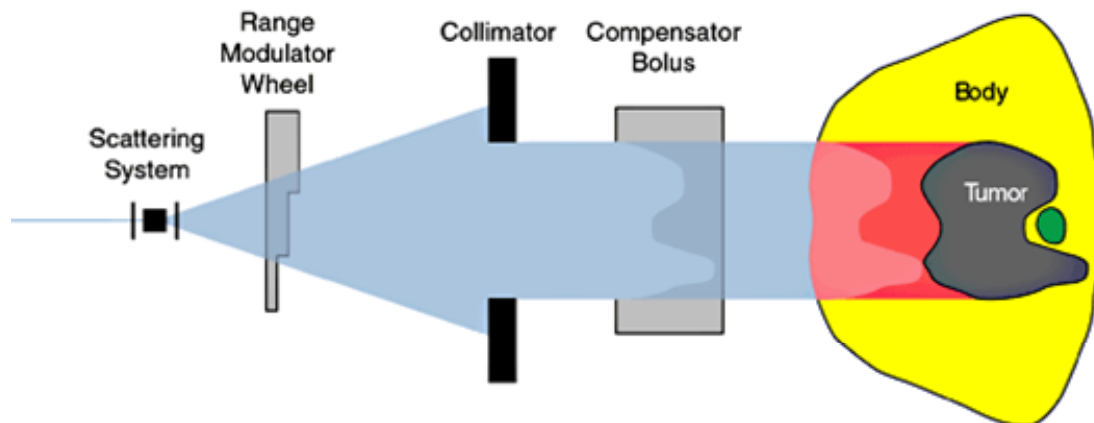


Figure 2-15. Formation of the conformal dose by a passive-scattering method

(Advanced Cancer Therapy 2009)

The examples of a broad beam and a pencil beam used at M.D. Anderson Cancer Center are shown in Figures 2-16 and 2-17. Figure 2-16.a shows the procedure how a

narrow beam passes through a rotating range modulation wheel; Figure 2-16.b shows how the scattered broad beam is shaped to the profile of the tumor; Figure 2-16.c shows how this “conformal” beam is aimed to the target tumor. Figure 2-17 shows the procedure that a pencil beam is “painting” on the tumor “actively”.

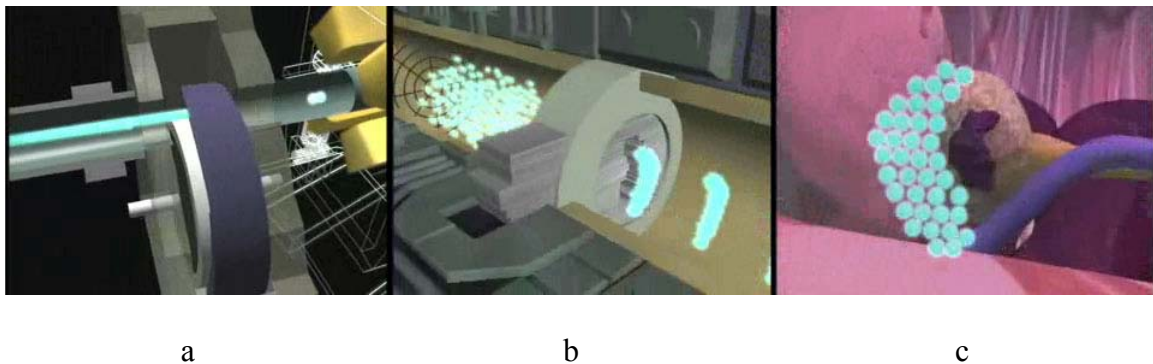


Figure 2-16. The delivery process of a broad beam

(Proton Therapy Center at M.D. Anderson Cancer Center 2008)

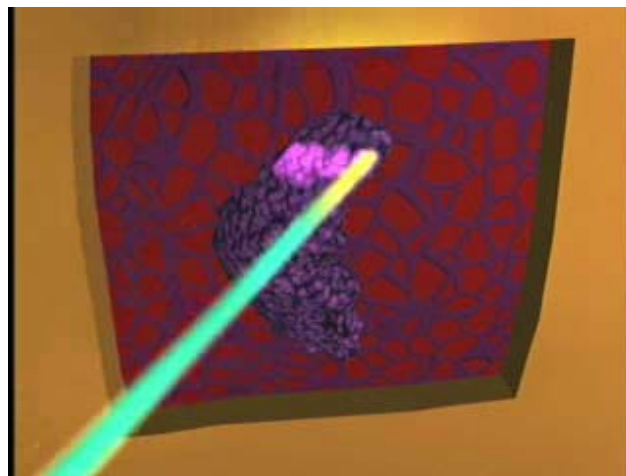


Figure 2-17. A scanning pencil beam

(Proton Therapy Center at M.D. Anderson Cancer Center 2008)

## 2.2.4 Patient Positioning System

Proton therapy is also a type of target-conformal technique (Paganetti and Bortfeld 2005). Patient positioning is very critical in the whole treatment procedure, which can affect the decision of treatment plan and the treatment effect. The tumor position and its surrounding structures, especially bones, on the treatment day should be identical to the position on the positioning day. Several factors can affect the result of the treatment. For example, if bone is suddenly moved into the beam on the treatment day, the high dose region will shift backwards, which will decrease the effect of the treatment. The physiologic motion of organs can also affect the therapy. For example, in the treatment of lung tumors, the dose will be disturbed due to the motion of lungs.

The schematic of patient positioning system (PPS) are shown in Figure 2-18 (Meggiolaro et al. 2004).

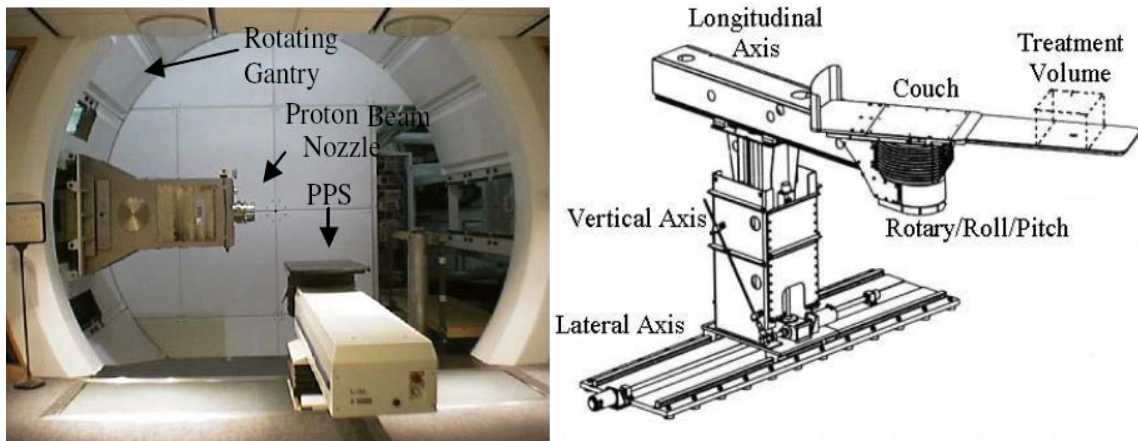


Figure 2-18. Patient positioning system

(Meggiolaro et al. 2004)

### 2.3 Biological Effects

The relative biological effectiveness (RBE) of protons is defined as the ratio of the absorbed dose from the reference radiation (photons) and the absorbed dose from protons producing the same biological effect. There are no proton RBE values based on human tissue response data, and there are no experimental data from *in vivo* systems supporting the specific RBE. Clinically, a generic RBE value of 1.1 is applied to all tissues in the direct proton beam path (ICRU 78 2007).

More information about the biological effects of proton beams in the therapeutic energy range can be found in ICRU Publication 78 (2007). Paganetti and Bortfeld (2005) also provided a systematic review on the biological effects.

### 2.4 Secondary Radiation

When protons pass through matter, they will slow down by interacting with the matter by Coulomb interactions and nuclear interactions (Paganetti and Bortfeld 2005). The secondary radiation, such as secondary neutrons and recoil nuclei, will be produced by nuclear interactions. Shielding for neutrons is necessary for any proton therapy facility. The production of neutrons cannot be avoided. Shielding can be used to reduce the effect of neutrons produced in the scattering system, the aperture and the compensator, but nothing can be done to avoid the neutrons produced in the patient's body (Paganetti and Bortfeld 2005).

The mass stopping power of protons in lead (Berger et al. 2005) is shown in Figure 2-19. The nuclear stopping power is much less than the electronic stopping power. Hence, we simplify the problem by omitting the production of neutrons and recoil nuclei. The purpose is to record the contribution from incident protons and secondary radiation separately. The effect of secondary radiation will be considered in future research.

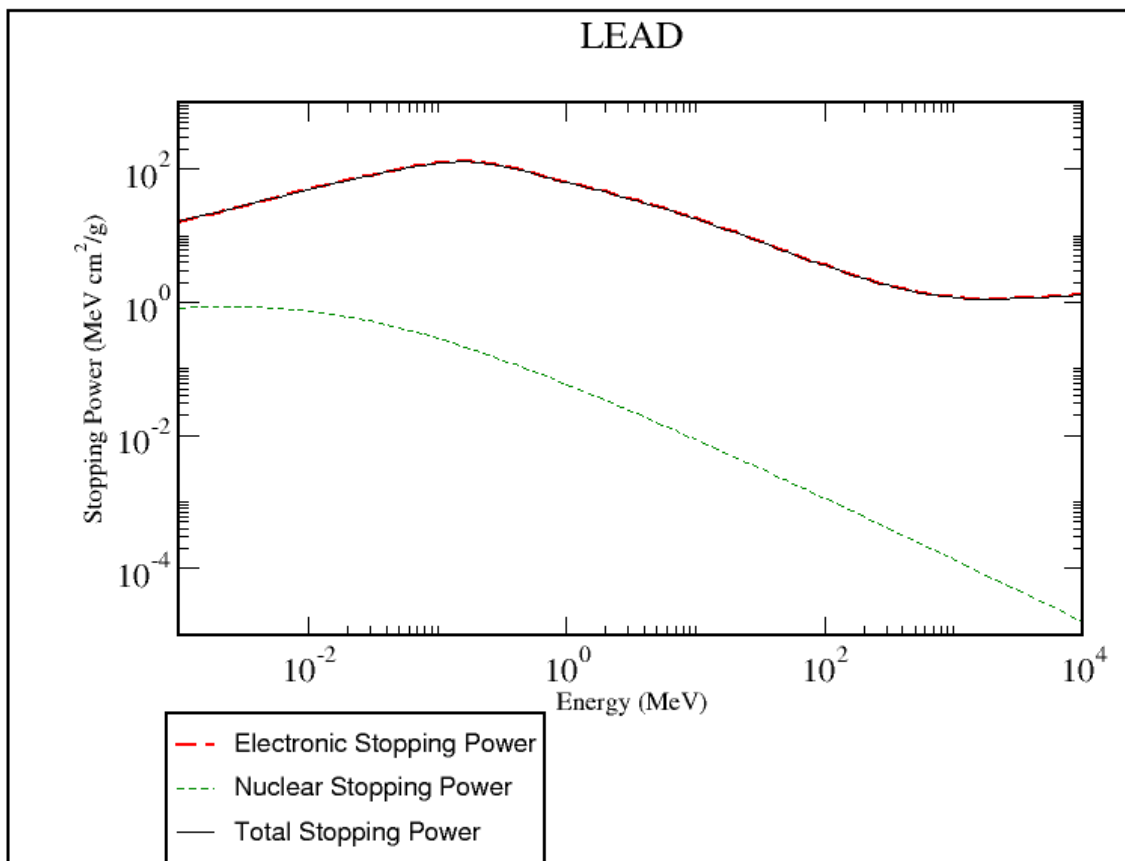


Figure 2-19. Mass stopping power of protons in lead

(Berger et al. 2005)

### 3. METHODS AND CODES USED IN DESIGN AND SIMULATION

In this Section, we will introduce a specialized design tool for a double-scattering system in a passive-scattering nozzle. We also will introduce some Monte Carlo particle transport codes. Finally, we will provide a method using static geometries to obtain simulation results for dynamic geometries. This is the most valuable part of this research. This method was realized by executing several auxiliary codes or scripts.

#### 3.1 NEU Codes Package

The NEU codes package is a specialized tool kit including several subroutines for designing the double-scattering system in a passive-scattering proton nozzle. NEU means “Nozzle with Everything Upstream,” which is also the design principle for a passive-scattering nozzle. Firstly, NEU was written by Bernard Gottschalk in FORTRAN language at HCL in the late 1980’s. During the next few years the program and designs were furnished to a number of proton therapy centers. NEU was tested at HCL and used to design the standard IBA nozzle and components for M.D. Anderson Cancer Center. Currently it is still very useful for the design and improvement of proton therapy nozzles.

The compressed package of NEU (named “BGware.zip”) can be obtained from Dr. Gottschalk’s website. NEU can be run on a Windows XP operating system. The

structure of the “BGware” directory is shown in Figure 3-1, and the contents are listed in Table 3-1.

We mainly used the files or codes in subdirectories “bpw”, “data”, “execs”, and “neu.” All the executable files, such as “bpw.exe” and “neu.exe,” are contained in the directory of “execs.” Input files and output files related with “neu.exe” are located in the directory of “neu.” The files related to “bpw.exe” are contained in the “bpw” directory. The fitted Bragg peak data files (\*.bpk) created by “bpw.exe” are copied from the directory of “bpw” to “data.” These \*.bpk files will be invoked by “neu.exe.” More detailed information can be found in NEU User Guide (Gottschalk 2006). The function and application of each code will be described in Section 4 with examples.

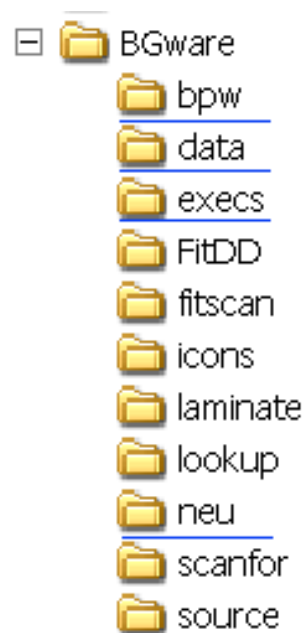


Figure 3-1. The structure of the BGware directory



Table 3-1. The list of files or codes in the BGware directory

Sub-directory	Description	File type or name
bpw	Original Bragg peak data	*.txt; *.dat; *.psi; *.scx
	Input files for bpw.exe	bpw.inp
	Fitted Bragg peak data	*.bpk
	Other output files	*.bmp
data	Range-energy table	*.ret
	Fitted Bragg peak data	*.bpk (copied from bpw directory)
execs	Executable files	Bpw.exe; CPO2.exe; FitDD.exe; fitscan.exe; laminate.exe; lookup.exe; neu.exe; scanfor.exe
neu	Catalog of NEU runs	NEU.CAT
	Input files for neu.exe	Neu.inp
	Beam line picture input file	BEAMPICT.INP
	Output files	*.mod; *.con; *.axd; *.axh; *.bmp; *.out; *.log; SOBP.DAT

### 3.2 Monte Carlo Codes

The Monte Carlo (MC) method was first used in the 1940s by physicists working on nuclear weapon projects at the Los Alamos National Laboratory (LANL). It is a random sampling computational algorithm (Wikipedia 2008). It is not limited by the dimensions of problems. It tracks the histories of a large number of individual events and records some aspects according to their average behavior. It has been applied in many fields, such as high-energy physics, radiation detection, space radiation, medical physics, and economics. With the computerization spreading, lots of MC codes have been developed. In the particles transport field, the popular MC codes include, MCNP(X), EGS, Geant4, Fluka, and PHITS.

Monte Carlo dose calculations are known to be more accurate in dosimetry than commercial treatment planning systems that are based on fast but approximating semi-empirical algorithms (Mohan and Nahum 1997). Except for EGS, any of the other MC codes mentioned above can be used in the simulation of proton therapy problems. The accuracy of MCNPX in predicting dose distributions in proton therapy have been confirmed by previous investigations (Newhauser et al. 2007), and the benchmarking of MCNPX applied to the nozzle at M.D. Anderson Cancer Center (Titt et al. 2008). The MCNPX code was adopted for this research. The other available codes will be applied in future work.

MCNP and MCNPX were developed by LANL. MCNPX is the extension of MCNP. MCNP can only be used to track neutrons, photons and electrons (X-5 Monte Carlo Team 2003), but MCNPX can be used to track nearly all particles at nearly all energies (Pelowitz 2008). MCNPX utilizes the latest nuclear cross section libraries and uses physics models for particle types and energies where tabular data are not available. Several different tally cards can be used to score different physical quantities. The tally results are tabulated in the pair of “mean and relative error”. The guidelines for interpreting the relative error in statistics from tallies are shown in Table 3-2.

Visual Editor is a visualization tool for MCNP or MCNPX. It was developed by Schwarz RA and maintained by Visual Editor Consultants (Schwarz et al. 2008). It integrates an `mcnp5.exe` or `mcnp5.exe` in its kernel. It can be used to aid the user to create the input files, show the geometry and particles tracks, and plot the cross section and tally results. The geometry visualization can be in two-dimensional (2D) or three-

dimensional (3D) mode, but the particle source and tracks can be shown only in 2D mode.

In this research, we will also use “gridconv”, a subroutine of MCNPX, to deal with the data from the mesh tally.

Table 3-2. Guidelines for interpreting the relative error  $R^a$

Range of R	Quality of the tally
0.5 to 1.0	Not meaningful
0.2 to 0.5	Factor of a few
0.1 to 0.2	Questionable
< 0.10	Generally reliable
< 0.05	Generally reliable for point detectors
<sup>a</sup> The guidelines are listed in MCNP5 Manual Volume I (X-5 Monte Carlo Team 2003).	

### 3.3 Other Codes and Scripts

Before we introduce the other codes and scripts, the problem of dynamic geometries must be discussed.

#### 3.3.1 Solution of Dynamic Geometries

MCNP(X) can be used only to simulate problems with static geometries and fixed settings for a single problem. The input parameters cannot vary with time. However, in a real passive-scattering proton therapy problem, in order to realize a SOBP dose curve, the range modulation wheel is rotating during the whole treatment process. It seems impossible to simulate the dynamic proton therapy problem using MCNPX. A common method used to produce a SOBP curve is to use the “Matlab” code to solve a matrix-equation to obtain a “weighting factor” for each pristine Bragg peak (Oertli 2006),

and to use the solution as the probability on the energy spectrum of the source. In this method, the particles after the range modulation wheel are assumed to be the incident source particles, because the wheel is not simulated physically. Actually, this method can be considered only to be an approximate solution, because not all the SOBPs have rational solutions, and in some cases, the obtained weighting factors are negative.

In this research, if the design parameters are chosen properly, the weighting factor and thickness of each step in a range modulation wheel can be obtained using NEU. Hence, it is not necessary to solve a matrix-equation. Assuming that a whole wheel is modeled in an MCNPX input file, can we make it rotate in the treatment procedure? The answer is obviously no. We cannot simulate a real rotating wheel using MCNPX, but we can try some methods to obtain the same results as if the wheel were “rotating.” Since parameters for each step are known, we can setup a series of input files, in each of which, only one step is used and the weighting factor of each step is used as the “weight” of source. After running each problem using MCNPX, we sum the results from different cases. We can declare that the sum is the same as the result from a rotating wheel. So, this simulation method can be called “replacing the dynamic by static.” As to the normalization of the simulation results, we will discuss this topic in Section 5.

### 3.3.2 Mcnp\_pstudy Script

To create the input files for each step one by one requires a great deal of time. In order to save time, we used a Perl script “mcnp\_pstudy” to set up input files in a batch.

This Perl script was developed by Brown et al (2004). It can be used to automate the setup, execution, and collection of results from a series of MCNP5 Monte Carlo calculations, each of which must contain varying input specifications. If the setting of this script is proper, it can also be used to run MCNPX problems, but currently it cannot be used to collect the mesh tally results from MCNPX. In this research, we used this script to create a series of input files, each of which contained the parameters from a single step in a modulation wheel, such as the materials, thicknesses and source weight.

Before using “mcnp\_pstudy”, the user should point the “\$MCNP” on the 152<sup>nd</sup> line to the location of user’s MCNP(X). In some cases, the users should also comment the 153<sup>rd</sup> line to run this script correctly. The modification is shown in Figure 3-2.

```
$MCNP      = "/usr/local/mcnpv26f/bin/mcnpv26f";
# die("*** change the line above to the location of MCNP on your system\n");
# line 152 and 153 were modified by Fada Guan, 2009 07 29
```

Figure 3-2. Modification of the “mcnp\_pstudy” script

To limit the float number to 4 for a result from an arithmetic expression, a line of script was added after the 654<sup>th</sup> line (`$val = eval( $expr );`) in original “mcnp\_pstudy” script. The added content is “`$val = sprintf("%.4f", $val); ,`” so that the contents in each line of the new produced input file will not exceed the limits of the 80<sup>th</sup> column required by MCNP(X).

### 3.3.2 Merge\_mctal Script

Even though “mcnp\_pstudy” cannot be used to collect the mesh tally results from MCNPX problems, we can solve it by using another Perl script “merge\_mctal.” This script is used to merge and average the tally results and statistical uncertainties. This script was developed by Brown and Sweezy in 2003 and was updated by Brown in 2004 (Brown 2008). It is designed only for MCNP5 tally results. Because there are some minor discrepancies (mainly the particle types) in the “mctal” files (tally data file) between MCNP5 and MCNPX, this script cannot be used to merge “mctal” files from MCNPX directly, especially for mesh tally results. However, if the mesh tally is modified to be a “F5” detector tally and the format of particle types is modified, the “merge\_mctal” script can be used to merge the “mctal” files from MCNPX. After the combination, the “F5” tally in the new combined mctal file should be changed back to the mesh tally. In other words, first the “mctal” file is changed from MCNPX format to MCNP5 format; then it is changed back.

### 3.3.4 VB Scripts Embedded in Excel

Additionally, referring to the VB scripts of output visualization (Schwarz 2007), we also developed some “VB scripts embedded in Excel” to read in the data from “mctal” files (in DOS format) from MCNP5 or MCNPX and distribute the data in a specified order and format. These VB scripts are very convenient for the users to deal with the data and plot the curves.

### 3.4 Systematic Flow of the Application of These Codes

The systematic flow of the application of all the codes is shown in Figure 3-3.

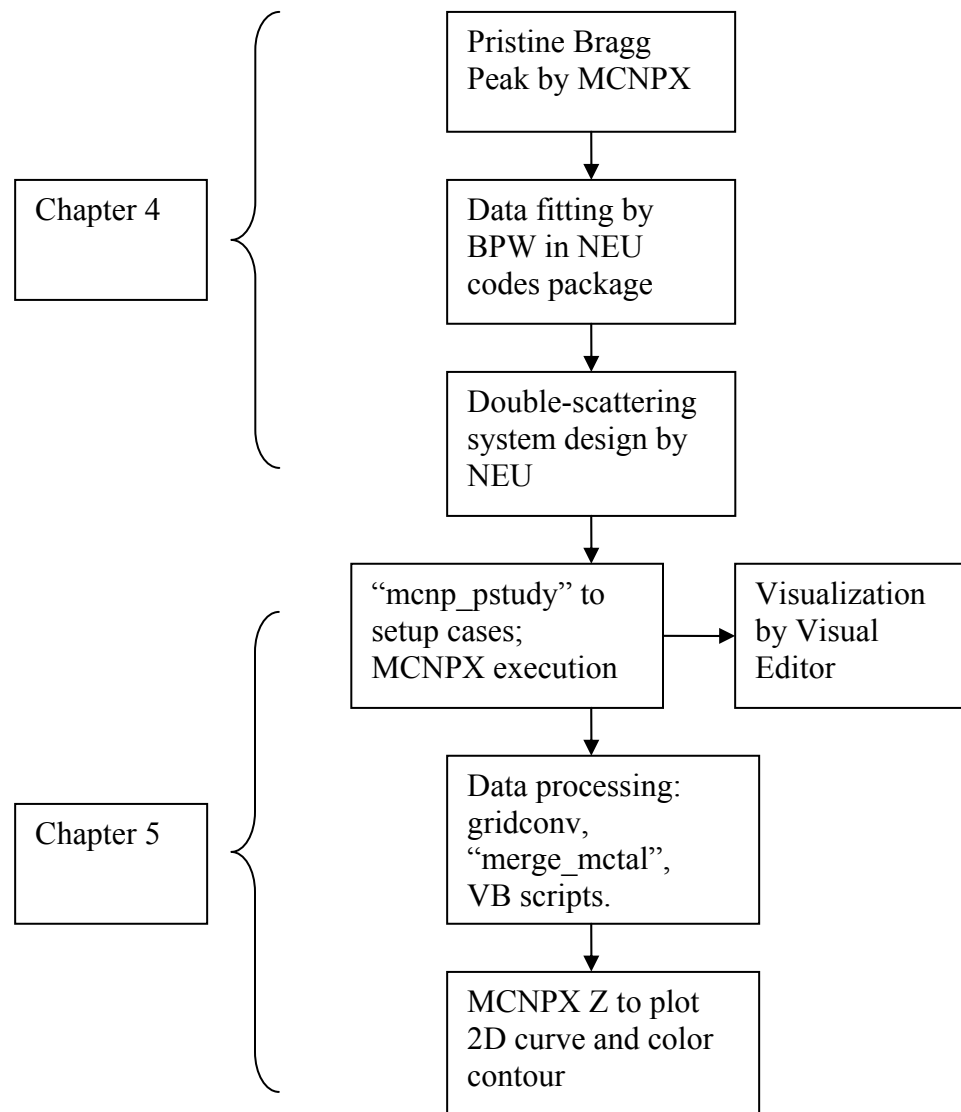


Figure 3-3. Flow chart of the application of codes

#### 4. DESIGN OF A DOUBLE-SCATTERING SYSTEM

In this section, we will introduce the whole design procedure of a double-scattering system in a passive-scattering nozzle in proton beam radiotherapy.

The first step is providing measured depth-dose data in a water phantom from a broad monoenergetic proton beam. In this research, because the clinical measurement data were not provided, we used the simulated depth-dose data in a water phantom from MCNPX to replace the measured data.

The second step involves using the BPW code in the NEU codes package to fit the depth-dose data from MCNPX. These data are read into the NEU code as the reference data to be used in designing the double-scattering system for a given SOBPs width.

The third step involves using the NEU code to design the double-scattering system, including a wheel (first scatterers and range modulation steps) and a contour-shaped scatterer, to meet the requirements in clinical applications.

##### 4.1 Simulation of Pristine Bragg Peak by MCNPX

The NEU code cannot be used to compute the effective stopping power theoretically, while it deduces it from the user's measured Bragg-peak data, in which, all the related effects such as nuclear effects, range straggling, and energy spread about the user's facility, are included automatically. This is why the user needs to provide the



measured Bragg-peak data first. An MCNPX simulation was used as an “experimental measurement.”

In this step, the necessary input parameters include the energy, radius and direction of the incident proton beam, the distance between source and the surface of the water phantom, and the size of the water phantom.

The synchrotron at M.D. Anderson Cancer Center can provide proton energies between 100 MeV and 250 MeV. The available eight energy intervals include 100, 120, 140, 160, 180, 200, 225 and 250 MeV (Zheng et al. 2008). The source particles were limited to a disc with a radius of 10 cm on the  $Z = -300$  cm plane and the emitting protons were assumed to fly along the  $Z$ -axis.

A recommended value of source surface distance (SSD) in ICRU Publication 78 (2007) is 300 cm. We also adopted 300 cm as the SSD in this research.

According to the range of protons in water, shown in Figure 2-6, a 40 cm depth is enough to stop the 250 MeV protons. In order to capture all the protons, the lateral side of the water phantom was set to 80 cm. The final size of the water phantom was 80 cm×80 cm×40 cm. The size of each water cell used to score the absorbed dose was 80 cm× 80 cm× 0.1 cm.

We used Visual Editor to check the geometry and the particle tracks. Figure 4-1 shows the tracks of 250 MeV broad-beam protons.

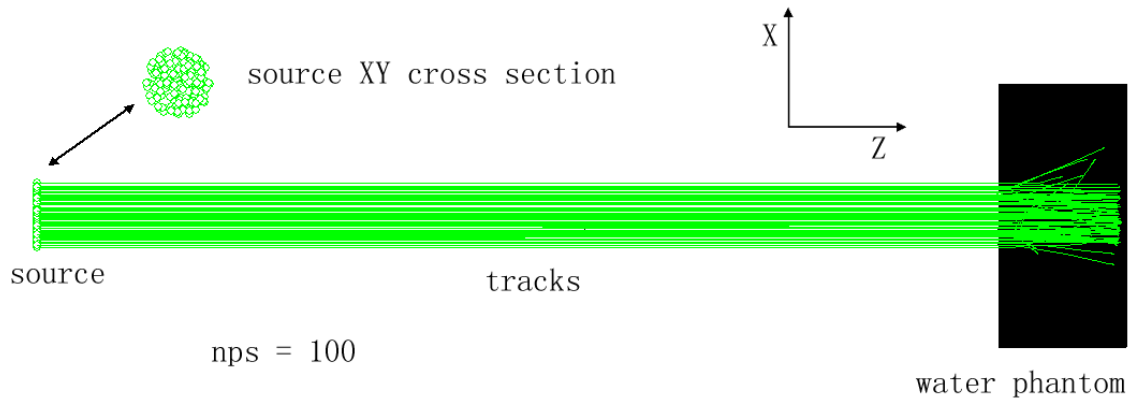


Figure 4-1. The tracks of 250 MeV broad-beam protons

The number of particles to be run was set to one million. The simulation results from MCNPX for 8 energy groups were plotted as depth-dose curves and are shown in Figure 4-2. Most of the tally results satisfied the statistical requirement shown in Table 3-2. However, several tally regions near the range of protons had higher relative errors due to the lower-sampling efficiency. The Bragg peak positions for 8 energies in this experimental setting are listed in Table 4-1. A comparison of projected ranges between MCNPX and NIST data (Berger et al. 2005) are also listed in this table. The MCNPX simulation can be accepted because the relative differences are lower than 5%.

In the following research, only 100, 180 and 250 MeV were used as the initial kinetic energies of the proton beams.

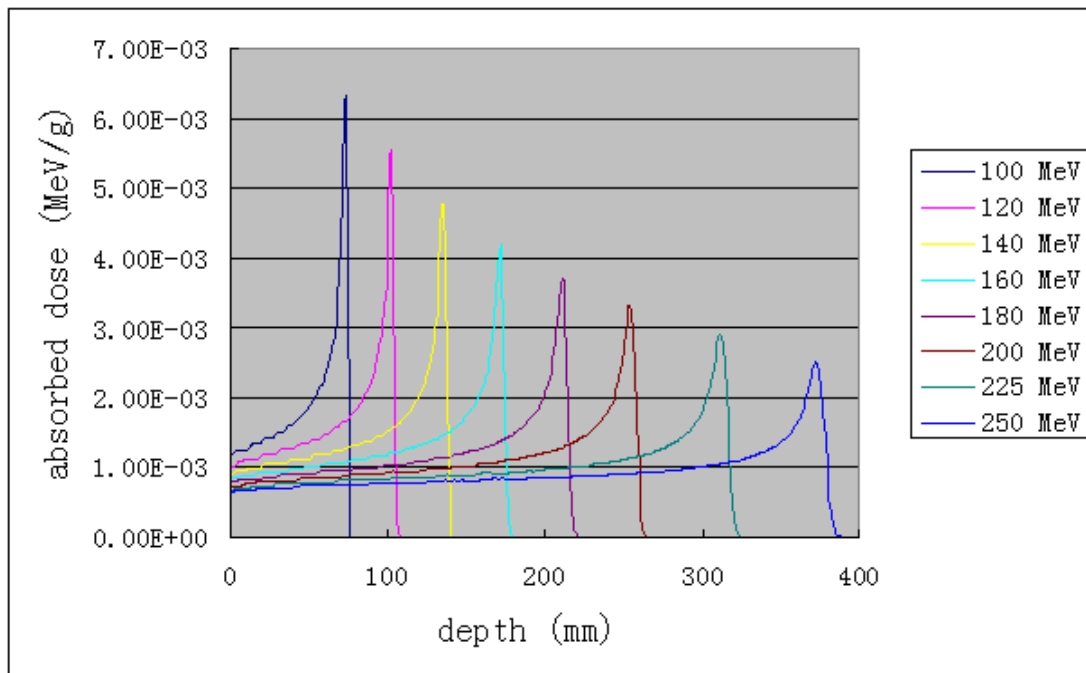


Figure 4-2. Depth-dose curves for broad-beam protons

Table 4-1. The Bragg peak position and range in water phantom

Energy (MeV)	Peak position (cm)	Projected range from MCNPX (cm)	Projected range from NIST (cm)	Relative difference (MCNPX-NIST)/NIST
100	7.4	7.9	7.707	2.50 %
120	10.3	10.9	10.65	2.35 %
140	13.6	14.3	13.96	2.44 %
160	17.2	18.2	17.63	3.23 %
180	21.1	22.3	21.63	3.10 %
200	25.4	26.9	25.93	3.74 %
225	31.2	33.1	31.71	4.38 %
250	37.3	39.6	37.90	4.49 %

## 4.2 Fitting Process of SOBP Data by BPW

The measured Bragg-peak data include a large number of data points. Before these can be read in by the NEU code, the data have to be fitted using the BPW code to decrease the number of data points. The function of the BPW code is to read in the measured data and convert them into a cubic spline form.

First, a data file, e.g. “MDA250.txt” containing the 400 pairs of depth-dose data from 100 MeV case by MCNPX, was set up.

Second, the “bpw.inp”, shown in Figure 4-3, was edited. Only two parameters were modified: the data file (“MDA250.txt”) and the distance from the source to the Bragg peak, shown in Table 4-2. On the new fitted SOBP curve, there are only 20 data pairs by default.

```
'MDA250.TXT'      data file
337.3             BP measurement source distance (cm)
0                depthCorr (cm H2O), added to x after scaling
.1              fraction of scan to extrapolate to 0 (may be 0)
0                # new points to add at beginning
----- SOBP fit (BSfit) -----
'T'             Terse or Verbose
'G'             fit method: GridParab (preferred), Marquardt, None
10 .0001        passes, convergence on rms
.5 10           reduction factor, initial lambda
.01 .01         delta x,y (best may depend on G,M)
20 0 0 0 0     pts/segment (pristine peak has only 1st segment)
2 1            deriv smoothing rms, # passes
.015 .005 .5   rms/avg < p1 -> 2 seg; distal cut; power
.05 .015       AB/AC < p1 -> 2 segs; rms/max > p2 -> add point
-----
Data files available for BPW tests in \BGware\bpw:
MDA 250MeV, to produce MDA250.bpk for NEU
Fada Guan 2009 08 02
```

Figure 4-3. Snapshot of “bpw.inp”

Table 4-2. Parameters modified in “bpw.inp”

Energy (MeV)	Data file name	Distance from source to peak (cm)
100	MDA100.txt	307.4
180	MDA180.txt	321.1
250	MDA250.txt	337.3

The routine “bpw.exe” was used to execute the fitting process. Figure 4-4 shows the execution of BPW code. Figure 4-5 shows the fitting results (marked by open squares) and related deviations (at bottom) for 250 MeV protons.

```

bpw
File Edit View State Window Help
Graphic1
working path      ?  \BGware\BPW\
data file         ?  MDA250.TXT
source distance (cm) ?  337.3
fraction for extrapolation ?  0.10
# initial points to add ?  0

Get data ...
Data file is handmade ...
# data points = 400
Fit data with cubic spline ...
nPairs,rms (%) = 20 0.221
Write .BPK file ...
Test .BPK file ...
x1 (cm H2O) = 0.10
d80 (cm H2O) = 37.64
MeV from d80 = 248.66
MeV from BP area = 245.80
tArea/td80 = 0.989
Bragg(0) = 3.899 MeV/(g/cm2)
Bragg(x1) = 3.911 MeV/(g/cm2)

<Enter> for graphics, again to continue ...

Running | Input pending in Graphic1

```

Figure 4-4. Execution of the BPW code

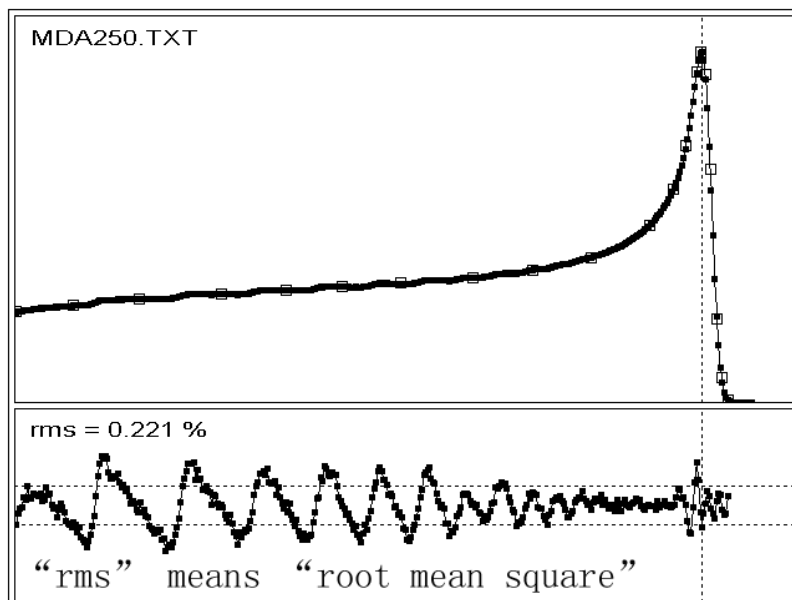


Figure 4-5. The fitted Bragg curve for 250 MeV protons

The fourth step involved copying the output data files “MDA100.BPK,” “MDA180.BPK,” and “MDA250.BPK” to the directory of “data.” The snapshot from “MDA250.BPK” is shown in Figure 4-6.

```

put a comment here ...
337.3 cm from source to Bragg peak
20 0.10000E+31 0.10000E+31 # pts, ypl, ypn
0.1000 3.1830 6.7302 11.2463 14.7008 17.7984 20.9604 cm H2O
24.8935 28.1274 31.2589 34.4288 35.7752 36.4336 37.0212
37.2115 37.4787 37.7541 38.1065 38.3570 38.7000
0.2591 0.2788 0.2955 0.3098 0.3209 0.3300 0.3407 rel dose
0.3566 0.3780 0.4145 0.5056 0.6108 0.7326 0.9451
1.0009 0.9378 0.6646 0.2397 0.0727 0.0057

```

Figure 4-6. Snapshot of “MDA250.BPK”

### 4.3 Design of Double-Scattering System Using NEU

The theory of multiple scattering is the design basis for a scattering system. When the beam particles pass through a medium, they can interact with the nuclei of the medium. Finally, the beam can be deflected in a small angle away from its original central trajectory. Elastic Coulomb scattering is the main reason for this small-angle deflection. The deflections lead to the creation of lateral-particle fluence distribution. The angular distribution of the deflected particles is roughly Gaussian for small deflection angles (Chu et al. 1993).

The current scattering system includes two components, so it is called “double-scattering” system. The first scatterer is usually a flat metal foil. According to “multiple scattering” theory, the particle fluence after the first scatterer is not distributed uniformly laterally but is best represented by a Gaussian distribution. Thus the dose distribution laterally will also be Gaussian-shaped. Particles in the central beam need more scattering to obtain a uniform fluence laterally. A Gaussian-shaped scatterer is added to the beam line to achieve this goal.

The range modulation wheel is also integrated into the double-scattering system. It is usually mated on the first scatterer. The benefits of this approach are two-fold: saving space and decreasing scattering near the patient. This design obeys the principle of “nozzle with everything upstream.” This combined component is called “S1” in the NEU code. The Gaussian-shaped (also called contour-shaped) scatterer is called “S2.”

A simplified schematic of the NEU design is shown in Figure 4-7. Notice that, in NEU design, except the double-scattering system, no other components in a real nozzle are provided.

The design is actually an interactive process. The user first specifies some approximate conditional parameters and fixed design goals. Then the user executes the NEU code to obtain the design results. If the design results are not satisfactory, the user can modify the input file and execute the code again until the design goals are achieved. It is a good way to approach the desired results by repeated execution of the code.

The fixed conditional parameters include beam energy, field radius  $r_1$  (on the surface of water phantom), SSD, SOBP width. The approximate conditional parameters include throw (distance between reference plane of S1 and the middle of SOBP), field radius  $r_2$  (on the plane at mid-SOBP, also called “useful radius”), and the scattering strengths of S1 and S2. In addition, the NEU code generally assumes the incident beam is an ideal beam (no size, no angular divergence and perfect steering). Notice that  $r_1$  or  $r_2$  is the radius at which the transverse dose falls 2.5% below the 100% level on the corresponding plane, but not the radius of the beam spot on the plane.



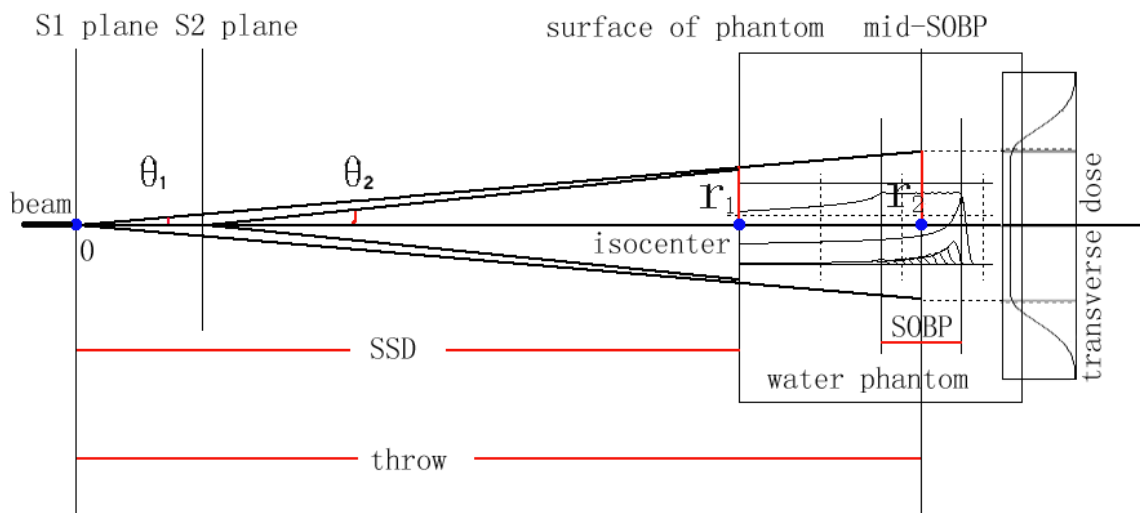


Figure 4-7. A simplified schematic of the NEU design

Both of S1 and S2 are bi-material components. The reference planes of S1 and S2 are also the mating surfaces of the two types of materials. In S1, on the left of mating surface ( $Z = 0$  cm) is the scatterer made of high-Z material, e.g., lead, and on the right is the range modulation wheel made of low-Z material, e.g., Lexan (plastic). In S2, on the left of mating surface ( $Z = 50$  cm) is the inner-contoured compensation part made of low-Z material, e.g., Lexan, and on the right is the contoured scatterer made of high-Z material, e.g., lead. The compositions and densities of the materials used in NEU design are listed in Table 4-3.

Table 4-3. Materials used in NEU design

Material	Composition	Density ( $\text{g cm}^{-3}$ )
Lead	Pb (100%)	11.35
Lexan	H (5.5%), C (75.6%), O (18.9%)	1.2
water	H <sub>2</sub> O	1.0

In a specified medium, the expected SOBP width cannot be larger than the maximum range of the beam. Hence, we set three different SOBP widths for the three beam energies listed in Table 4-4.

Table 4-4. Expected SOBP width

Energy (MeV)	SOBP width (cm)
250	10
180	8
100	2

As to the field radius  $r_1$ , we referred to the design parameters (Zheng et al. 2008) from M.D. Anderson Cancer Center, listed in Table 4-5. However, in NEU, the useful radius  $r_2$  is used as a parameter, rather than  $r_1$ . The value of  $r_2$  can be approximated. In Figure 4-7, the tangent of the scattering angle  $\theta_1$  is:

$$\tan(\theta_1) = \frac{r_1}{SSD} = \frac{r_2}{throw}, \quad (1)$$

so,

$$r_2 = \frac{r_1}{SSD} \cdot throw. \quad (2)$$

Table 4-5. Uncollimated field sizes

Field	radius $r_1$ (cm)
Large	Up to 17.7
Medium	Up to 12.75
Small	Up to 7.05

Considering the combinations of energy, SOBP-width, and field size, we designed nine double-scattering systems listed in Table 4-6.

Table 4-6. NEU-Case No. for different settings

Energy (MeV)	SOBP width (cm)	Field size	NEU-Case No.
250	10	Large	1
		Medium	2
		Small	3
180	8	Large	4
		Medium	5
		Small	6
100	2	Large	7
		Medium	8
		Small	9

The quantities related to SOBP are shown in Figure 4-8. AB is the proximal dose rise region; BC is the flat region (SOBP); CD is the decreasing distal dose region. The distance m100 is the projected distance between B and C, and m90' (or m90) is the width between points of 90% of the nominal “full” dose (average value in BC). The parameter d100 is the depth at C; d90' is the depth specified clinically; d80 is used to identify the range of protons. NEU analyzes the SOBP depth-dose distribution produced by S1 and S2 to find the values for these quantities, especially the three depths: middle of BC, C and B to calculate the transverse doses.

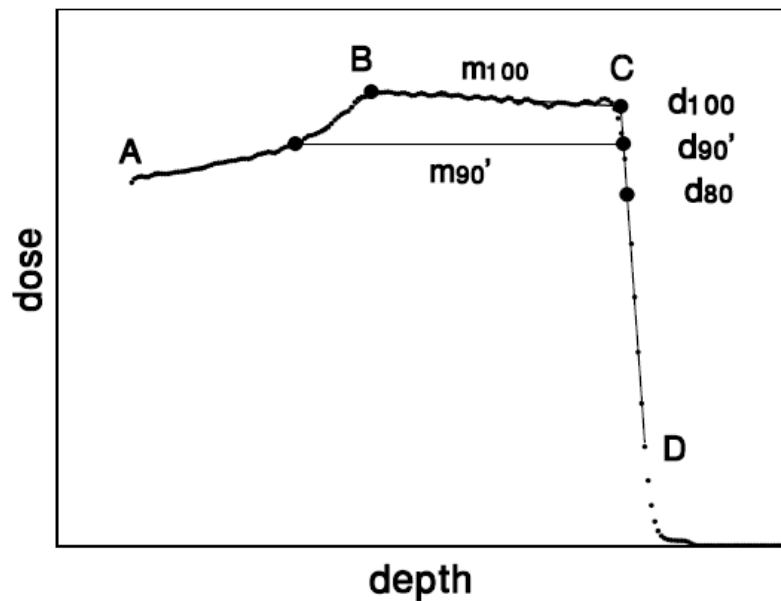


Figure 4-8. Various quantities on a SOBP curve

The default input file is “neu.inp.” Besides the parameters mentioned above, the pristine Bragg-peak data, such as “MDA250.BPK”, dose tolerance interval set at  $\pm 2.5\%$ , some fitting methods, and the number of scan points on the SOBP curve are also specified in it.

“BEAMPICT.INP” is an optional input file for the NEU code to show a picture of a schematic beam line for the current simulation. It provides some components possibly used in a real nozzle, and puts a hypothetical spherical tumor in the water phantom.

The input files for NEU-Case 1 are shown in Figure 4-9 and 4-10.

```

'DESIGN.MOD'          (filename or DESIGN).MOD
'DESIGN.CON'          (filename or DESIGN).(ANN or CON) or '
'MIXED.RET'          range-energy table in \BGWARE\DATA\
'MOLIERE'            MOLIERE or HIGHLAND scattering theory
'MDA250.BPK'         Bragg peak file in \BGWARE\DATA\
'MARQUARDT'         MARQUARDT or GRID or RANDOM
'NONE'              measured data file or 'NONE'
----- eight elements: mat'l, upstream z (cm), thickness (g/cm2) -----
'      '            -2   -999   1: precat mat'l,z,g/cm2 or blank
'      '            -999  -999   2: preabs ditto
'LEAD'              -999  -999   3: S1 A mat'l (simple scatterer)
'LEXAN'             0    -999   4: S1 B mat'l,z
'      '            -999  -999   5: postscat mat'l,z,g/cm2 or blank
'      '            -999  -999   6: postabs ditto
'LEXAN'             -999   .12   7: S2 A mat'l,z,MIN g/cm2
'LEAD'              50   -999   8: S2 B ditto (use for ANN)
----- major design parameters -----
322.4 250 0         throw (cm), energy (MeV), beam theta0 (mrad)
19.0 99 10         design radius (cm), d100, m100 (cmW)
2.5 -1 1          dose +/-%; step factor (- unlocks); cm/file unit
0                zoom (cm), added to z(1-6)
0 0 0            depth linear,quad coefft; transv quad coefft (%)
----- major design parameters -----
322.4 250 0         throw (cm), energy (MeV), beam theta0 (mrad)
19.0 99 10         design radius (cm), d100, m100 (cmW)
2.5 -1 1          dose +/-%; step factor (- unlocks); cm/file unit
0                zoom (cm), added to z(1-6)
0 0 0            depth linear,quad coefft; transv quad coefft (%)
----- scattering system -----
0.92 4 2 0        u0, # fit radii, # between, # passes
0 1E30 .07        spline y' at rMin,rMax; contour trim fraction
0 0 0 2           starting radii (max 4/line)
.8433 .6108 .2124 .2608 starting proj scat strengths (max 4/line)
.02 .02 .02 .02   parameter deltas for fit (0 means lock that one)
15 15 10000      Q steps QA->QB, PHI steps 0->pi, lookup table size
15 20 11         # steps, infty mult norm int; # pts for FOM calc
.5 99 .005       MGR: lambda,n/a,eps factor,quit,eps nrep,miss,eps
1.000 -0.007 1.80 mod: aa(j) = (p1 + p2*(j-1))*aa(j), cofM = p3
.001 5           z0 loop conv test (g/cm2 gm3,gm4), max # loops
----- Broken Spline fit to SOBP -----
201              # of scan points to compute (odd)
'T'              Terse or Verbose
'G'              fit method: GridParab, Marquardt, None
10 1E-8          passes, convergence on rms (%)
.5 10            reduction factor, initial lambda
.1 .0001         delta x,y (best may depend on fit method)
25 0 0          pts/segment (pristine peak)
10 2 2          ditto modulated (MAX pts for seg1)
8 1             deriv smoothing rms, # passes
.005 .1 .5      rms/avg < p1 -> 2 seg; distal cut; power
.05 .015        AB/AC < p1 -> 2 segs; rms/max > p2 -> add point
1 -.9 -.8       3 levels (neg -> horizontal fiducial line)
----- graphics -----
0 -.2 0         middle,deep,shallow scan depth corrections
-5 30           z limits modulator picture (absolute cm)
-5 2 -10 10     z,y limits 2nd scat picture (cm, z rel to zz(7))
0 40            z limits depth dose (cm rel to zz(8) = skin)
-40 40          x limits transverse scans (cm)
1 60 0          playback mod range, norm const

```

Figure 4-9. Contents of "neu.inp" for NEU-Case 1

```

'NOSKIP'          'SKIP' beam picture, else draw
'GRID'            'GRID' on picture, else none
0 100 25          picture x1,x2,y2
1 6 5             modulator x1,x2,y2
13 18 4          second scatterer x1,x2,y2
50               tank x1
-36 -26          depth-dose y1,y2
91 100           transverse dose x1,x2
33 37 5.0 10     apert x1,x2,radius,y2 (r>y2 -> no apt)
23 9 15          snout x1,y1,y2
22.4 5.0         target depth in tank,radius
10000 6 2        # points, power (x,y)
.001             fraction for projection test
2005            random seed

```

Figure 4-10. Contents of “BEAMPICT.INP” for NEU-Case 1

The output plots of “neu.inp” for NEU-Case 1 are shown in Figure 4-11. They were obtained by repeated modifications and simulations. The abscissas in all plots are in the unit of cm. Plot 1 is the schematic of S1, in which the horizontal length is the thickness of each step, and the vertical width represents the “weight” of that step. The sum of all weights is equal to 1. There are eleven steps in S1 for NEU-Case 1. Plot 2 is the cross-section of S2, in which the thickness and radius are marked in the unit of cm. The ordinates in plot 3 through plot 6 represent the dose values. The dose value at the depth of mid-SOBP point is normalized to 1. The high-dose regions are enclosed in the  $\pm 2.5\%$  tolerance interval marked by dashed lines. Plot 3 shows the formation of a SOBP curve by several “weighted” pristine Bragg-peak curves, in which the abscissa represents the depth in the water phantom. Plots 4 through 6 are the transverse dose

distributions at the scanning depths of proximal, middle and distal points on the SOBPs curve. The abscissas in these three plots represent the radial dimensions.

The beam line is shown in Figure 4-12. It was also obtained by repeated modifications and simulations. This figure is included only for demonstration, and it does not reveal the real geometrical size of each component. In addition, the density of dots standing for the values of dose is exaggerated to make it easy to be understood by the users.

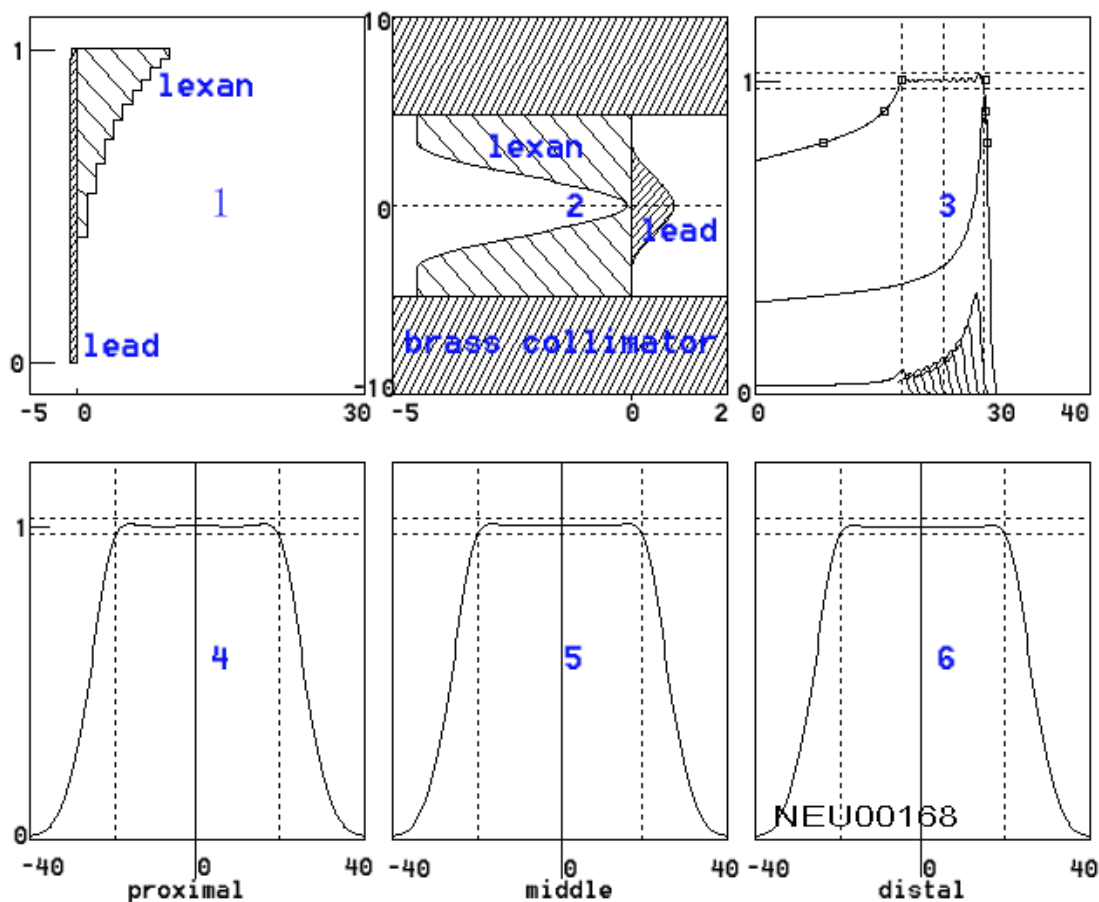


Figure 4-11. Output plots of “neu.inp” for NEU-Case 1

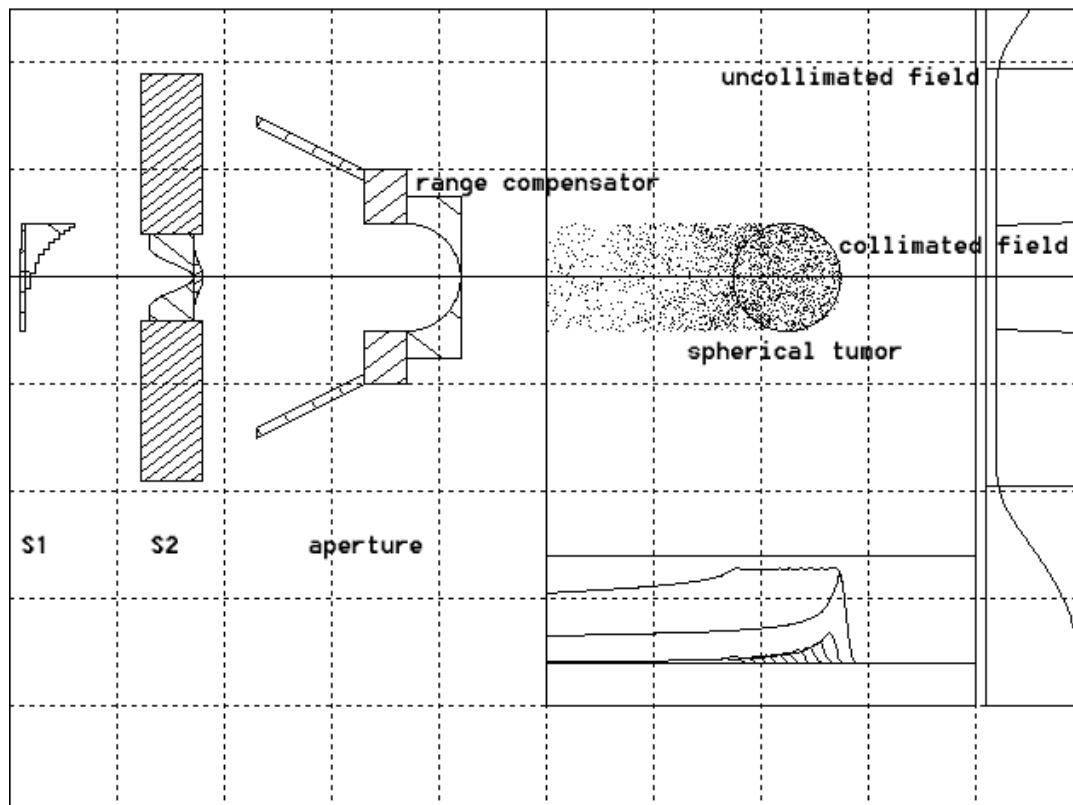


Figure 4-12. Output plot of “BEAMLIN.INP” for NEU-Case 1

One enlarged cross-section of the lead part in S2 for NEU-Case 1 is shown in Figure 4-13. The abscissa represents the design radius, and the ordinate represents the corresponding thickness of lead. There are ten design radii numbered from 1 to 10 by default, but only the first seven thicknesses have non-zero values. Hence, we can use seven laminated cone frustums marked by 1 to 7 to compose the lead part.



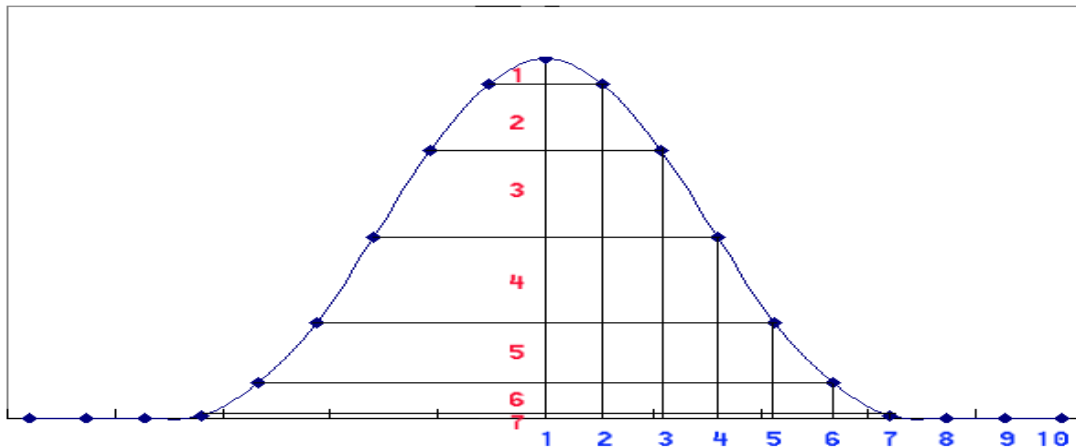


Figure 4-13. Cross-section of lead part in S2 for NEU-Case 1

The final satisfactory output parameters for nine NEU cases are listed in Table 4-7. The results for NEU-Case 5 and 9 are shown in Appendix A.

Table 4-7. The final parameters for 9 NEU cases

NEU-Case No.	Throw (cm)	Useful radius $r_2$ (cm)	Scattering strength of $S1(u_0)$	# of steps in S1	# of lead cells in S2
1	322.4	19.0	0.92	11	7
2	326.7	13.9	0.92	11	7
3	330.5	7.9	0.92	11	6
4	311.65	18.3	0.92	15	7
5	314.0	13.4	0.92	15	7
6	316.2	7.5	0.92	15	6
7	304.5	18	0.92	9	7
8	305.4	13	0.90	9	6
9	306.1	7.2	0.88	9	6

Before proceeding it is necessary to discuss some design tips and limitations of the NEU code. The most common problem with a double-scattering system is that S1 is

poorly matched to S2. The scattering strength of S1 ( $u_0$ ) plays a key role in the forming of the shape of the transverse dose distribution. Figure 4-14 shows a comparison of transverse dose distributions from different settings of the scattering strengths of S1 in NEU-Case 8. If the transverse dose curve looks ‘domed,’ it is because the scattering strength of S1 is too weak to scatter enough particles laterally. The solving method is to make S1 a little thicker to increase the scattering strength of S1, at a cost of penetration, or to move it farther from S2. On the other hand, if the transverse dose curve looks ‘dished,’ it is because the scattering strength of S1 is so strong that too many particles are scattered from the central line. The remedies are just opposite. One limitation of the NEU code is that it can be used to deal with only S1 and S2, without any additional objects or situations that are not cylindrically symmetric.

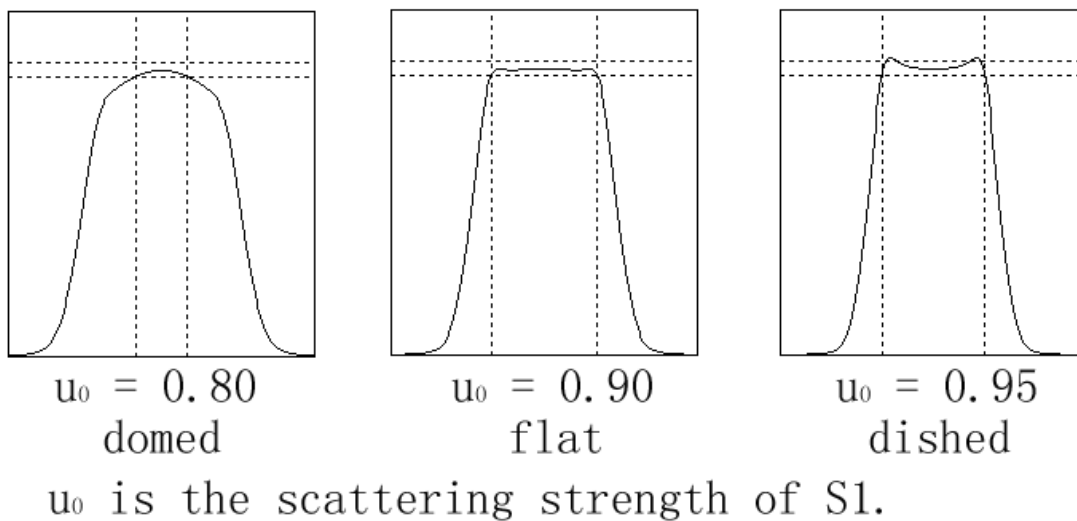


Figure 4-14. Comparison of transverse doses from different scattering strengths of S1

## 5. MONTE CARLO SIMULATION OF SIMPLIFIED NOZZLE

In this section, we will describe the whole simulation procedure, including modeling, transport of particles, results visualization, etc., for several nozzles with different settings. The purpose of Monte Carlo simulation is to verify the design results from Section 4.

### 5.1 MCNPX Input File for a Passive-Scattering Nozzle

The contents in an MCNPX input file for a passive nozzle is listed in Table 5-1. We used “mcnp\_pstudy” script to invoke MCNPX, so there was a block for “mcnp\_pstudy” input parameters.

Table 5-1. Contents in an MCNPX input file for a passive nozzle

Block No.	contents
1	“mcnp_pstudy” parameters
2	Cell cards to describe all the cells in the problem
3	Surface cards to describe all the surfaces
4	Data cards
4.1	Mode
4.2	Materials
4.3	Coordinate transformation
4.4	Void card to make some unused cells “void”
4.5	Source definition
4.6	Physics setting to specify the Energy ranges and Physics model
4.7	Tally cards to record energy deposition in water phantom, energy and angular distribution of protons incident on the front surface of water phantom
4.8	Mesh tally cards Particles flux and energy deposition in a designated mesh Multiplication factors for results
4.9	Running control cards

### 5.1.1 Geometry Models

The geometry model used in this research included two parts: a proton beam delivery system - a nozzle, and a dose measurement system - a water phantom.

The components of a nozzle have been described in Section 2, but the coordinate system in the MCNPX model was changed. The nozzle was aligned along the X-axis, and the size of water phantom was 40 cm × 80 cm × 80 cm, located at the isocenter. The components and the materials are listed in Table 5-2. To simulate a proton nozzle as close as possible to a clinical one, besides the scattering system, some shaping devices must be added. An outer shell of the nozzle was also needed, which was used to enclose the beam in a limited space. The thicknesses of dose monitors are usually very small, and their influence is minor, so they were omitted in the MCNPX model.

The materials listed in Table 5-2 are not the only available ones for a nozzle. For example, some vendors also use tungsten alloy or brass in first scatterers, and aluminum alloy is also an alternative for Lexan or ABS resin in the modulation wheels. Table 5-3 lists the composition of materials used in the MCNPX simulations and the CSDA ranges of 250 MeV protons (Berger et al. 2005).

Table 5-2. Components used in a typical passive-scattering-nozzle problem

Name	Designed by NEU	Illustrated by NEU	Modeled in MCNPX	Material
Vacuum window			Yes	
Profile monitor				
Reference monitor				
First scatterer	Yes	Yes	Yes	Lead
Range modulation wheel	Yes	Yes	Yes	Lexan
Second scatterer	Yes	Yes	Yes	Lead and Lexan
Range shifters			Yes	ABS resin
Collimators			Yes	Brass
Sub dose monitor				
Main dose monitor				
Final aperture		Yes	Yes	Brass
Range compensator		Yes	Yes	ABS resin
Shielding shell			Yes	Steel
water phantom		Yes	Yes	Water

Table 5-3. Composition of materials and CSDA ranges of 250 MeV protons

Material	Composition (weight fraction by percent or atomic fraction by number)	Density (g cm <sup>-3</sup> )	CSDA Range of 250 MeV Protons (g cm <sup>-2</sup> )	CSDA Range of 250 MeV Protons (cm)
Lead	Pb (100%)	11.35	76.69	6.76
Lexan	H (5.5%): C (75.6%): O (18.9%)	1.2	38.98(Lucite)	32.5
ABS resin	(C <sub>3</sub> H <sub>3</sub> N) <sub>2</sub> :(C <sub>4</sub> H <sub>6</sub> ) <sub>3</sub> :(C <sub>8</sub> H <sub>8</sub> ) <sub>5</sub>	1.04	37.94(H <sub>2</sub> O)	36.5
Brass	Cu (67%): Zn (33%)	8.35	56.62(Cu)	6.78
Steel	Fe(100%), other ingredients are omitted	7.86	54.54	6.94
air	N (75.6%): O (23.1%): Ar (1.3%)	0.001225	42.90	3.50E+04
water	H <sub>2</sub> O	1.0	37.94	37.94

In order to observe the influence of different components, three conditions were set for a nozzle, listed in Table 5-4. First, only S1 and S2 were included in a nozzle to

see the dose distribution from an uncollimated scattered beam. Second, a steel outer shell, several square collimators, and a final cylindrical aperture were added to see the dose distribution from a collimated broad beam. Third, a “hemi-spherical tumor” was assumed to be located at a depth in the water phantom, so a range shifter and a patient-specific range compensator were added to make the high-dose region “conformal” to the tumor. Notice that the shape of tumor in the MCNPX model was different from the spherical tumor demonstrated in the NEU model. The parameters of the beam-modifying devices and the radii of dose-recording regions for different field sizes are listed in Table 5-5.

Table 5-4. Components-in-nozzle conditions in a nozzle for MCNPX

Condition No.	Components used in a nozzle
1	S1, S2
2	All except range shifter and patient-specific range compensator
3	all

Table 5-5. Parameters for different field sizes

Field	Uncollimated radius of field (cm)	Area of square collimators (cm <sup>2</sup> )	Inner radius of the final aperture or the patient-specific range compensator (cm)	Radius of dose-recording region in the water phantom (cm)
Large	Up to 17.7	25×25	10	15
Medium	Up to 12.75	18×18	7	10
Small	Up to 7.05	10×10	4	6

In order to understand the different characteristics of the treatment beam, several types of tallies were used in MCNPX to obtain the desired quantities in selected regions in the water phantom. The tally-geometry conditions are listed in Table 5-6.

Figure 5-1 is a schematic of the rectangular mesh tally in the water phantom. Figure 5-2 is a schematic of the meshes used to score depth dose along the central axis (X-axis), and transverse doses at different depths. The whole model is cylindrically symmetric along the X-axis, so any symmetric layer along the X-axis can be used to score the transverse dose. Here, we used X-Z (fixed-Y) layer. Red meshes are used to score and show depth-dose distribution (SOBP-curve) along the central axis (X-axis), and blue meshes are used to score and show the transverse-dose distribution crossing the proximal, middle, and distal points on the SOBP-curve. The transverse layer in Figure 5-3 was used to show the contoured distribution of fluence or dose. Figure 5-4 is a general schematic of the scoring planes (or layers) in a mesh tally. The numbers of meshes shown in these figures are only used to illustrate the rectangular meshes, and they are not the real number of meshes used in the mesh tally.

Table 5-6. Tally-geometry conditions

Condition No.	tally region	Tally type	Quantity to score	Unit <sup>a</sup>
1	The front surface	F1 (E1, Fm1)	Current (Energy spectrum of incident protons)	# in the specified energy bin
		F11 (*C11, Fm11)	Current (Angular distribution spectrum of incident protons)	# in the specified angle bin
	Whole water phantom	F6 (Fm6)	Absorbed dose	MeV g <sup>-1</sup>
2 (Figure 5-2)	X: 300~340 Y: -0.5~0.5 Z: -39.5~39.5 The number of meshes is 80×1×79.	Mesh tally (type 1) Flux <sup>b</sup>	Fluence in a mesh	# cm <sup>-2</sup>
		Mesh tally (type 1) PEDEP <sup>c</sup>	Energy deposition in a mesh	MeV cm <sup>-3</sup>
3 <sup>d</sup> (Figure 5-3)	X: 321~321.5 Y: -39.5~39.5 Z: -39.5~39.5 The number of meshes is 1×79×79.	Mesh tally (type 1) Flux	Fluence in a mesh	# cm <sup>-2</sup>
		Mesh tally (type 1) PEDEP	Energy deposition in a mesh	MeV cm <sup>-3</sup>
4 (Figure 5-4)	The number of meshes is listed in Table 5-7.	Mesh tally (type 1) Flux	Fluence in a mesh	# cm <sup>-2</sup>
		Mesh tally (type 1) PEDEP	Energy deposition in a mesh	MeV cm <sup>-3</sup>
<sup>a</sup> The tally results are normalized to the results by one source particle. <sup>b</sup> In MCNPX, the result from “Flux” is “fluence” if omitting time. <sup>c</sup> The result from “PEDEP” (MeV cm <sup>-3</sup> ) equals to absorbed dose (MeV g <sup>-1</sup> ) if the medium is water. <sup>d</sup> Condition 3 is used to see the function of a single step of S1 (scattering part) alone and combined S1 (a single step of scattering part) and S2.				



Table 5-7. Parameters for mesh tally by collimated beams

Field size	Energy (MeV)	Axis	Coordinate range (cm)	Number of bins <sup>a</sup>
large	250	X	300 ~ 330	60
	180	X	300 ~ 320	40
	100	X	300 ~ 308	16
	All <sup>b</sup>	Y	-15 ~ 15	30
	All	Z	-15 ~ 15	30
medium	250	X	300 ~ 335	70
	180	X	300 ~ 325	50
	100	X	300 ~ 309	18
	All	Y	-10 ~ 10	20
	All	Z	-10 ~ 10	20
small	250	X	300 ~ 340	80
	180	X	300 ~ 325	50
	100	X	300 ~ 309	18
	All	Y	-6 ~ 6	12
	All	Z	-6 ~ 6	12

<sup>a</sup> All the meshes have the same size: 0.5 cm×1 cm×1 cm.  
<sup>b</sup> “All” represents all the energies of 100, 180 and 250 MeV.

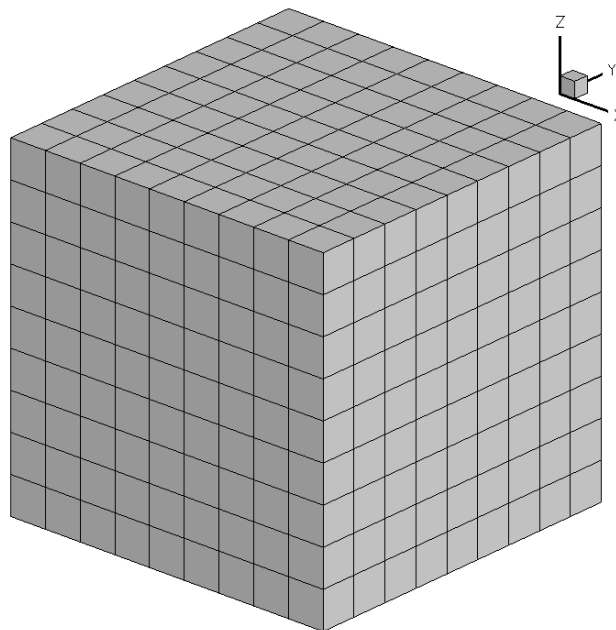


Figure 5-1. Illustration of the geometry for a rectangular mesh tally

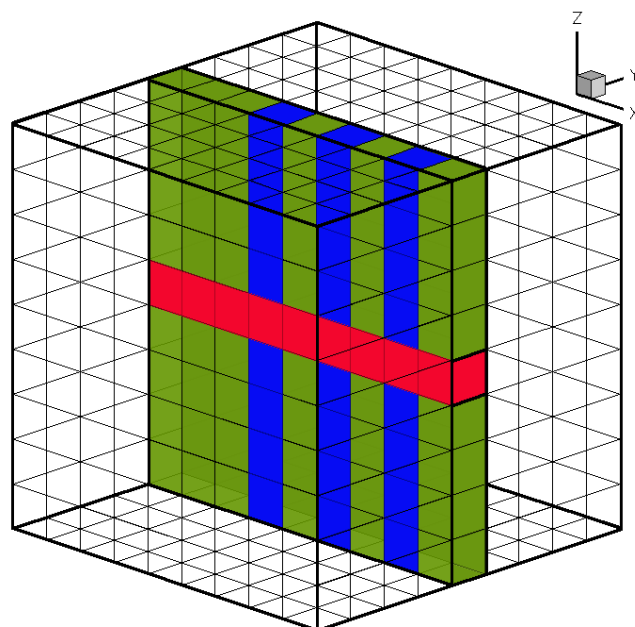


Figure 5-2. Fixed-Y layer for depth-dose and transverse dose

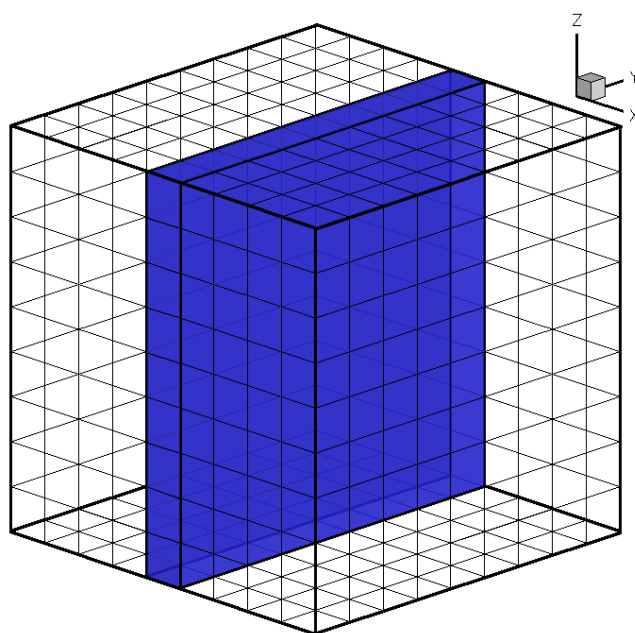


Figure 5-3. Fixed-X layer for transverse dose

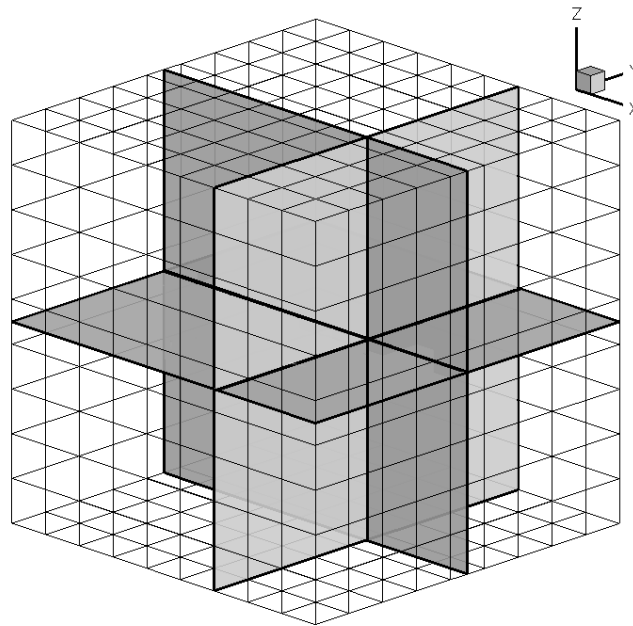


Figure 5-4. Planes to show dose or fluence distribution

Fifteen MCNPX cases were used to simulate the problems with different nozzle settings. A parallel-broad-beam case without a nozzle was also simulated, numbered by MCNPX-Case 0, in which the size of each mesh was  $2\text{ mm} \times 2\text{ mm} \times 2\text{ mm}$ . All the sixteen MCNPX cases are listed in Table 5-8.

Table 5-8. Cases simulated by MCNPX

Energy (MeV)	Components-in-nozzle condition No.	Field size	Tally-geometry condition No.	MCNPX-Case No.
100	None	Beam radius is 2 cm	Mesh: 2 mm×2 mm×2 mm	0
250	1	Large	1, 2, 3	1
		Medium	1, 2	2
		Small	1, 2	3
	2	Large	4	4
		Medium	4	5
		Small	4	6
	3	Large	4	7
		Medium	4	8
		Small	4	9
180	2	Large	4	10
		Medium	4	11
		Small	4	12
100	2	Large	4	13
		Medium	4	14
		Small	4	15

The distance between the source and the front surface of the water phantom, the center of which was defined as isocenter, was 300 cm. The parameters of S1 and S2 were obtained from the NEU-design results. The results for NEU-Case 1 (250 MeV, large field), used in MCNPX-Case 1, 4 and 7, are listed in Table 5-9. The modeling of S2 was the most difficult part in the whole modeling process. A series of superimposed “truncated right-angle” cones were used to compose the S2. The parameters of other components can be found in Appendix B.

Table 5-9. Parameters of S1 and S2 from NEU-Case 1

S1				S2		
Step No.	weight	Thickness (cm)		Radius (cm)	Thickness (cm)	
		Lead	Lexan		Lexan	Lead
1	0.4006	0.8829	0	0	0.1002	0.8731
2	0.1378	0.8679	0.9377	0.5331	0.4135	0.811
3	0.0965	0.8515	1.8827	1.0662	1.2238	0.6505
4	0.0739	0.8344	2.8317	1.5993	2.2867	0.4397
5	0.0608	0.8166	3.7842	2.1324	3.3258	0.2334
6	0.0513	0.7982	4.7402	2.6655	4.071	0.0853
7	0.0437	0.7791	5.6996	3.1986	4.4633	0.0073
8	0.0379	0.7593	6.6627	3.7317	4.5001	0
9	0.0343	0.7388	7.6294	4.2648	4.5001	0
10	0.0287	0.7176	8.6003	4.7979	4.5001	0
11	0.0344	0.6956	9.5754	To avoid the sharpness in the first cell, the radius of 0 is set to 0.001 in MCNPX input.		
To avoid the input error, if the thickness is 0, it is set to 0.0001 in MCNPX.						

### 5.1.2 Proton Beam Source Definition

As stated in Section 2, the production of secondary neutrons and recoil nuclei was not considered in this research. “Mode H” was used to indicate that protons were the only tracked particles. In MCNPX, there is no explicit generation of “delta-ray” knock-on electrons as trackable particles for heavy charged particles, and delta rays are produced only for electrons.

The particle-source description was based on the source at M.D. Anderson Cancer Center. One surface vertical to X-axis was set as the source plane. The spatial distribution of the proton source was described by two Gaussian distributions with a FWHM value of 0.54 cm in the Y-direction and 1.22 cm in the Z-direction. A “cookie-cutter” cell was defined to limit the position of the starting source particles to a radius of 3 cm around the beam axis. The energy of the source was also described by a Gaussian

distribution. The FWHM ranges from 0.23% to 0.3% for nominal energies from 250 MeV to 100 MeV (Titt et al. 2008). The FWHM for 180 MeV proton beam was assumed to be 0.27%. In MCNPX, the formats for the spatial and energy Gaussian distribution functions are different. For the spatial distribution, the input parameters are the mean value and the FWHM ( $2.35\sigma$ ), but for energy, the input parameters are the mean value and  $1.414\sigma$ . The parameters for the energy Gaussian distributions are listed in Table 5-10. The descriptions of the spatial and energy distributions in MCNPX are shown in Figure 5-5. The spatial distribution of the proton source (one thousand particles) is shown in Figure 5-6, in which the radius of the circle is 3 cm. The probability density function of energy (250 MeV) distribution is shown in Figure 5-7.

Table 5-10. Parameters for energy Gaussian distributions

Energy (MeV)	FWHM= $2.35\sigma$ (MeV)	$1.414\sigma$ (MeV)
100	0.3	0.184
180	0.486	0.292
250	0.575	0.346

```

sp1 -41 0.54 0      $Y position Gaussian distribution
sp2 -41 1.22 0      $Z position Gaussian distribution
sp3 -4 0.184 100     $Energy Gaussian distribution
sp3 -4 0.292 180     $Energy Gaussian distribution
sp3 -4 0.346 250     $Energy Gaussian distribution

```

Figure 5-5. The spatial and energy distribution functions in MCNPX



Figure 5-6. The spatial Gaussian distribution of proton source on Y-Z plane

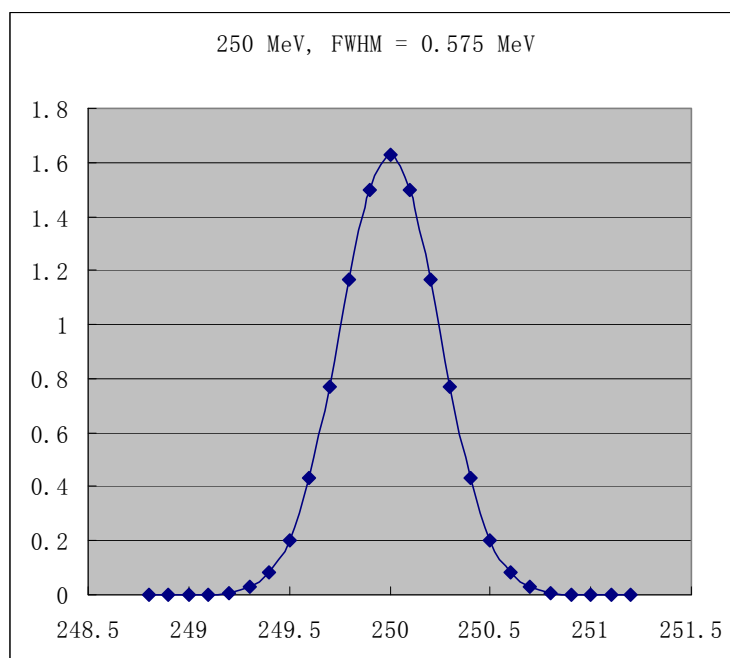


Figure 5-7. The energy Gaussian distribution (250 MeV)

However, if the spatial and energy distributions of the source are complex, i.e., both are Gaussian distributions, the sampling process of particles can require a large amount of time. To save time but not to affect the simulation results seriously, an ideal

beam source (the same beam as in NEU code) was used instead of the Gaussian-distributed source in all the cases with a nozzle.

### 5.1.3 Physics Settings

The “phys:h” card can be used to control some physics settings in proton-transport problems. The default upper energy limit is 100 MeV for protons. It must be increased to a higher value to simulate problems with 250 MeV protons. The parameter “emax” was changed to “300 MeV” to cover all the problems to be simulated. For the “table-based physics cutoff”, the “mix and match” option was still kept by default to assure either tables or models can be used when available. For protons, “la150h” is the data library containing the data tables for several isotopes. In addition, the “Vavilov” model, which is the default, was kept to control charged-particle straggling, rather than “CSDA”, and the production of recoil light ions was inhibited by default.

The “cut:h” card can be used to set the energy cutoff. The default lower energy cutoff for protons is 1 MeV. This cutoff was retained in MCNPX-Case 0. In other cases, to save the computation time, the cutoff was increased to 20 MeV, within a range of 4.3 mm in water, which was shorter than the maximum size of a mesh (5 mm) in mesh tally.

In MCNPX, five cards (LCA, LCB, LCC, LEA and LEB) can be used to control physics parameters for the “Bertini”, “ISABEL”, “CEM03”, “INCL4” and “FLUKA” options (the version of FLUKA in MCNPX is kept for legacy purposes.). In this research, only “CEM03” or “INCL4” models can be chosen to run the problem correctly. The problems with “Bertini” or “ISABEL” models aborted in the running in this research.



The reason for failure is still unclear. Simulation results from the “CEM03” and “INCL4” models had a minor discrepancy, while the problems with “CEM03” took less time, so the “CEM03” model was adopted finally. Only the LCA, LCB and LEA cards were used in this research.

#### 5.1.4 Tally Types

As given in Table 5-1 and Table 5-6, in this research the F1 tally was used to record the current of particles across a surface. The F1 tally and energy bin (E1) were used to obtain the energy spectrum, and the F1 and angle bin (\*C1) were used to obtain the angular distribution of the protons incident on the front surface of water phantom. The F6 tally (energy deposition in unit mass) was added also to record the average absorbed dose in the water phantom.

To obtain the “cell-wise” fluence and dose distributions, we used Mesh tally (type 1) in MCNPX. Mesh tally is a method to score the quantities of interest, such as fluence, energy deposition, and number of tracks in the volume of the mesh. One mesh can contain several different materials, so the mesh tally result is not averaged over the mass of the mesh, but over the volume of the mesh. There are three types of mesh geometries: rectangular mesh, cylindrical mesh and spherical mesh. A rectangular-mesh geometry built on the water phantom was used in this research.

Actually, to obtain unbiased tally results, we set the number of steps in S1 as the multiplication factor for all the tallies by using Fm cards. The reason will be explained in Section 5.4.

### 5.1.5 NPS Settings

In a mesh tally, the continuity of the results among adjacent meshes and the precision of results are strongly dependent on the size of the mesh. If the mesh is set too fine, a large NPS will be required to meet the precision requirement. If the mesh is set too coarse, it is not necessary to set NPS so large, but the smoothness of the dose or fluence distributions curve will be unacceptable, and the values on a contour line will vibrate seriously.

Several experiments were simulated with different choices of the mesh size and the NPS to observe the effects. Considering the computation time and precision of the results, finally, the mesh size was set to 0.5 cm×1 cm×1 cm in most cases. The NPS was set to  $10^7$  in most cases to make the tally results in the regions of interest meet the requirement of 10% uncertainty in statistics. In MCNPX-Case 1, the mesh size was set to 2 mm×2 mm×2 mm, and NPS was set to  $10^8$ .

### 5.2 Mcnp\_pstudy Parameters and Execution

An MCNP(X) (MCNP or MCNPX) input file, which includes “symbolic” parameters and real values, should be prepared before the execution of “mcnp\_pstudy” script. The parameter line should begin with “c @@@” and the options line should begin with “c @@@ OPTIONS”. The MCNP(X) input parameters that would use the “mcnp\_pstudy” variables should be the corresponding symbolic variables defined in the parameter lines for “mcnp\_pstudy”.

Before the execution of this script, the user should confirm if “Perl” is installed under the UNIX system. The user can use the command “which perl” to check it, and usually the default location is “/usr/bin/perl.” If Perl cannot be found, it must be installed before using “mcnp\_pstudy”. The execution command of “mcnp\_pstudy” is “perl mcnp\_pstudy -i inputfile.” Several other optional parameters can be added after this command or specified in an input file.

The execution process of “mcnp\_pstudy” includes three stages: setup, invoking and data collection. If “-setup” is specified, a series of case directories will be created and the real values are assigned to the symbolic parameters to produce one real MCNP(X) input file named “inp” in each case directory. If “-run” is specified, an MCNP(X) command will be invoked to execute all the “inp” files successively. If “-collect” is specified, the data will be collected and the average tally results and relative errors will be created and shown. Currently, the specification of “-collect” is invalid to the mesh tally results in MCNPX but valid to the tallies in MCNP.

### 5.3 Visualization of Models

The purpose of visualization of these models is to find the errors in the geometry and to observe the tracks of particles.

In each MCNPX case, one input file in a sub-directory was chosen randomly as the input file for Visual Editor of MCNPX to see the geometries and tracks. Sub-case 5 in MCNPX-Case 7 was taken as an example. The 3D model is shown in Figure 5-8. Due to a deficiency of Visual Editor, the final aperture and patient-specific compensator

cannot be shown in the 3D model. The 2D cross sectional profiles of different components are shown in Figure 5-9 through Figure 5-12. The fifth step in S1 included two parts: a scattering foil and a range modulation part. The bi-material contour-shaped S2 can be seen clearly. The collimator for S2 was a cylindrical ring surrounding S2, but the other three collimators are rectangular frames. The final aperture is a rectangular solid with a cylindrical hole. The patient-specific compensator is set to be a rectangular solid with a subtraction of a hemisphere. The compositions of the components were consistent with the data in Table 5-3.

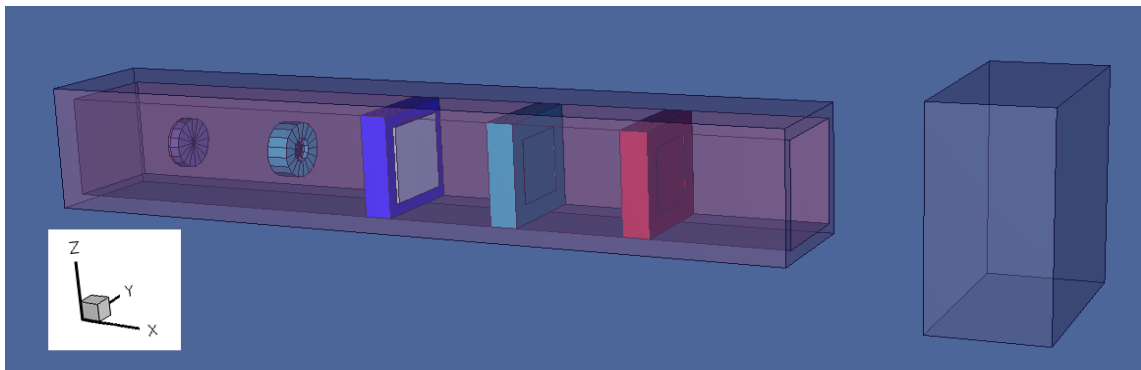


Figure 5-8. 3D model of a passive-scattering nozzle in MCNPX-Case 7

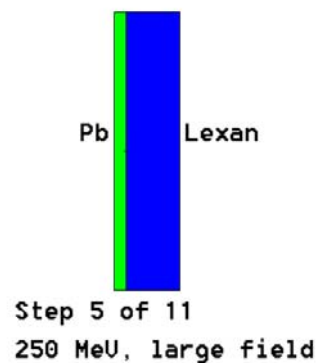


Figure 5-9. Schematic of one step on S1

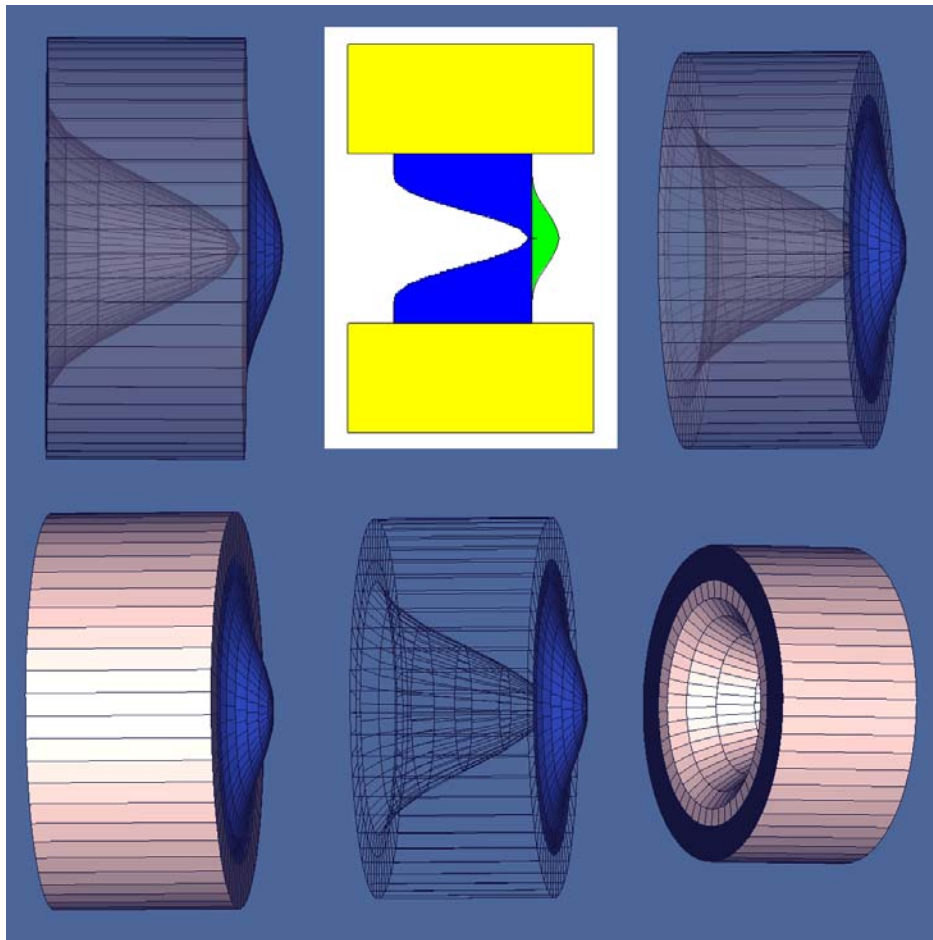


Figure 5-10. Schematic of a contour-shaped S2

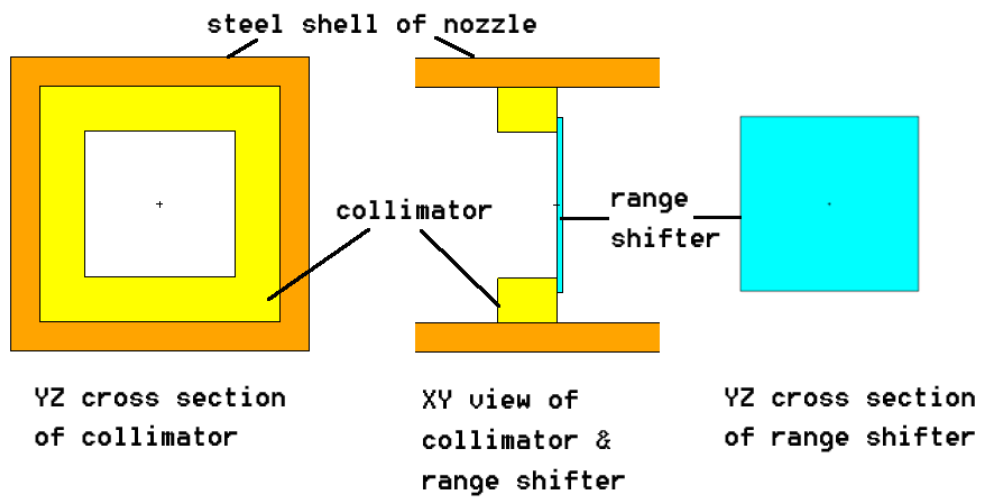


Figure 5-11. Schematic of collimator and range shifter

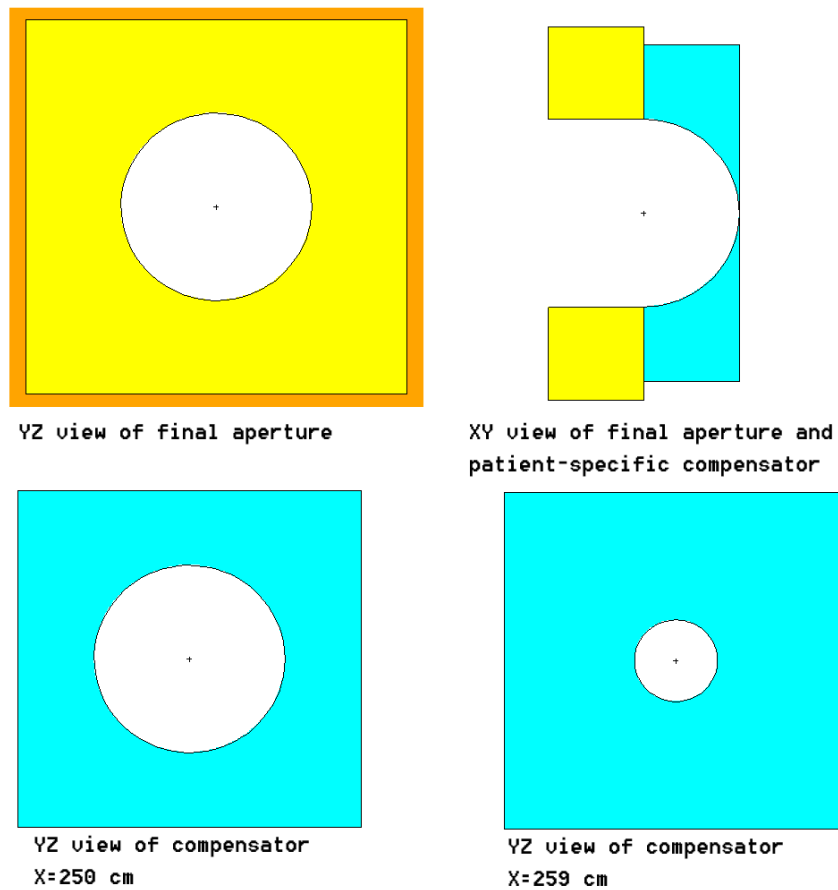


Figure 5-12. Schematic of final aperture and patient-specific range compensator

The tracks of proton in MCNPX-Case 1, 4 and 7 were plotted by Visual Editor and are shown in Figures 5-13, 5-14 and 5-15. The region in white represents a “void” region, in which protons can pass through without any interactions (This applies to all the figures with particle tracks). Track changes are clearly seen in the different cases with different geometry settings. In Figure 5-14, the lateral profile of the tracks is “collimated” comparing with the “uncollimated” beam in Figure 5-13. The laterally “collimated” beam is “shaped” to a hemispherical profile longitudinally by a range compensator in Figure 5-15.

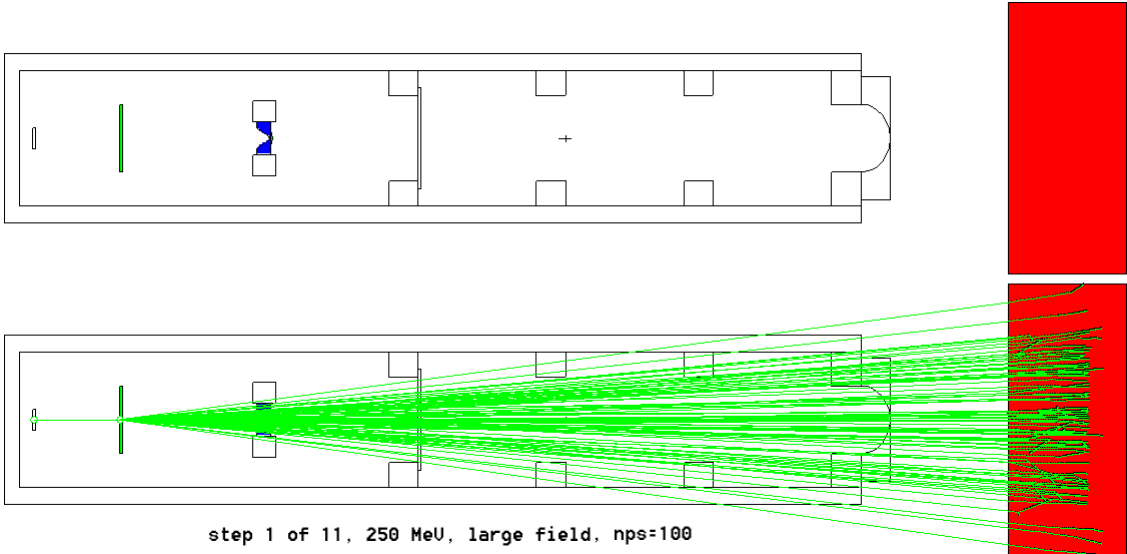


Figure 5-13. Nozzle geometry and proton tracks in MCNPX-Case 1

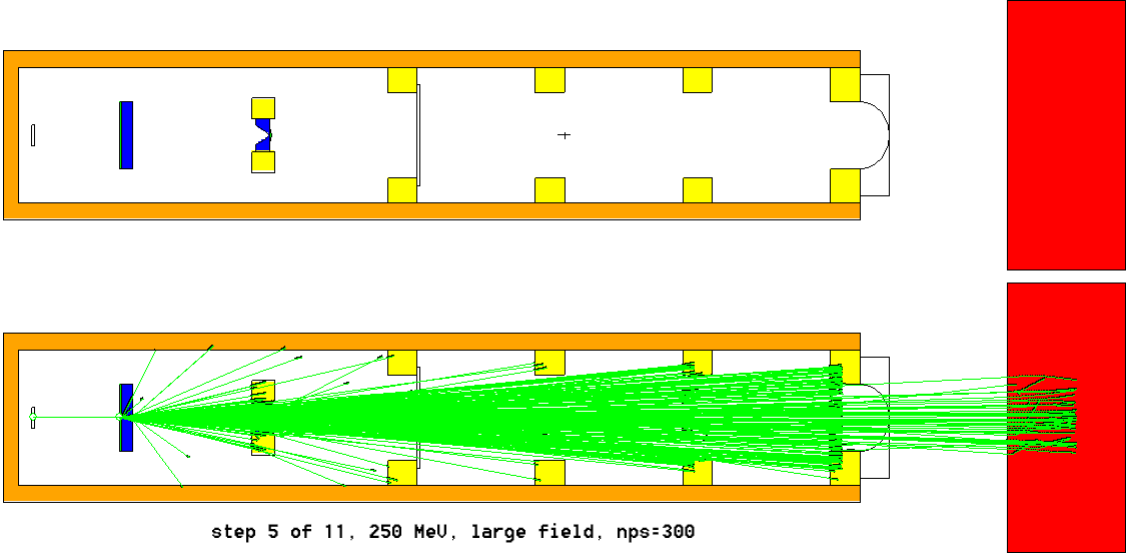


Figure 5-14. Nozzle geometry and proton tracks in MCNPX-Case 4

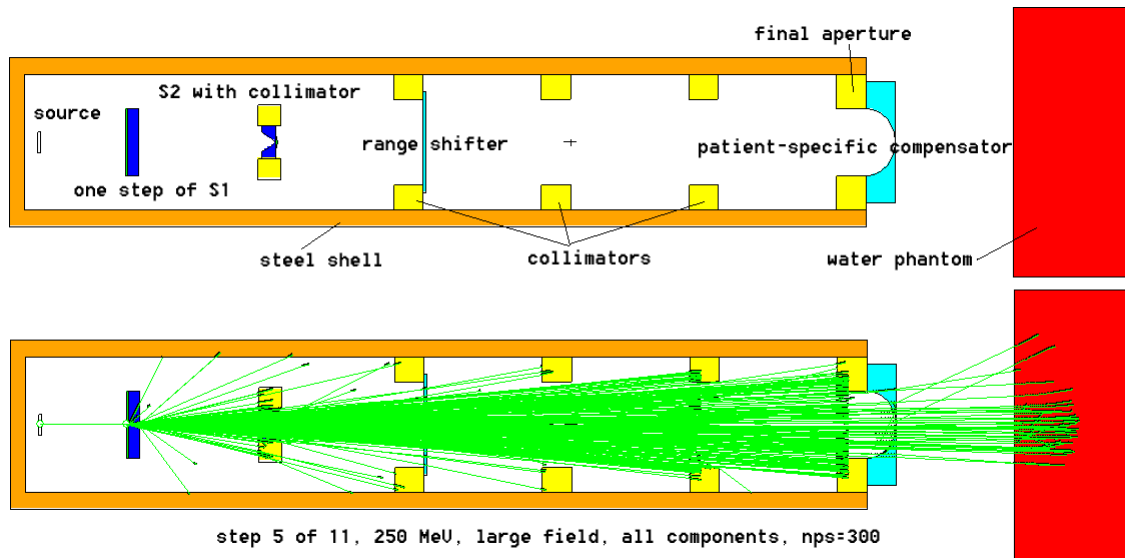


Figure 5-15. Nozzle geometry and proton tracks in MCNPX-Case 7

#### 5.4 Execution of Merge\_metal

As mentioned previously, in a problem including an N-step range modulation wheel, the problem is separated into N cases, so the expected result from all the cases should be:

$$\bar{D} = \sum_{i=1}^N W_i D_i, \quad (3)$$

where  $W_i$  is the weight of the  $i_{\text{th}}$  step;  $D_i$  is the corresponding score, such as fluence or dose;  $W_i D_i$  is the weighted score from the  $i_{\text{th}}$  step; N is the total steps in a

wheel; the known condition is  $\sum_{i=1}^N W_i = 1$ .



The function of “merge\_mctal” script is to sum the “mctal” data from N cases and average the sum by N to create a new “mctal” data file. If so, the results from “merge\_mctal” will be:

$$D_{merge} = \frac{1}{N} \sum_{i=1}^N W_i D_i = \frac{1}{N} \bar{D}, \quad (4)$$

which is not the expected result. The expected result has been reduced by a factor of  $\frac{1}{N}$ . If a multiplication factor N is set for each tally, the  $i_{th}$  result for the  $i_{th}$  case will be

$W_i D_i N$ . The result from “merge\_mctal” will be:

$$\frac{1}{N} \sum_{i=1}^N W_i D_i N = \sum_{i=1}^N W_i D_i = \bar{D}, \quad (5)$$

which is the expected result. Hence, to obtain unbiased results, a multiplication factor N must be set in Fm cards for all the tallies. For problems whose results did not need to be merged by “merge\_mctal”, this multiplication factor was not set.

After execution of “mcnp\_pstudy,” the tally results for each sub-case are stored in a UNIX-format file “mctal.” All the “mctal” files were renamed with successive numbers, such as “mctal01”, “mctal02”, etc; copied to a folder named “merge” containing “merge\_mctal” and an executable script “mergecsh” including the command to execute “merge\_mctal.”

Because the alignment format of the contents in a “mctal” file from MCNPX is different from the one from MCNP, “merge\_mctal” cannot be directly used to merge “mctal” files from MCNPX. The codes on the 368<sup>th</sup> line of the source code of “merge\_mctal” revealed that “merge\_mctal” did not read the parameters after “f” for F5

tally. In addition, the format of source particle types was the main difference between MCNP and MCNPX. Hence, the solving method was to change the particle types to MCNP format and modify the mesh tally in MCNPX to “F5” tally.

After the execution of “merge\_mctal,” a new “mctal” file was produced. To make this new “mctal” file to be read in by MCNPX, it must be changed back to MCNPX format. The method was to replace the first paragraph with the contents from an original “mctal” file from MCNPX.

## 5.5 Simulation Results

Several methods can be used to view the simulation results. The first is to execute the command “mcnpx z” in the directory containing the “mctal” file to view the results graphically. The second method is to read in the data by a “VB script embedded in Excel” edited by the author of this thesis. Before using VB script, the “mctal” file must be converted to “DOS” format from “UNIX” format. The third method is to invoke one subroutine of MCNPX - “gridconv” to convert the “mdata” file (a binary file storing the mesh tally data) to an ASCII file compatible with several external graphics packages, such as “PAW”, “IDL”, “Tecplot” or “GNU PLOT.” The packages “IDL” and “Tecplot” were preferred because the formats for these packages are more readable.

### 5.5.1 Results No. 1

The first set of results came from MCNPX-Case 0. Only a broad beam and a water phantom were provided. The proton energy was 100 MeV, and the beam radius was 2 cm. The contour depth-dose distribution on the central layer and the contour transverse dose distribution at a depth of 5 cm are shown in Figure 5-16 and Figure 5-17. The depth-dose distribution along the central axis (X-axis) and the transverse dose distribution at a depth of 7.6 cm along the Y-axis are shown in Figure 5-18 and Figure 5-19. These results are consistent with the theoretical expectations qualitatively.

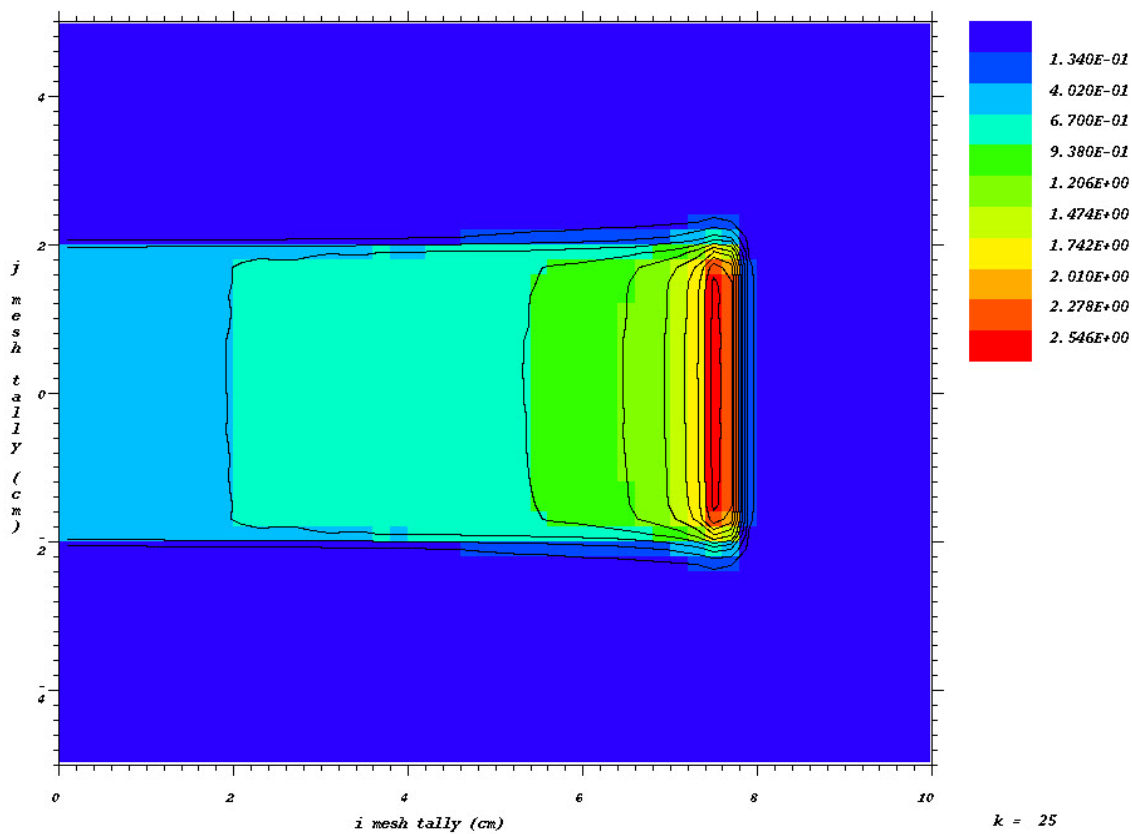


Figure 5-16. Contour depth-dose distribution on the central layer

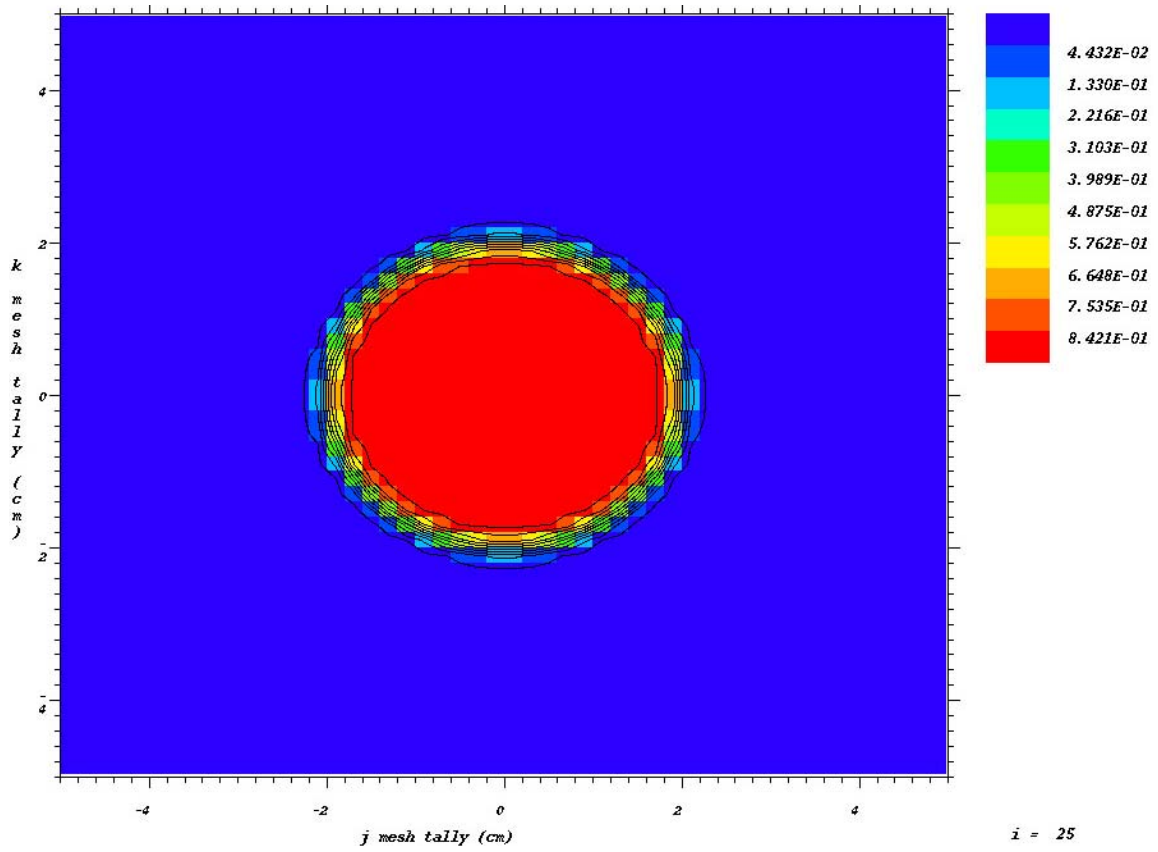


Figure 5-17. Contour transverse dose distribution at the depth of 5 cm

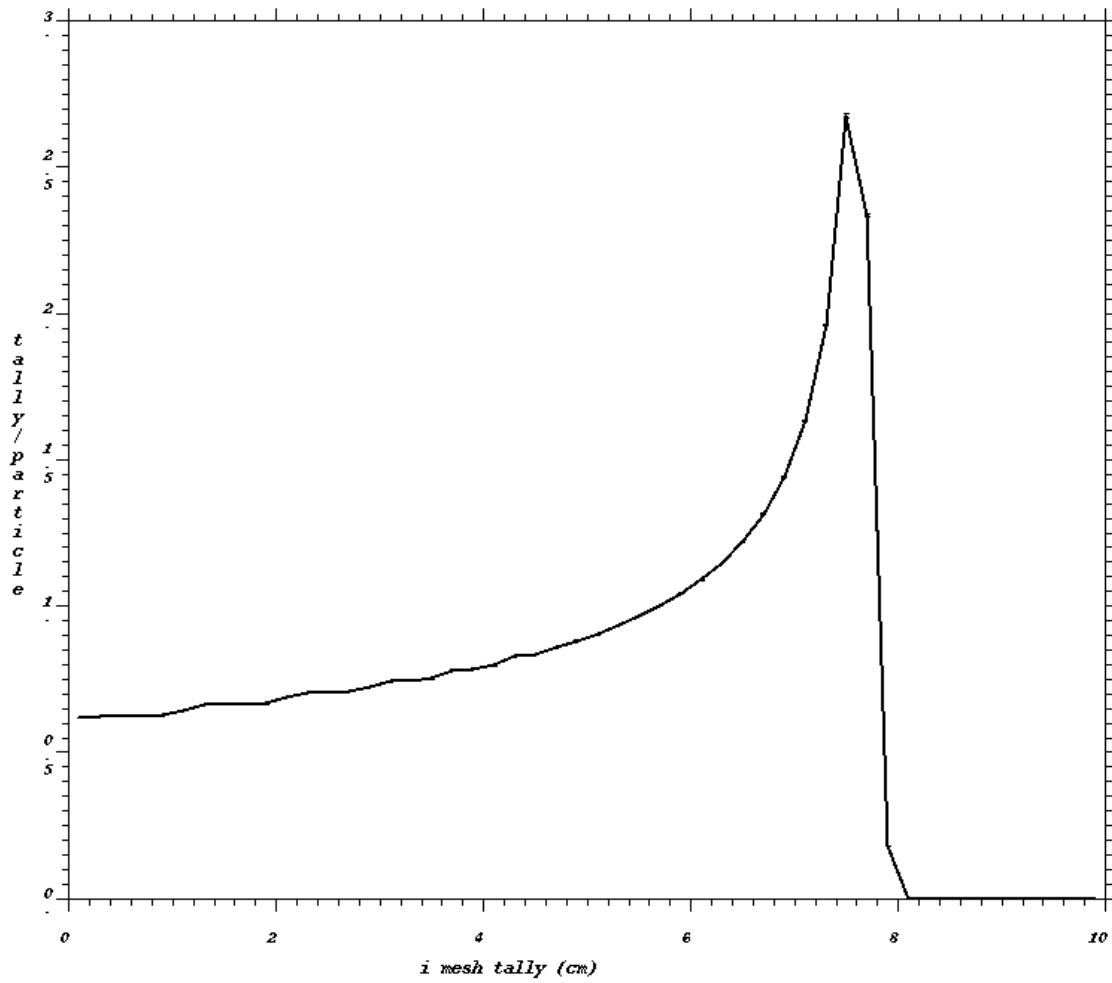


Figure 5-18. The depth-dose distribution along the central axis (X-axis)

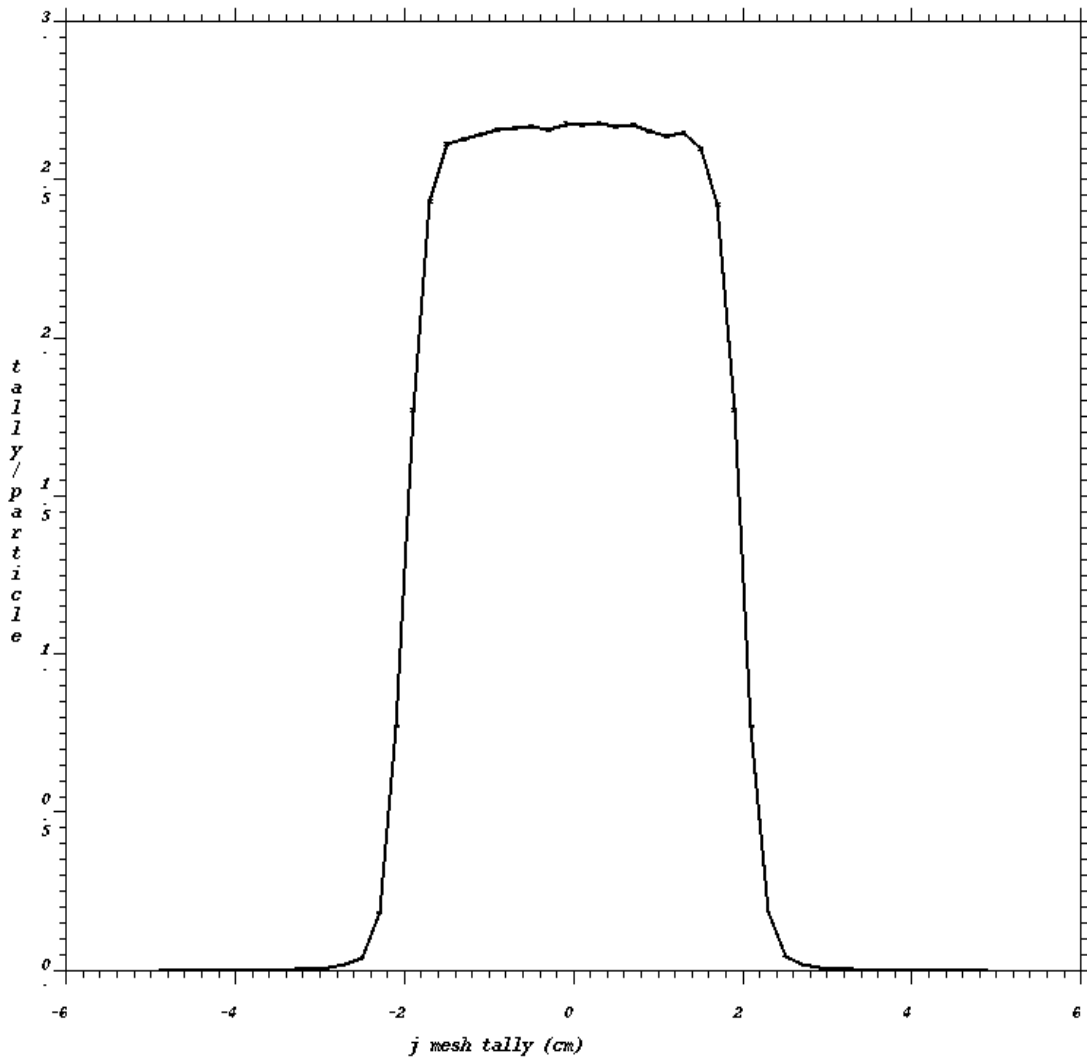


Figure 5-19. The transverse dose distribution at the depth of 7.6 cm along the Y-axis

### 5.5.2 Results No. 2

The second set of results came from MCNPX-Case 1 with tally-geometry condition No. 3. A narrow beam with an energy of 250 MeV and the first step of S1 (scattering part) were provided. MCNPX and “gridconv” were invoked to deal with the data. The converted data from “gridconv” were plotted using Excel in 2D or 3D mode.

The geometry and tracks are shown in Figure 5-20. The contour transverse dose distribution (X: 321 ~ 321.5 cm) from MCNPX is shown in Figure 5-21. The results plotted using Excel are shown in Figure 5-22 in 2D mode and Figure 5-23 in 3D mode. The transverse dose along the Y-axis (X: 321 ~ 321.5 cm, Z: -0.25 ~ 0.25 cm) from MCNPX is shown in Figure 5-24.

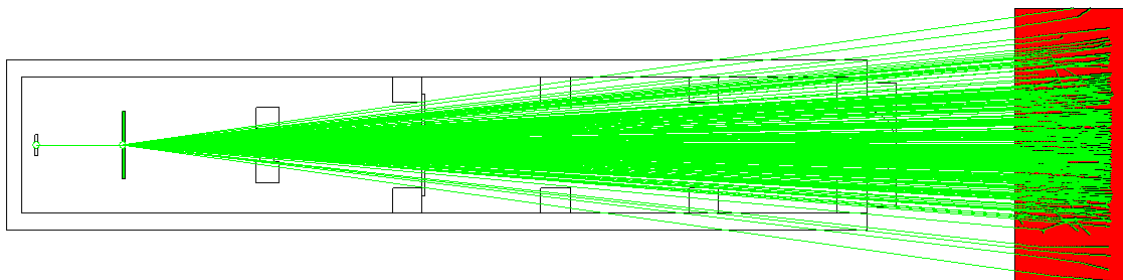


Figure 5-20. Geometry and tracks of protons scattered by a scattering foil

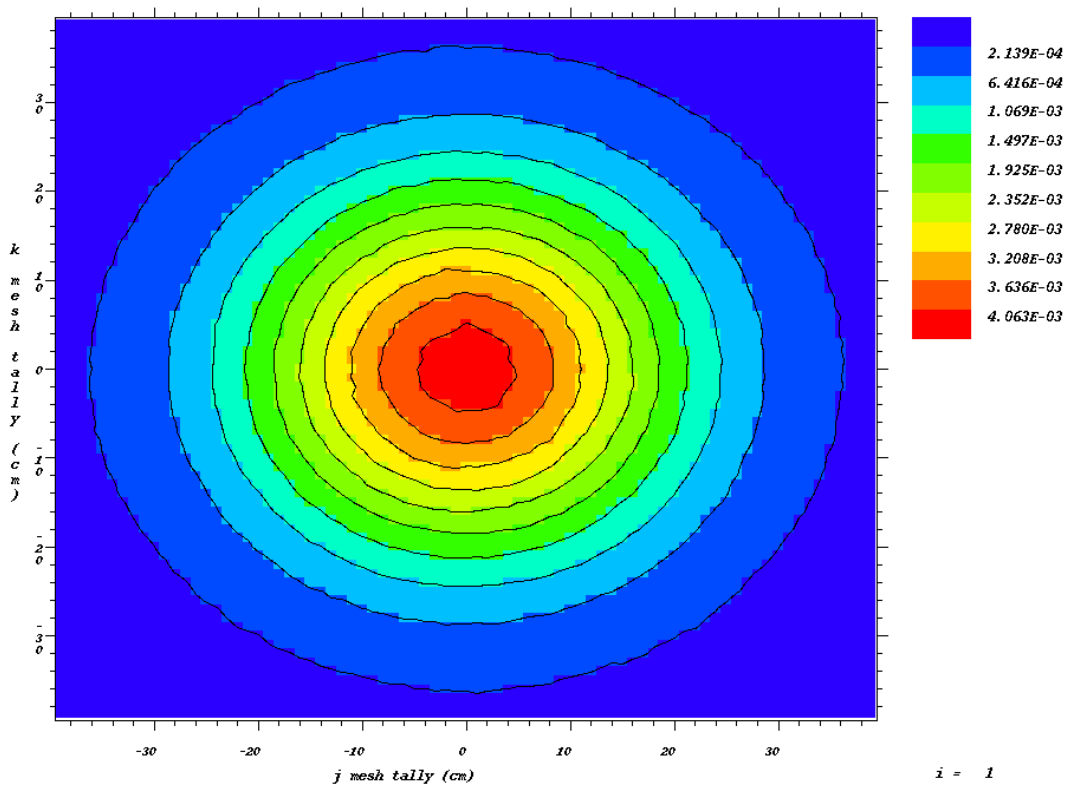


Figure 5-21. The contour transverse dose distribution for Results No. 2

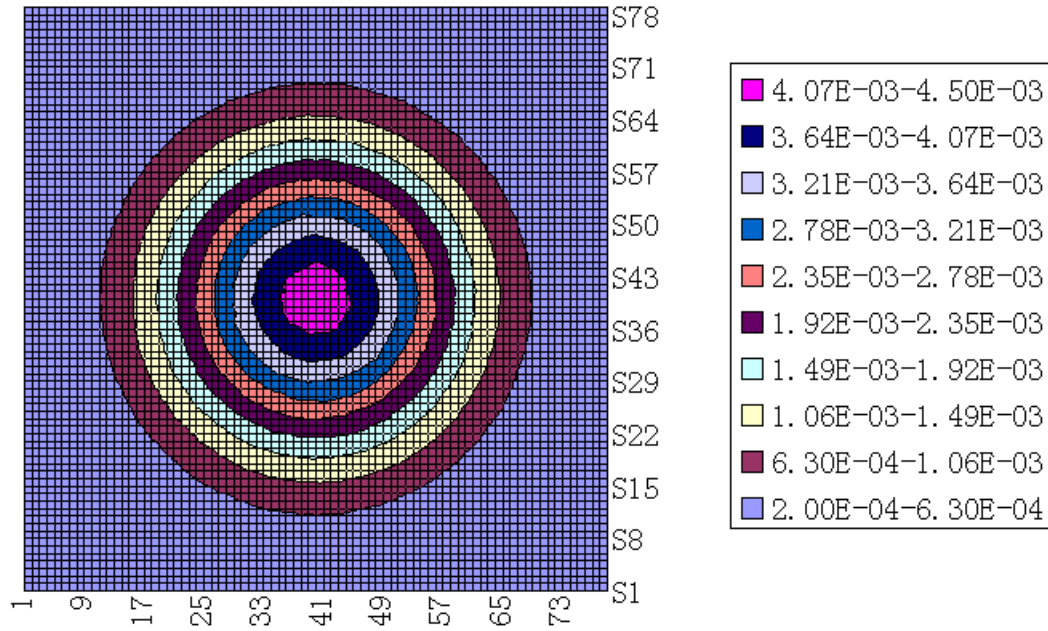


Figure 5-22. The contour transverse dose distribution for Results No. 2 in 2D mode

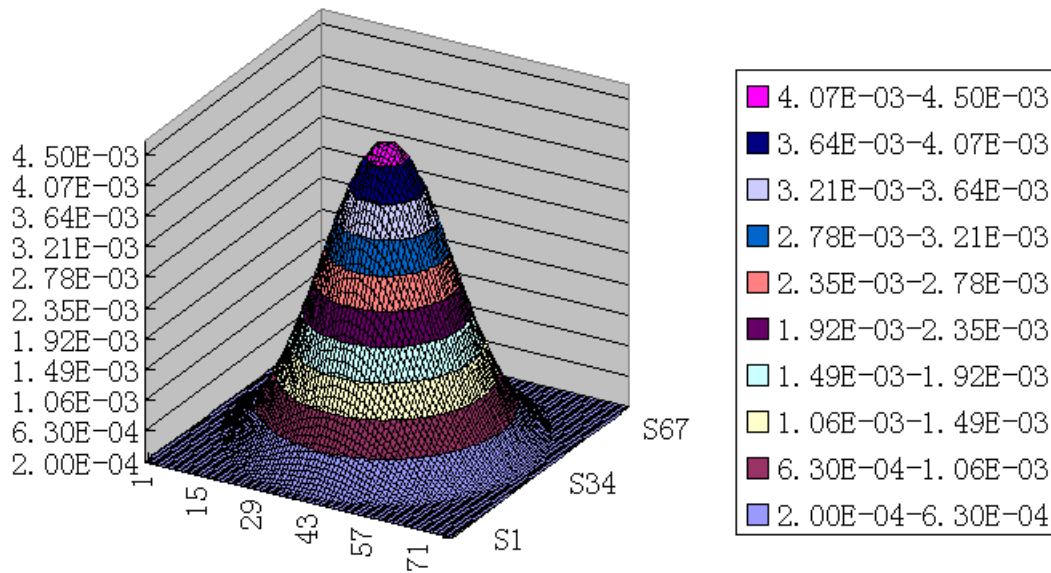


Figure 5-23. The contour transverse dose distribution for Results No. 2 in 3D mode



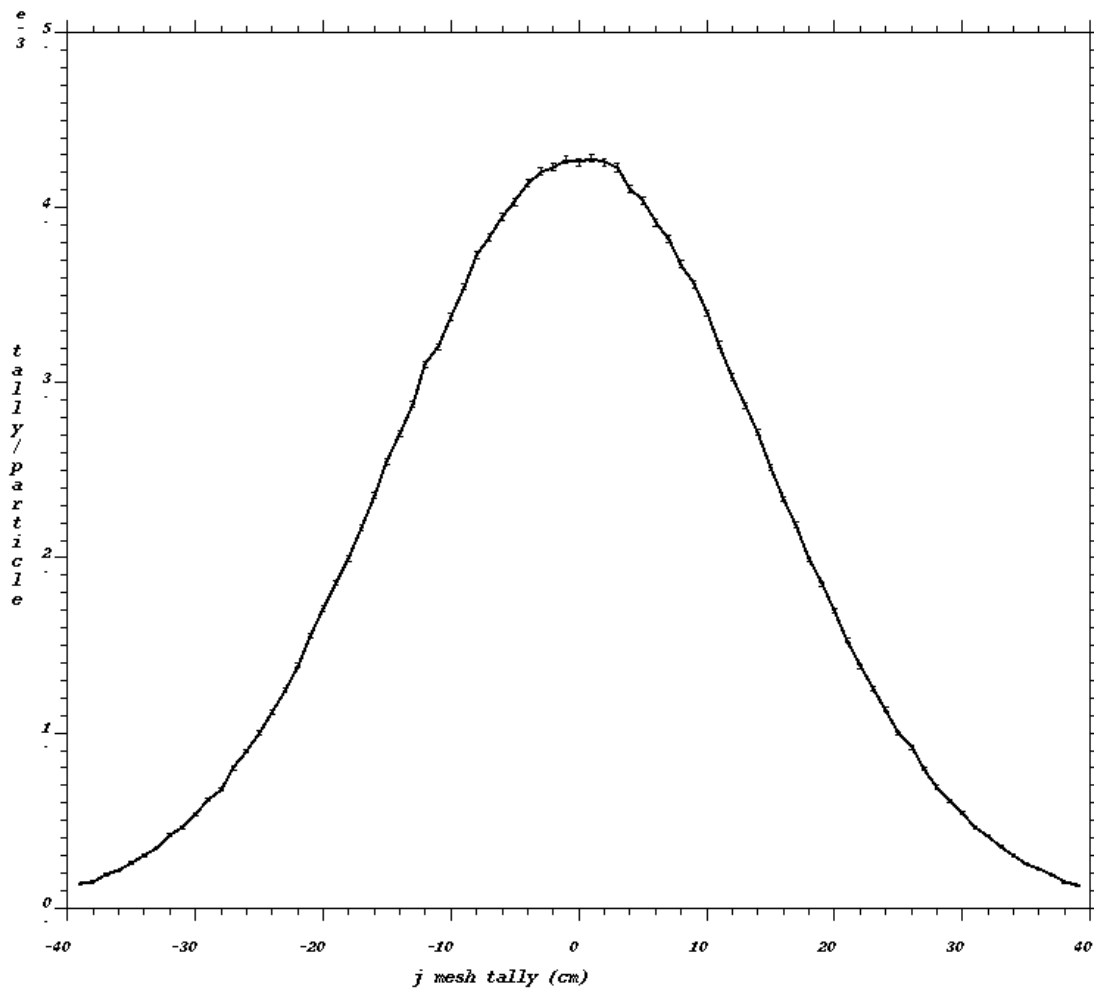


Figure 5-24. The transverse dose along the Y-axis for Results No. 2

From these figures, it was determined that the transverse dose distribution had a Gaussian shape rather than a uniform-spreading shape if only a single scattering foil was used. The high-dose region from this scattered beam was so narrow that this beam possibly can be used to only treat tumors with small sizes.

### 5.5.3 Results No. 3

Based on the settings for the second set of results, an S2 was added to produce the third simulation results. The geometry and tracks are shown in Figure 5-25. The contour transverse dose distribution (X: 321 ~ 321.5 cm) is shown in Figure 5-26. The transverse dose along the Y-axis (X: 321 ~ 321.5 cm, Z: -0.25 ~ 0.25 cm) is shown in Figure 5-27.

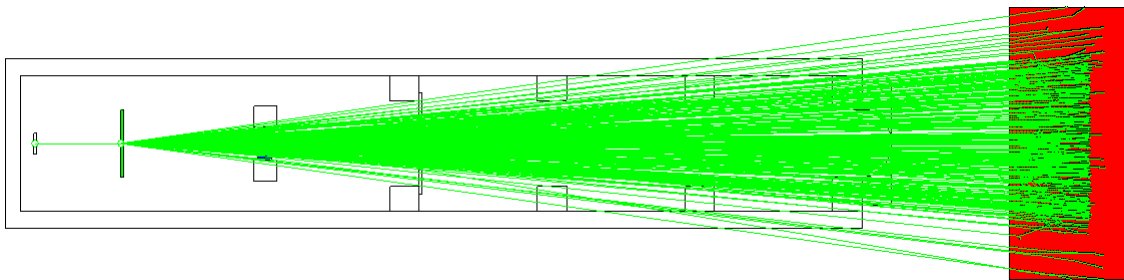


Figure 5-25. Geometry and tracks of protons scattered by a scattering foil and S2

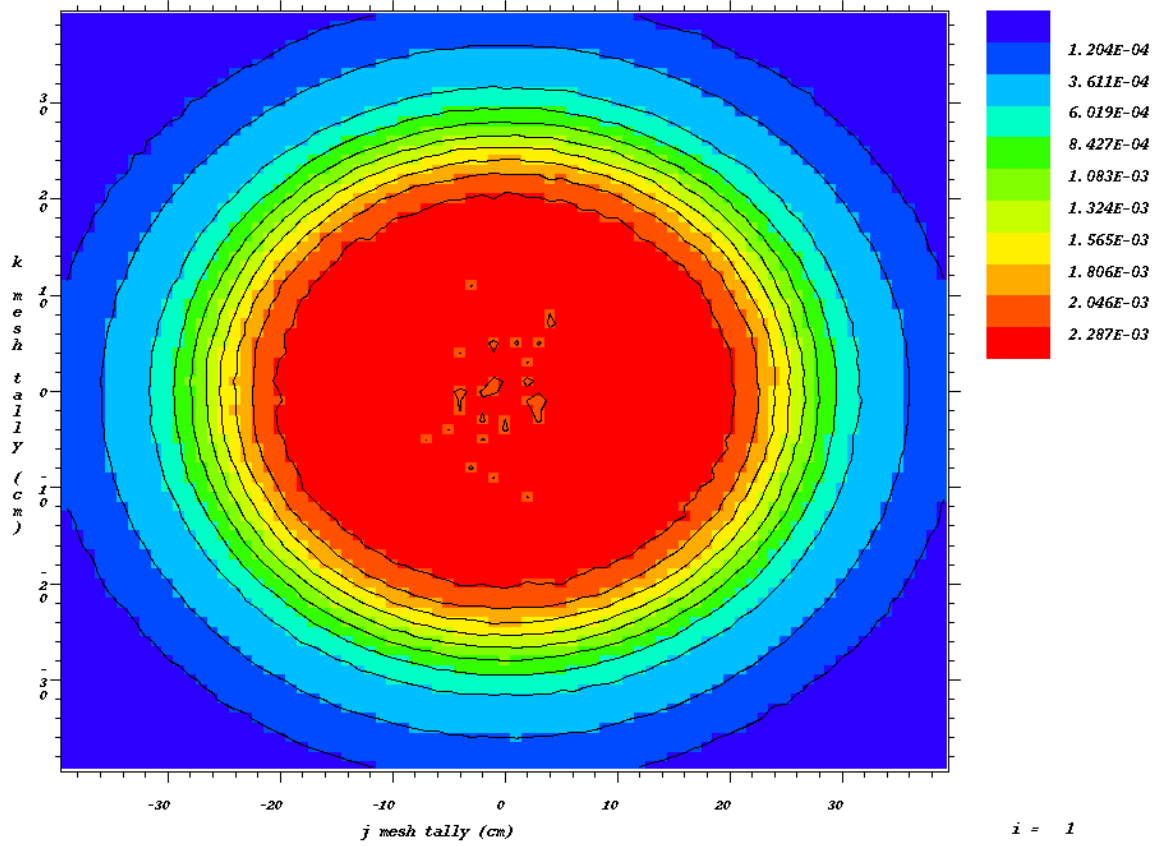


Figure 5-26. The contour transverse dose distribution for Results No. 3

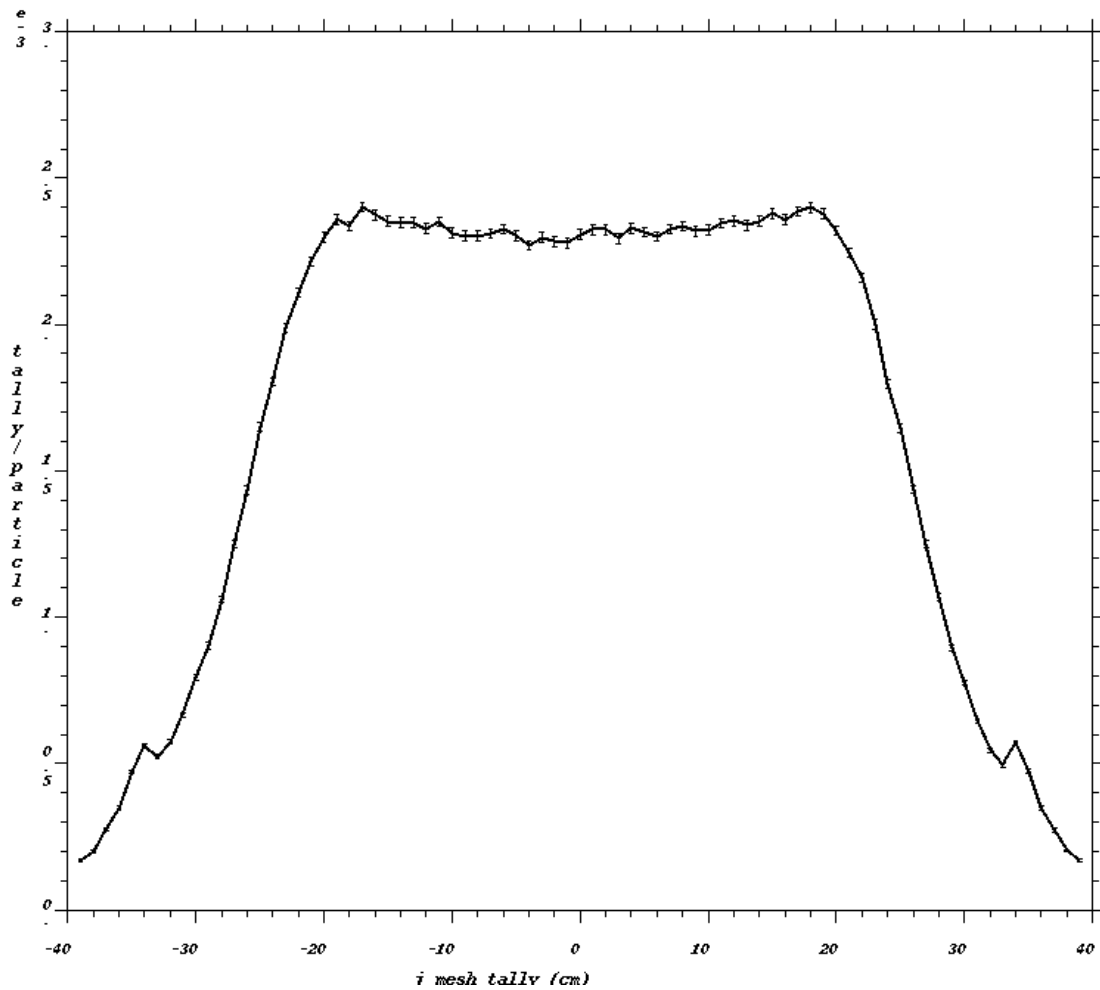


Figure 5-27. The transverse dose along the Y-axis for Results No. 3

From these figures, it can be seen that the introduction of S2 helped to flatten the beam to produce a flat transverse dose distribution. The high dose at the middle point seemed to have been added to the lateral points. This flat beam can be used to treat tumors with relative large sizes.

#### 5.5.4 Results No. 4

The entire double-scattering system (S1 and S2) was used to produce the fourth simulation results. This case was MCNPX-Case 1 with the tally-geometry condition No. 1 and No. 2.

The energy spectrum is shown in Figures 5-28, from which it can be seen that the energies of most of the particles are between 150 and 230 MeV. In an MCNPX model, the normal direction of the front surface of the water phantom is along the opposite direction of the X-axis. The angular distribution spectrum is shown in 5-29. The abscissa is the cosine of the angle between the direction of an incident particle and the normal of the front surface. From this figure, it can be concluded that most of the incident particles fly into the water phantom perpendicularly. Besides, some low-probability backscattered protons are distributed nearly uniformly between 0 and 90 degrees. The depth-dose distribution and transverse dose distributions at three depths on SOBP are shown in Figure 5-30. The contour depth-dose distribution in the central layer (Y: -0.5 ~ 0.5 cm) is shown in Figure 5-31. It can be concluded that the double-scattering system over-scattered the beam because the transverse dose distributions look “dished.”

The average dose in the water phantom (result from F6 tally) was  $6.26 \times 10^{-4}$  MeV  $g^{-1}$  per particle. Hence, the integral dose (the production of average dose and the mass of the water phantom) was 160.2 MeV per particle. The average current (F1 tally) over the front surface of the water phantom is 0.8588 per particle. It can be concluded that 35.92%  $((250-160.2)/250)$  of the initial kinetic energy was lost in the double-scattering

system in average, and 85.88% of the source particles can deposit energy in the water phantom.

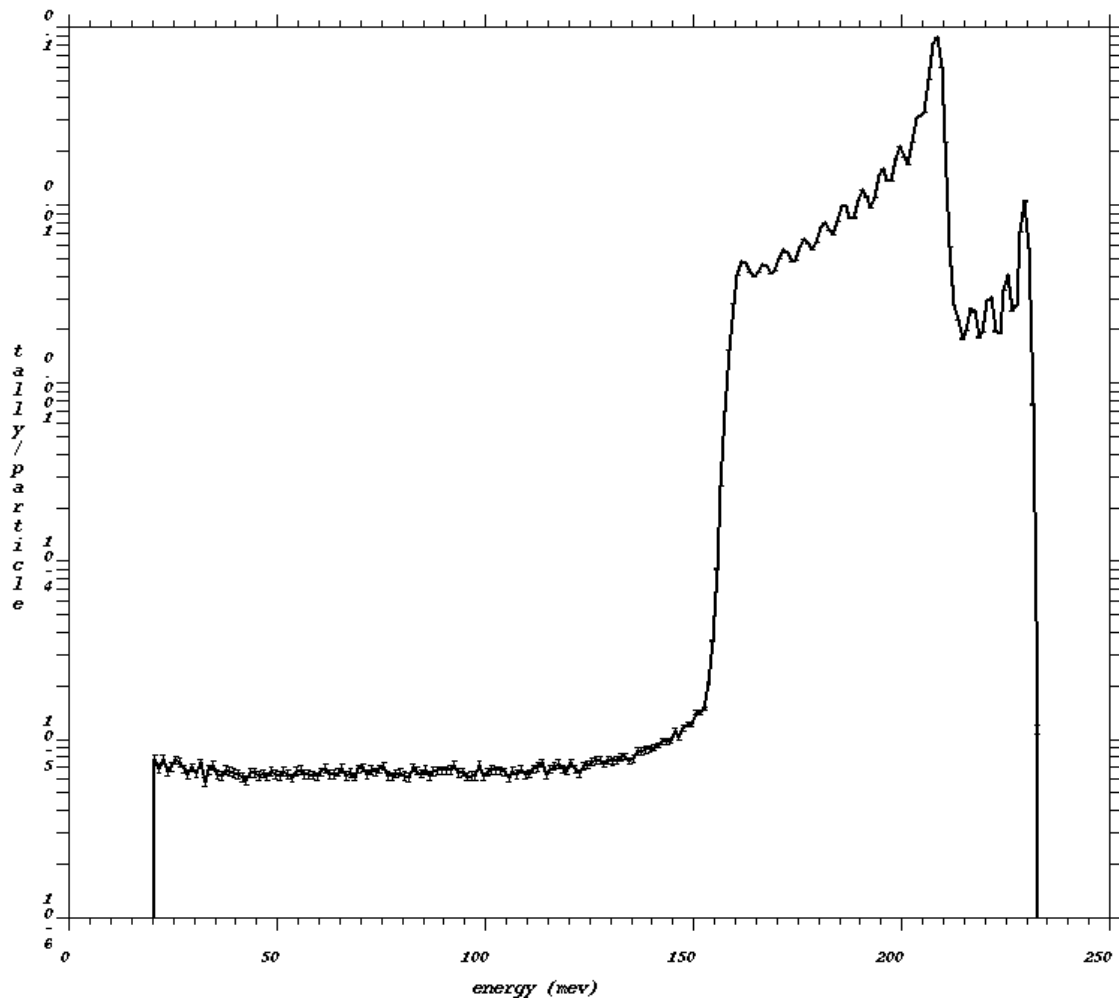


Figure 5-28. The energy spectrum of incident protons in water phantom

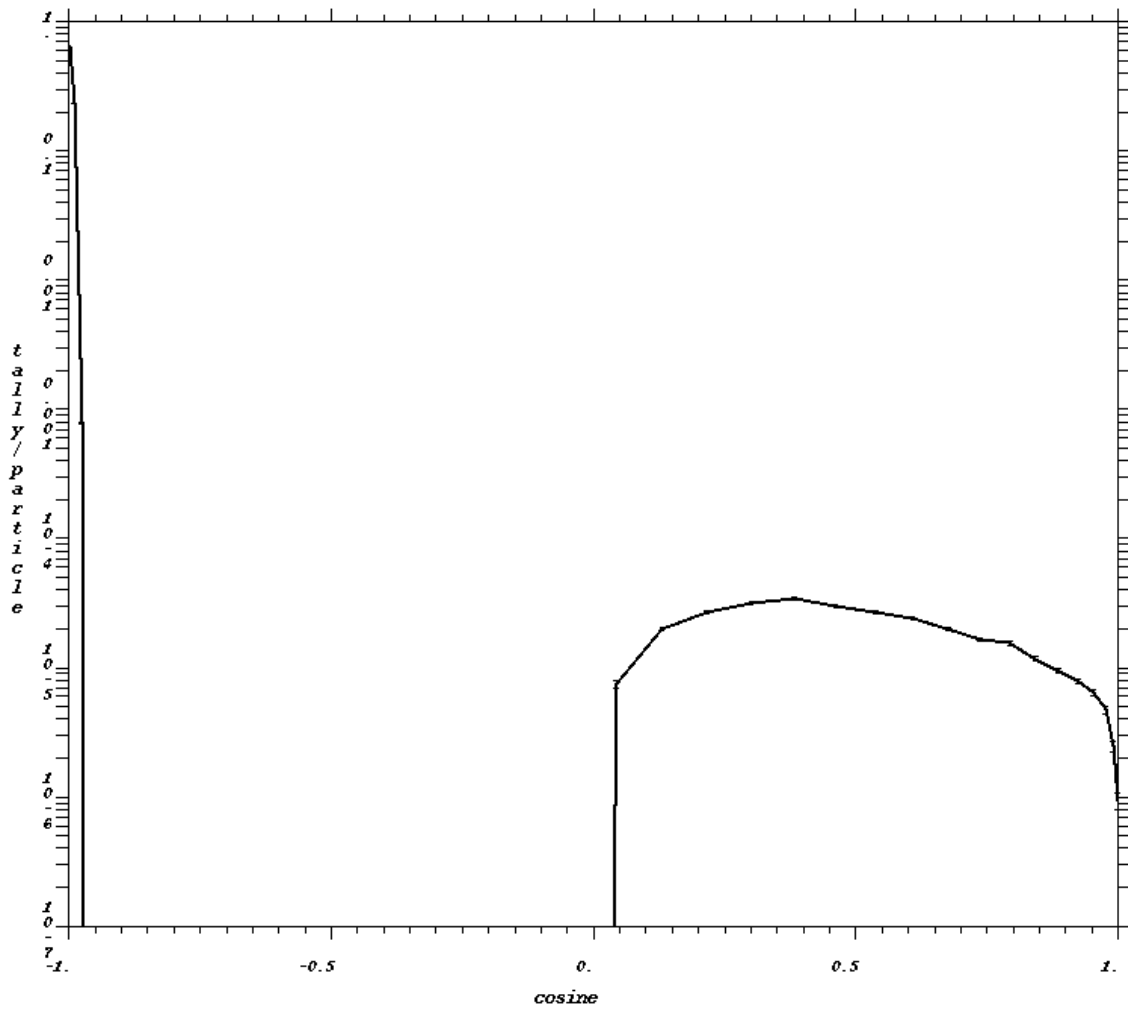


Figure 5-29. The angular distribution spectrum of incident protons in water phantom

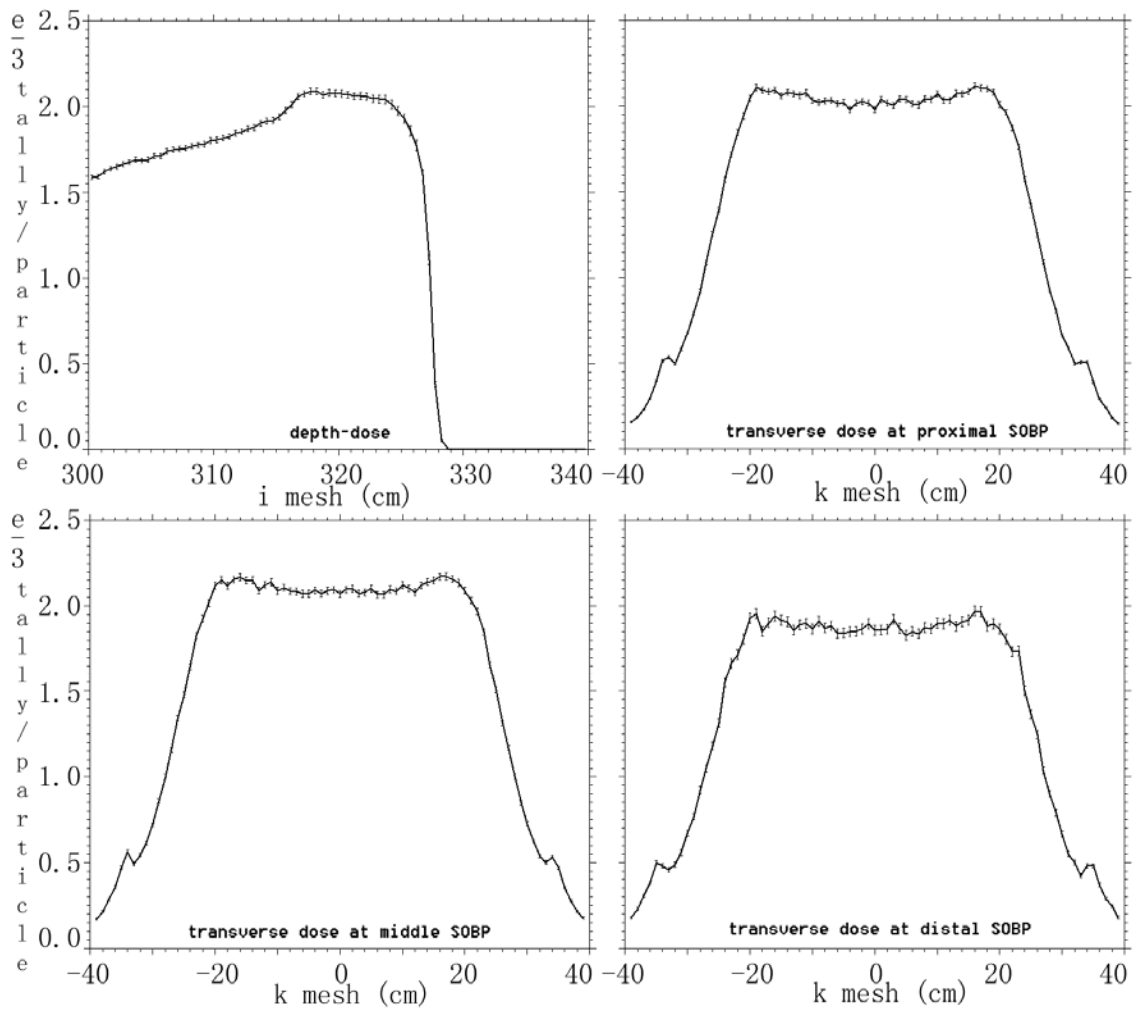


Figure 5-30. The depth-dose and transverse dose distributions for Results No. 4



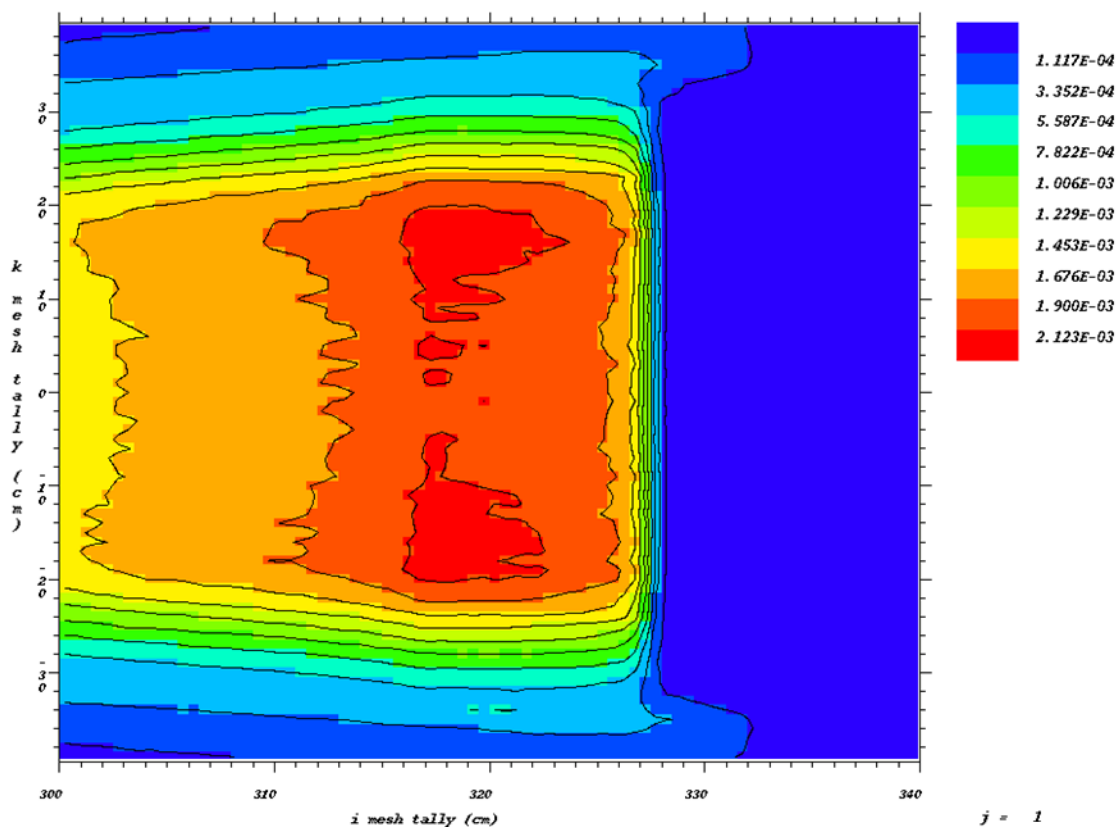


Figure 5-31. The contour depth-dose distribution in the central layer for Results No. 4

The formation of the SOBP curve is shown in Figure 5-32. A comparison of SOBPs from the NEU design result and the MCNPX simulation result is shown in Figure 5-33. In the dose plateau region, the MCNPX results are higher, but in the decreasing distal region, the MCNPX results are lower. Generally, the MCNPX results are very close to or consistent with the NEU results, especially at the proximal point of the SOBP and at the depth of the range of protons. One reason for this divergence is that the size of each dose scoring cell is different. Each depth-dose region in MCNPX is the central part of the water cell perpendicular to X-axis, but the dose region in NEU is the

whole water cell. In Figure 5-33, if the dose is averaged over a whole water cell, the depth-dose values much closer to the NEU results can be obtained.

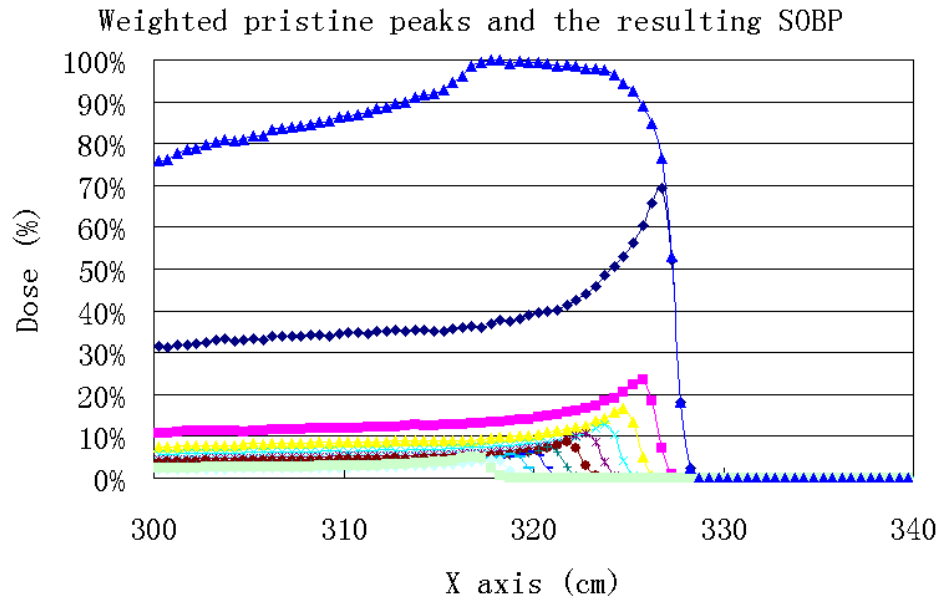


Figure 5-32. The formation of the SOBP by weighted pristine peaks

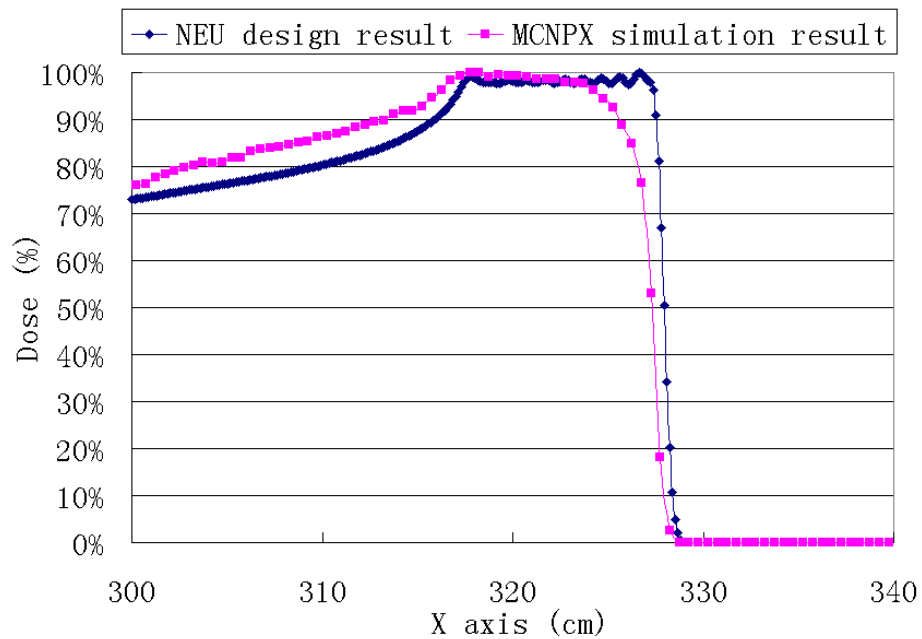


Figure 5-33. The comparison of SOBP between NEU and MCNPX results

The depth-fluence distribution along the X-axis is shown in Figure 5-34. The trend of fluence is decreasing with the increasing of depth as expected.

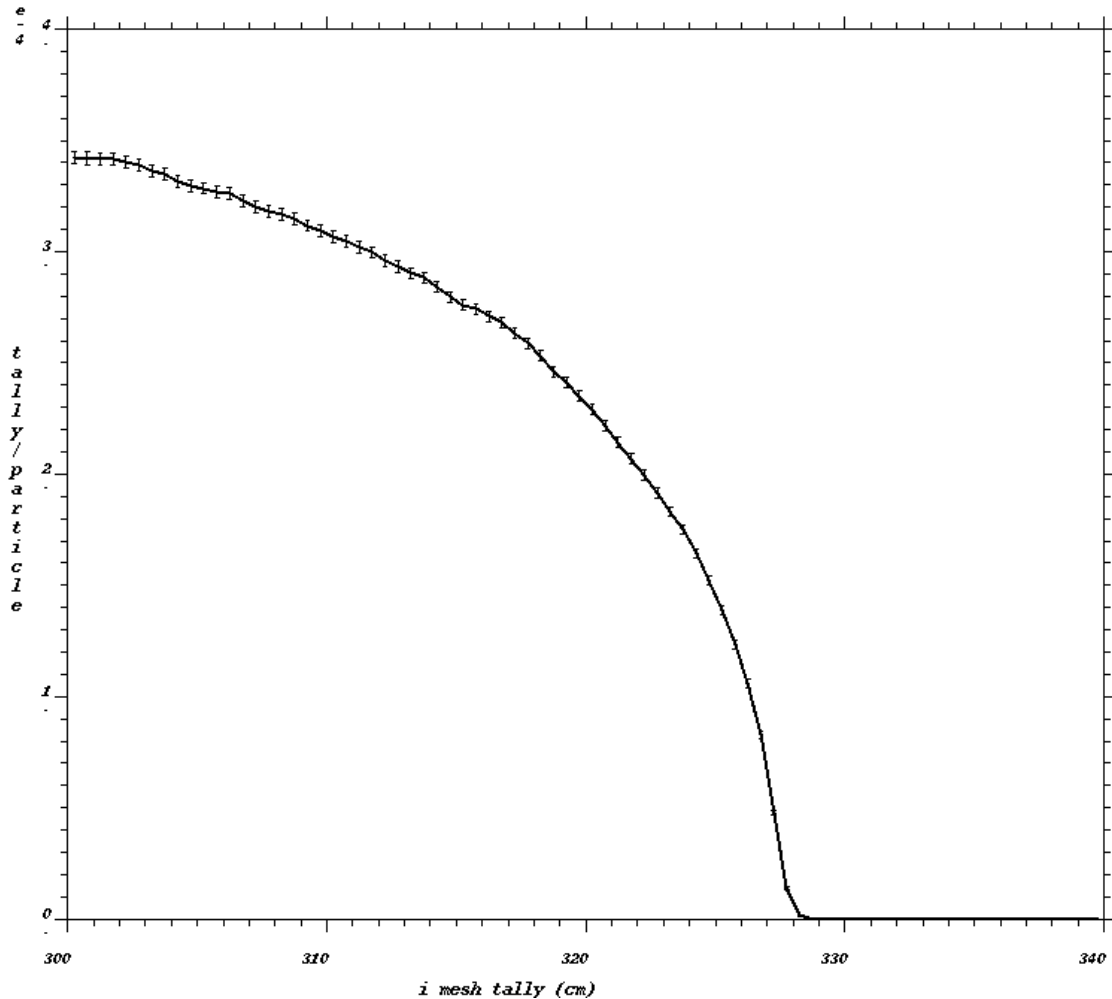


Figure 5-34. The depth-fluence distribution along X-axis

### 5.5.5 Results No. 5

Using the settings for the fourth results, collimators, a final aperture, and a steel outer shell were added to produce the fifth results. This case was MCNPX-Case 4. The radius for the final aperture was 10 cm. The linear and contour depth-dose and transverse

dose distributions are shown in Figure 5-35 and Figure 5-36. From these figures, it can be seen that the field radius is about 12 cm, a little larger than the radius of the final aperture due to the scattering effect; the range of protons is 28 cm; the uniformity of the transverse dose at the distal SOBP is the worst in the three transverse dose plots.

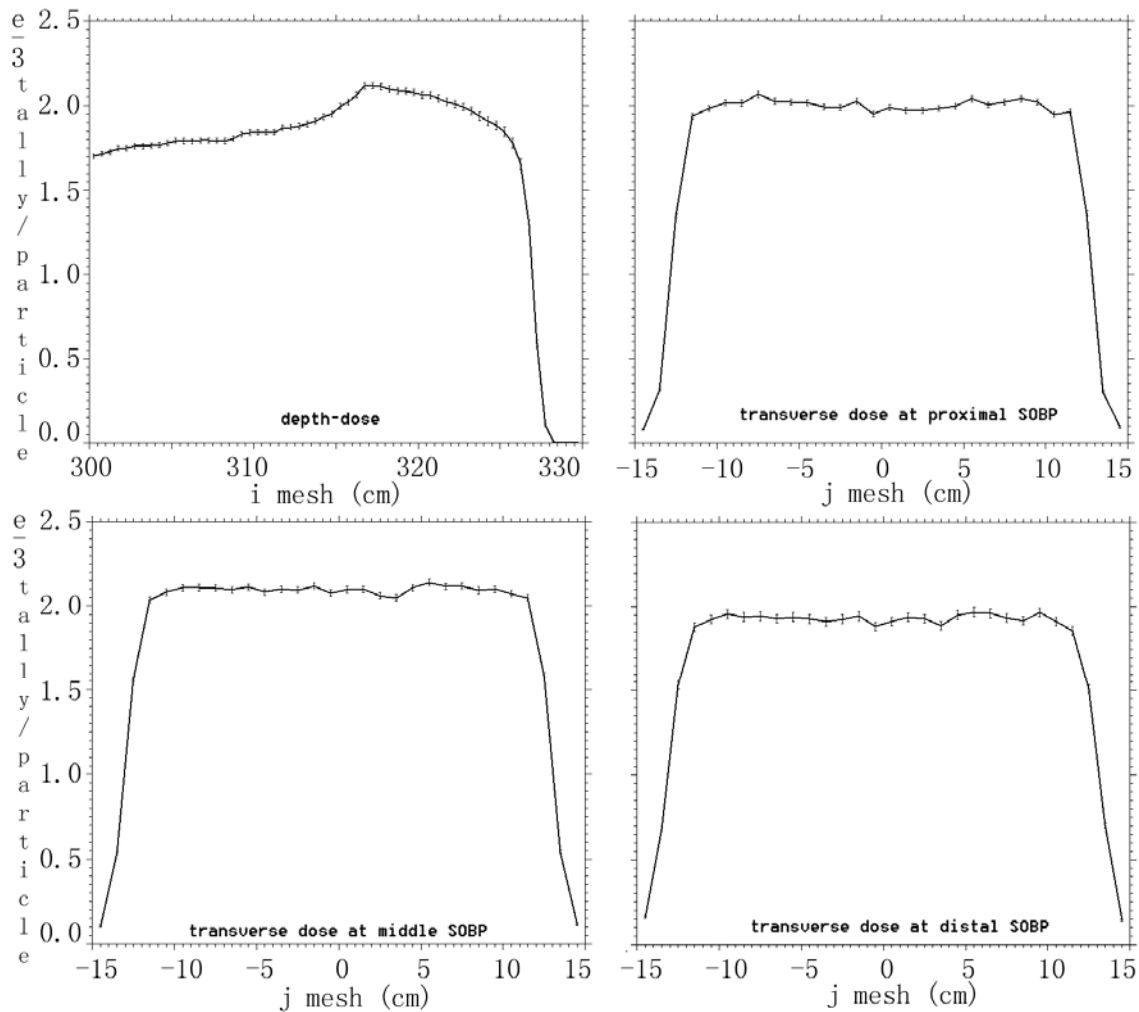


Figure 5-35. The depth-dose and transverse dose distributions for Results No. 5

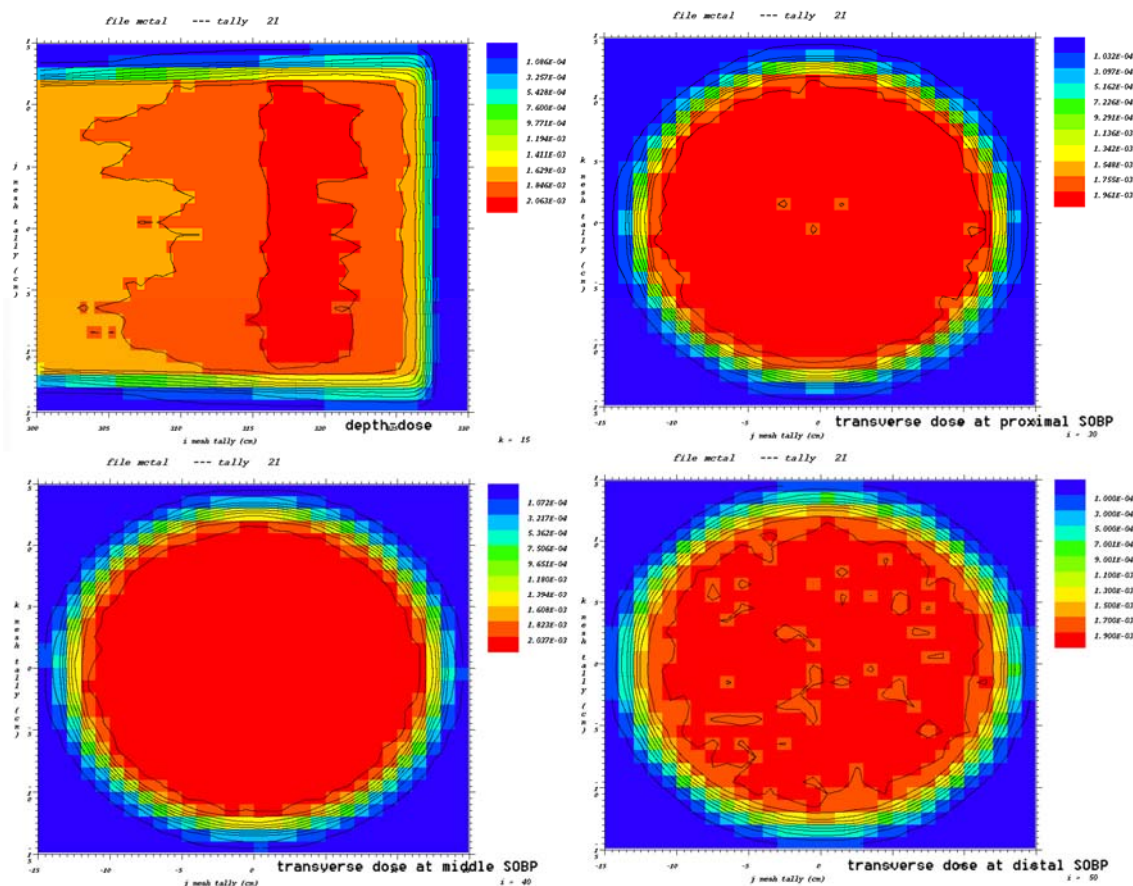


Figure 5-36. The contour dose distributions for Results No. 5

### 5.5.6 Results No. 6

Using the settings for the fifth results, a range shifter and a patient-specific compensator were added to produce the sixth results. This case was MCNPX-Case 7. A hypothetical hemispherical tumor with a radius of 10 cm was located in the water phantom centered at 315 cm on the X-axis. A 1 cm long range shifter (made of ABS resin with a density of  $1.04 \text{ g cm}^{-3}$ ) and an inner-hemispherical patient-specific compensator with a radius of 10 cm were added in the nozzle. The linear and contour depth-dose distributions are shown in Figure 5-37 and Figure 5-38. The contour

transverse dose distributions are shown in Figure 5-39. From these figures, it was concluded that the proton range was about 27 cm and profiles of the outer-dose contours were hemispherical. Hence, the high-dose region was nearly “conformal” to the hemispherical tumor.

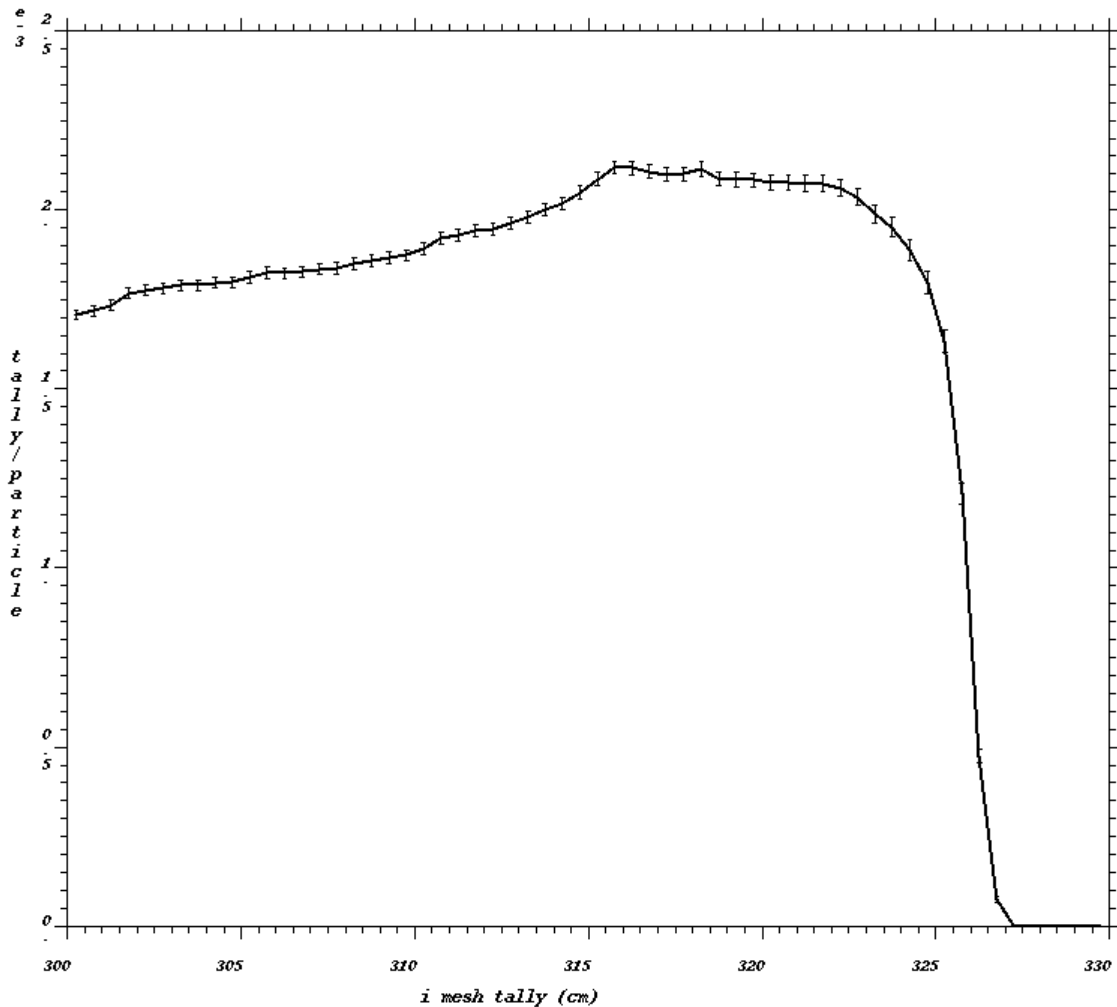


Figure 5-37. The depth-dose distribution for Results No. 6

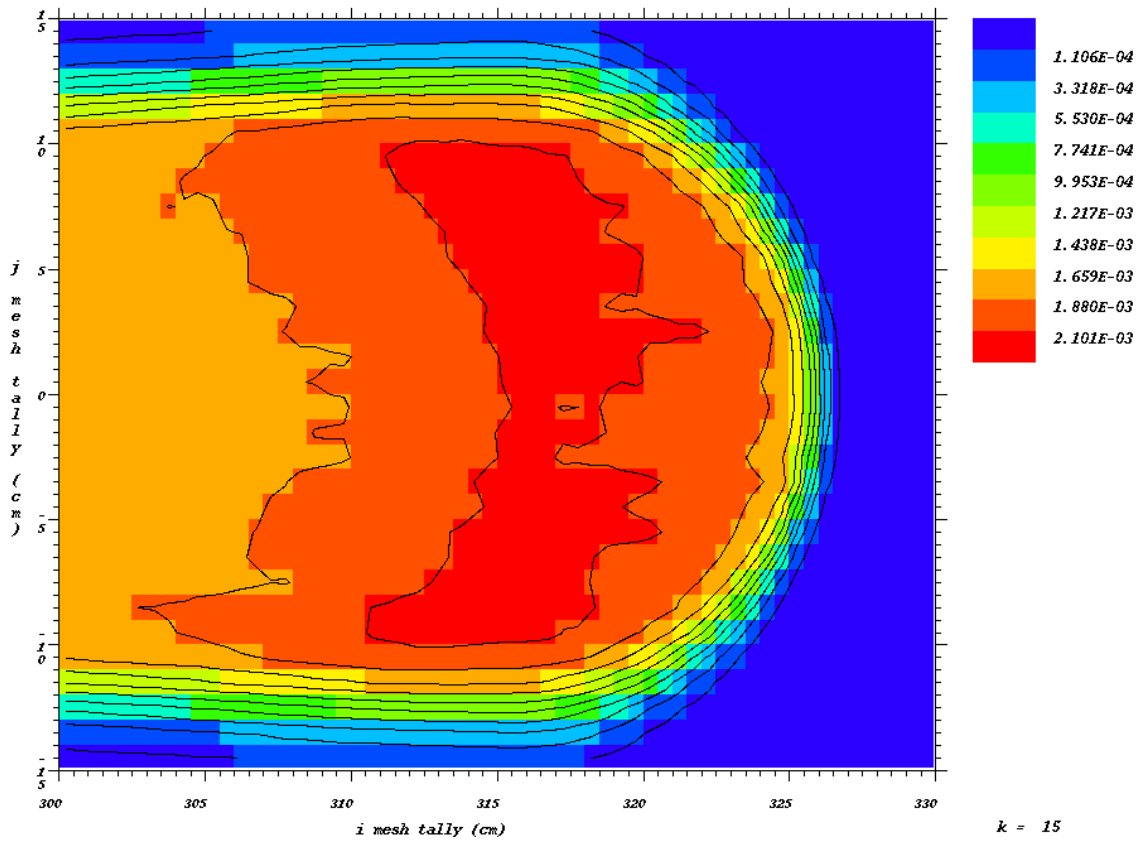


Figure 5-38. The contour depth-dose distribution for Results No. 6

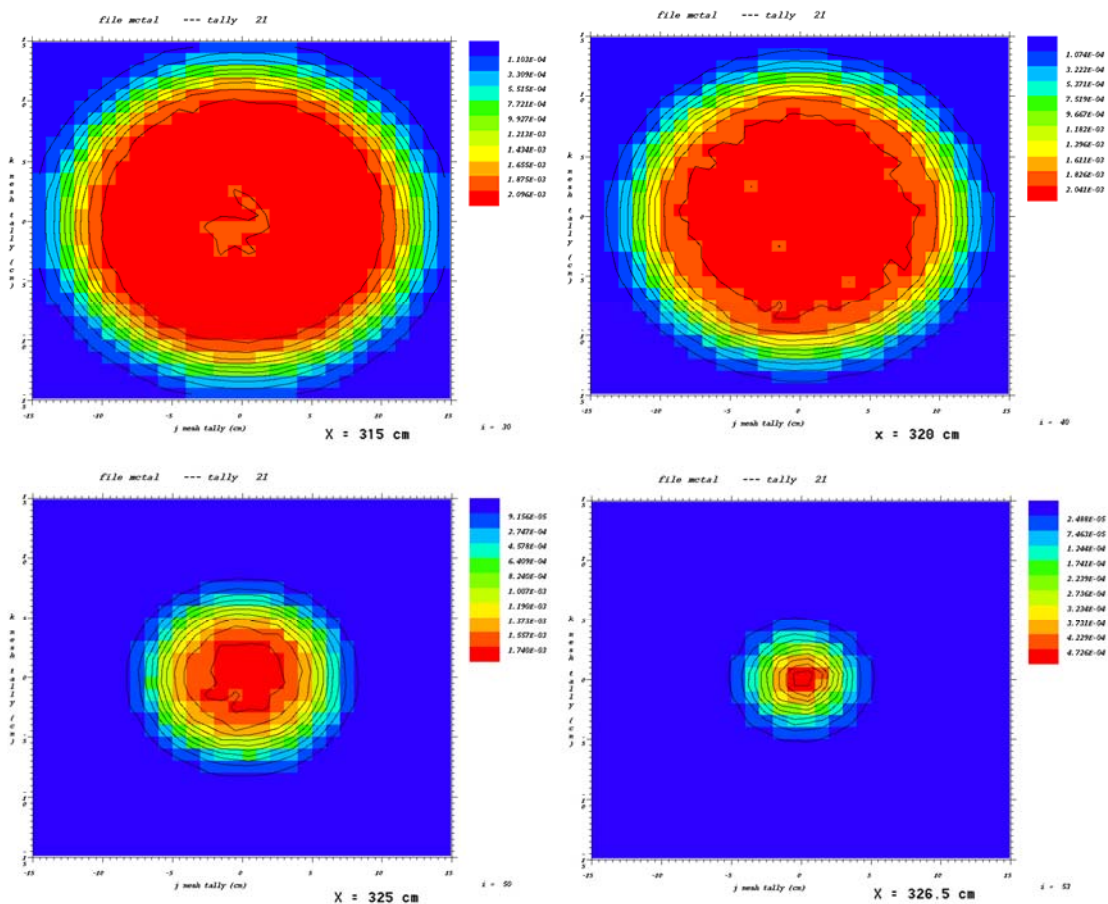


Figure 5-39. The contour transverse dose distributions at different depths

### 5.5.7 Results No. 7

The seventh results came from MCNPX-Case 11. The beam energy was 180 MeV; the radius of the final aperture was 7 cm; the expected SOBP width was 8 cm. The VB script embedded in Excel was used to analyze this case. The dose distributions in the central layer are shown in Figure 5-40 and the relative errors are shown in Figure 5-41.

“i” or “j” is the mesh number in the corresponding dimension. Only near the decreasing



distal part were the relative errors higher than 10%. The depth-dose distribution plotted using MCNPX is shown in Figure 5-42.

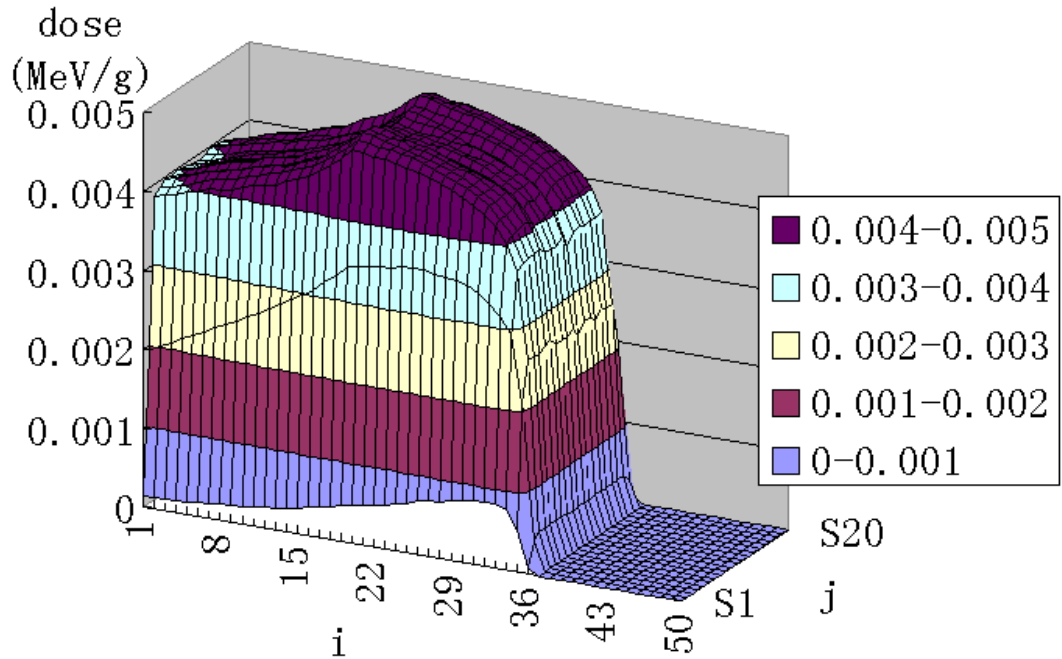


Figure 5-40. The dose distributions in the central layer plotted by Excel

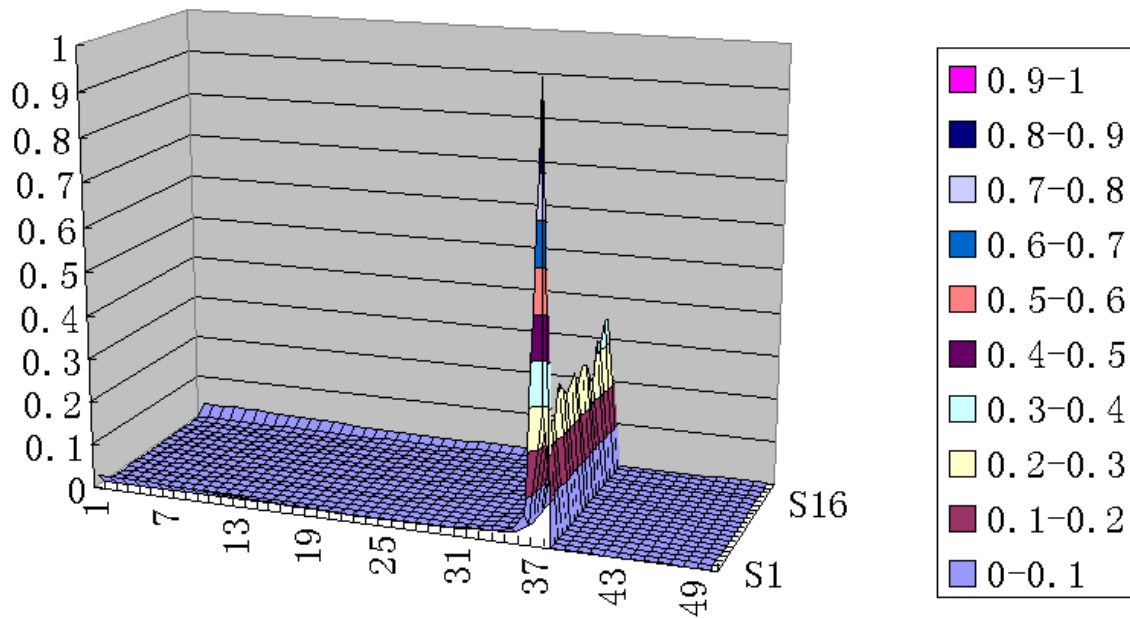


Figure 5-41. The relative errors for dose distributions in the central layer

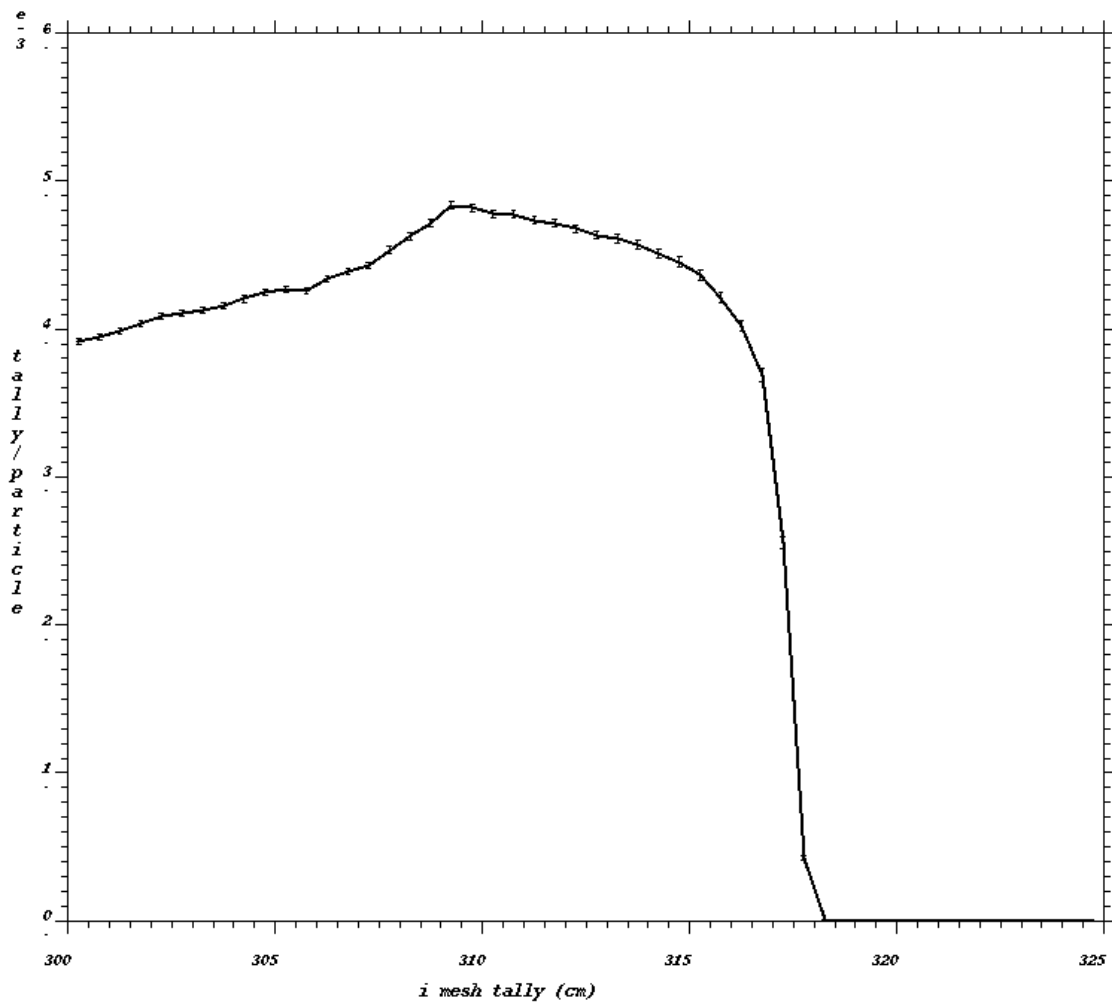


Figure 5-42. The depth-dose distributions in the central layer plotted by MCNPX

#### 5.5.8 Results No. 8

The eighth results came from MCNPX-Case 15. The beam energy was 100 MeV; the radius of the final aperture was 4 cm; the expected SOBP width was 2 cm. The depth-dose distribution is shown in Figure 5-43. The contour transverse dose distribution at mid-SOBP is shown in Figure 5-44. It can be found that the shape of SOBP was not as flat as expected with the NEU design, but the transverse dose was still flat.

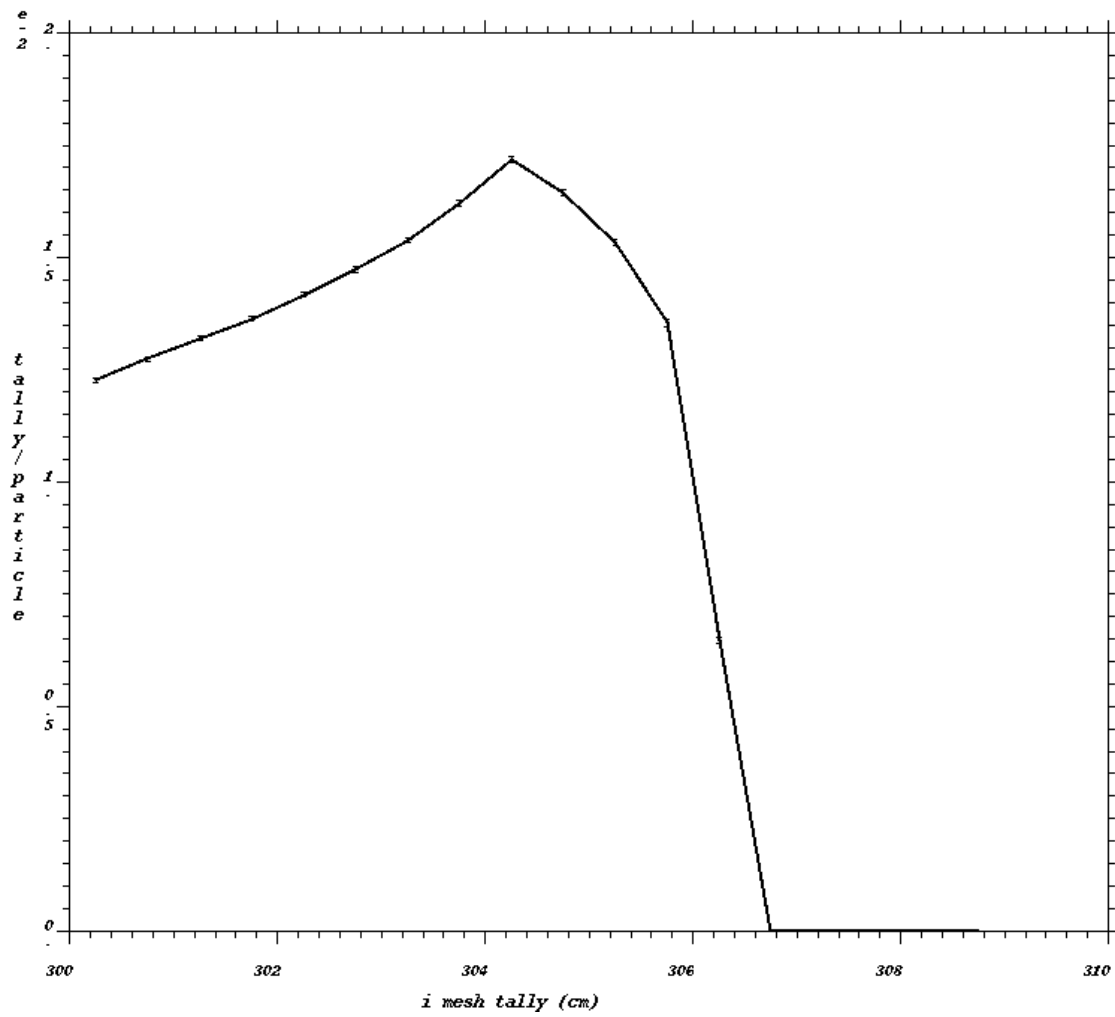


Figure 5-43. The depth-dose distribution for Results No. 8

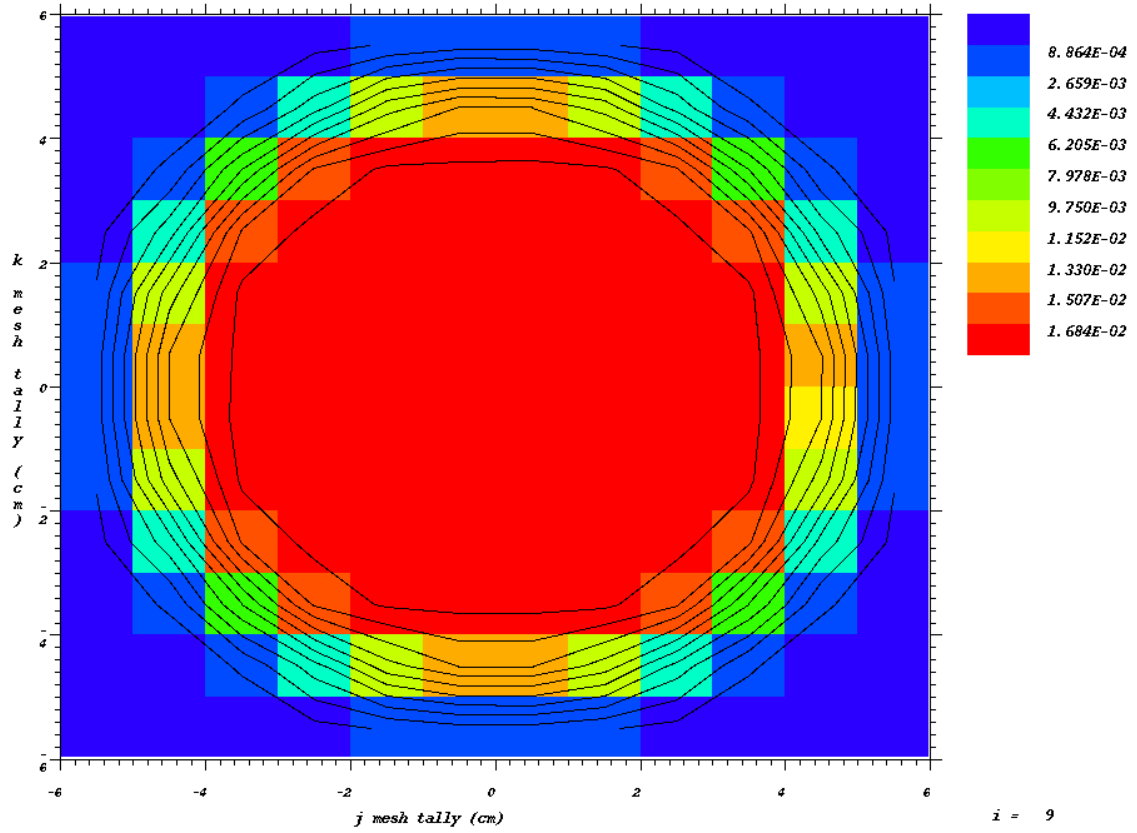


Figure 5-44. The contour transverse dose distribution for Results No. 8

## 6. SUMMARY AND FUTURE WORK

### 6.1 Summary

Even though the scanning technique in proton therapy is more advanced, the passive-scattering nozzle is still used more widely. It is very significant to concern the design and simulation of a passive-scattering nozzle.

The double-scattering system is a key component in a passive-scattering nozzle for proton therapy. The first scatterer is used to broaden the narrow beam, and the second scatterer is used to flatten the broad beam. After passing through this double-scattering system, a narrow beam can be used to produce broad and flat transverse dose and fluence distributions in the water phantom. A rotating multiple-step range modulation wheel can be used to modify the beam to produce a SOBP distribution in the depth-dose curve. Currently, the popular design modes are positioning the double-scattering system upstream in the nozzle to decrease unwanted scattering and combining the first scattering foil and range modulation step together to save space.

In this research, we used the NEU codes package to design the double-scattering system that can meet the requirements of the dose distribution. The parameters of the double-scattering system were used in the simulation procedure using MCNPX. After comparing the simulation results with the design goals, we concluded that NEU is a very useful and powerful tool in designing the double-scattering system, and MCNPX can be successfully used to simulate the proton therapy-related problems.

The application of “mcnp\_pstudy” script successfully solved the problem that MCNP(X) cannot be used to simulate the transport problems with varying parameters and dynamic geometries. In this research, it was used to create input files with different parameters and invoke MCNPX to simulate problems with a rotating range modulation wheel.

The “merge\_mctal” script can be used to merge and average the “mctal” files including the data in the exact same format from MCNP. If some parameters in the “mctal” files were modified first, it can also be used to deal with the “mctal” files from MCNPX. The application of “mcnp\_pstudy” and “merge\_mctal” scripts makes it possible to obtain real simulation results from a proton therapy nozzle mounted with a rotating range modulation wheel.

The VB script embedded in an Excel file edited in this research can also be used to read in the scores from most tally types in the “mctal” files created by either MCNP or MCNPX and spread the data in a desired arrangement format. The user can flexibly edit different scripts to meet different requirements.

## 6.2 Future Work

Even though several problems in the design and simulation of a passive-scattering nozzle in proton beam radiotherapy have been solved, there are still lots of problems that should be solved. These represent extensions of the existing research and should be pursued.

In this research, only MCNPX was used to verify the design results from NEU. Actually, several other codes could be used to simulate the transport of protons. It would be possible to use Geant4 or/and Fluka to verify the design results from NEU and compare the simulation results with the results from MCNPX in future research.

The current “merge\_mctal” script can only be used to merge the “mctal” files from MCNP, but cannot be used to deal with the “mctal” files from MCNPX directly. MCNPX can be used to track more types of particles than MCNP, and the output formats in particle types and mesh tally are different between MCNP and MCNPX. It should be possible to modify the current script or edit a new script suitable to the “mctal” files from MCNPX.

The current VB script embedded in Excel used to read in the data from the “mctal” files cannot be used to read in the coordinates of mesh tally. This function should be added in a future VB script.

In this research, the effect of secondary radiation such as scattered neutrons was omitted. It should be considered in future research.

The clinical effective dose for tumor control and tolerance dose for critical tissues were also not mentioned in this research. Clinical measurement data were not provided in this thesis, so there was no comparison between simulation results with measurements. The current design and simulation results need further verification experimentally in the future.

The effect of the distance between S1 and S2 on the dose and fluence distribution was not studied in this research. The distance was set a fixed value of 50 cm in all the

cases. The separate type of S1 (the scattering foil and the range modulation part are separated by a distance) was not used, but rather the integrated type of S1 (the scattering foil and the range modulation part are connected together) was used in this research. The effect of distance should be studied in the future.

Only a water phantom was used to score the dose and fluence distributions. A more realistic mathematical human phantom should be added in the radiation field to obtain the dose and fluence distributions, so that the interaction mechanism of protons with human body can be better understood.



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## APPENDIX A

## SIMULATION RESULTS FROM NEU-CASE 5 AND 9

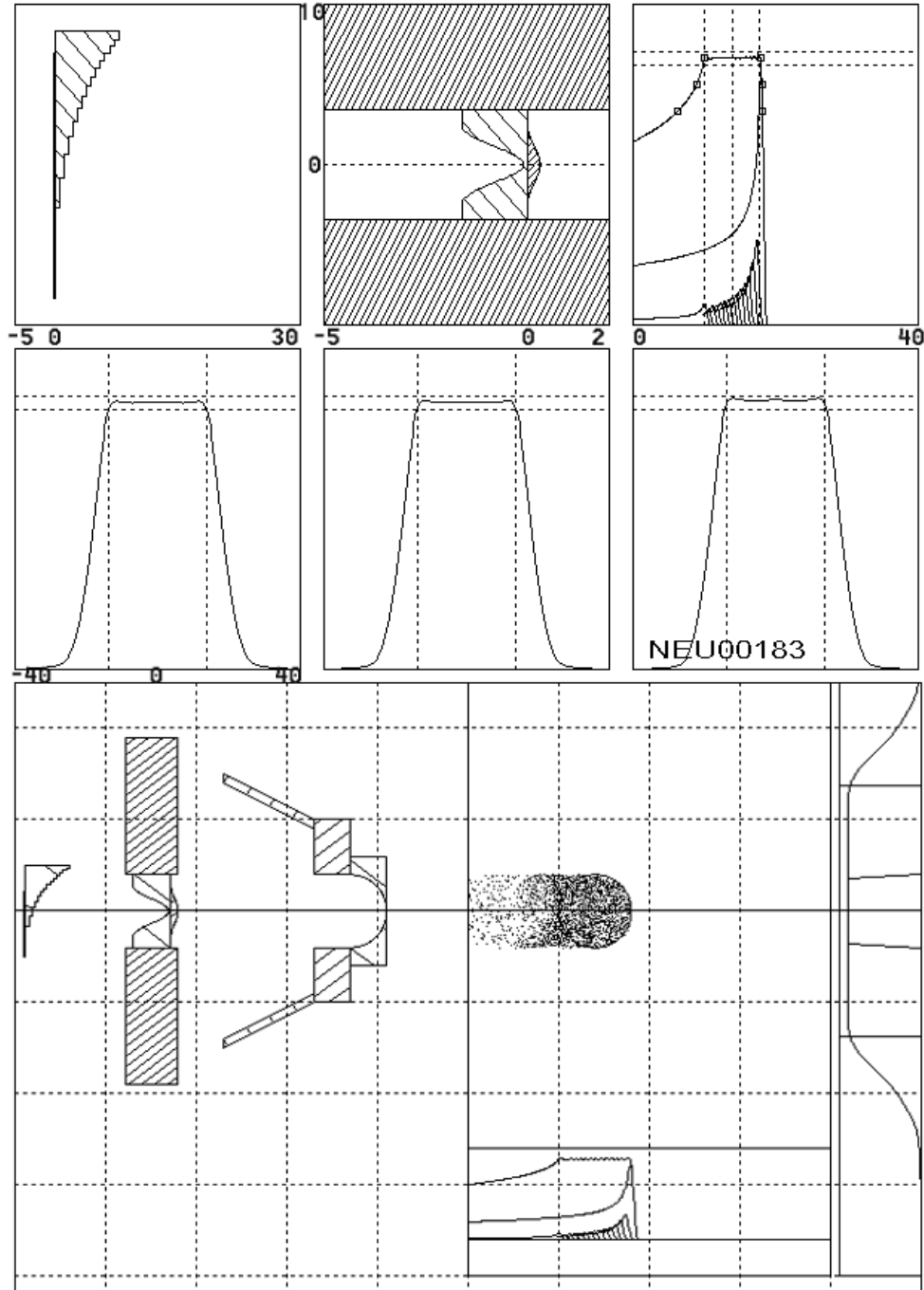


Figure A-1. Results from NEU-Case 5 (180 MeV, medium field)



## APPENDIX B

## EXECUTION OF “MCNP\_PSTUDY” AND “MERGE\_MCTAL” SCRIPTS

## B.1 “Mcnp\_pstudy” Input File for MCNPX-Case 7

```

c   Created on: Wednesday, August 5th, 2009 at 19:00
c beam energy H_Eng
c @@@ H_Eng = 250
c s1 parameters: weight, pb and Lexan thicknesses, number of steps
c @@@ No_Step = 11
c @@@ Step_Wgt = 0.4006 0.1378 0.0965 0.0739 0.0608 \
c @@@      0.0513      0.0437 0.0379 0.0343 0.0287 0.0344
c @@@ S1_Pb = 0.8829      0.8679 0.8515 0.8344 0.8166 \
c @@@      0.7982 0.7791 0.7593 0.7388 0.7176 0.6956
c @@@ S1_LE = 0.0001 0.9377      1.8827 2.8317 3.7842 \
c @@@      4.7402 5.6996 6.6627 7.6294 8.6003 9.5754
c s2 parameters: radius, Lexan and pb thicknesses, shielding size
c @@@ r0 = 0.001
c @@@ r1 = 0.5331
c @@@ r2 = 1.0662
c @@@ r3 = 1.5993
c @@@ r4 = 2.1324
c @@@ r5 = 2.6655
c @@@ r6 = 3.1986
c @@@ r7 = 3.7317
c @@@ r8 = 4.2648
c @@@ r9 = 4.7979
c @@@ r10 = 11.0
c @@@ t101 = 0.1002
c @@@ t102 = 0.4135
c @@@ t103 = 1.2238
c @@@ t104 = 2.2867
c @@@ t105 = 3.3258
c @@@ t106 = 4.071
c @@@ t107 = 4.4633
c @@@ t108 = 4.5001
c @@@ t109 = 4.5001
c @@@ t110 = 4.5001
c @@@ t201 = 0.8731
c @@@ t202 = 0.811
c @@@ t203 = 0.6505

```

```

c @@@ t204 = 0.4397
c @@@ t205 = 0.2334
c @@@ t206 = 0.0853
c @@@ t207 = 0.0073
c @@@ t208 = 0
c @@@ t209 = 0
c @@@ t210 = 0
c @@@ air101 = ( (t102 - t101) )
c @@@ air102 = ( (t103 - t102) )
c @@@ air103 = ( (t104 - t103) )
c @@@ air104 = ( (t105 - t104) )
c @@@ air105 = ( (t106 - t105) )
c @@@ air106 = ( (t107 - t106) )
c @@@ air107 = ( (t108 - t107) )
c @@@ air108 = ( (t109 - t108) )
c @@@ air109 = ( (t110 - t109) )
c @@@ pb201 = ( (t202 - t201) )
c @@@ pb202 = ( (t203 - t202) )
c @@@ pb203 = ( (t204 - t203) )
c @@@ pb204 = ( (t205 - t204) )
c @@@ pb205 = ( (t206 - t205) )
c @@@ pb206 = ( (t207 - t206) )
c @@@ pb207 = ( (t208 - t207) )
c @@@ pb208 = ( (t209 - t208) )
c @@@ pb209 = ( (t210 - t209) )
c collimators and aperture's inner size, large field
c @@@ Rcoll = 12.5
c @@@ Raper = 10
c @@@ OPTIONS = -inner
c @@@ OPTIONS = -mcnp 'mpirun -np 20 /usr/local/mcnpv-v26f/bin/mcnpv.mpi'
c @@@ OPTIONS = -mcnp_opts 'o=1e7out'
c @@@ OPTIONS = -setup
c c @@@ OPTIONS = -run
c c @@@ OPTIONS = -collect
  1  0  1 $void
 101  0  -101 $cookie cutter
 201 252 -11.35 -201 $s1 compensator Pb
 202 289 -1.2 -202 $s1 range modulator
 301 289 -1.2 -301 311 312 313 314 315 316 317 $s2 Lexan
 302 252 -11.35 -321 :-322 :-323 :-324 :-325 :-326 :-327 $s2 pb
 303 290 -8.35 331 -332 $collimator brass
 401 256 -1.04 -401 $range shifter, ABS resin
 501 290 -8.35 -501 503 $final aperture, brass
 502 256 -1.04 -504 505 $patient specific compensator, ABS

```

```

503 290 -8.35 -501 502 trcl=22 $2nd collimator
504 290 -8.35 -501 502 trcl=3 $3rd collimator
505 290 -8.35 -501 502 trcl=4 $4th collimator
601 244 -7.86 -602 601 $steel shell
701 280 -1 -701 $water tank
801 204 -0.001225 -1 #101 #201 #202 #301 #302 #303 #401 #501 #502
      #503 #504 #505 #601 #701 $air

1 so 400 $void sphere
101 rcc -30.5 0 0 1 0 0 3 $cookie cutter
201 rcc 0 0 0 -S1_Pb 0 0 10 $s1 scattering power compensator Pb
202 rcc 0 0 0 S1_LE 0 0 10 $s1 range modulator Lexan
301 1 rcc 0 0 0 -t110 0 0 r9 $s2 Lexan outer
311 1 trc -t102 0 0 air101 0 0 r1 r0 $s2 Lexan air trunc1
312 1 trc -t103 0 0 air102 0 0 r2 r1 $s2 air trunc2
313 1 trc -t104 0 0 air103 0 0 r3 r2
314 1 trc -t105 0 0 air104 0 0 r4 r3
315 1 trc -t106 0 0 air105 0 0 r5 r4
316 1 trc -t107 0 0 air106 0 0 r6 r5
317 1 trc -t108 0 0 air107 0 0 r7 r6
c 318 1 trc -t109 0 0 air108 0 0 r8 r7
c 319 1 trc -t110 0 0 air109 0 0 r9 r8
c $s2 pb trunc 1
321 1 trc t201 0 0 pb201 0 0 r0 r1
322 1 trc t202 0 0 pb202 0 0 r1 r2
323 1 trc t203 0 0 pb203 0 0 r2 r3
324 1 trc t204 0 0 pb204 0 0 r3 r4
325 1 trc t205 0 0 pb205 0 0 r4 r5
326 1 trc t206 0 0 pb206 0 0 r5 r6
327 1 trc t207 0 0 pb207 0 0 r6 r7
c 328 1 trc t208 0 0 pb208 0 0 r7 r8
c 329 1 trc t209 0 0 pb209 0 0 r8 r9
331 1 rcc 2 0 0 -8 0 0 r9 $s2 inner shielding
332 1 rcc 2 0 0 -8 0 0 r10 $s2 outer shielding
401 2 rpp 0 1 -15 15 -15 15 $range shifter
501 5 rpp -10 0 -20 20 -20 20 $collimator outer
502 5 rpp -10 0 -Rcoll Rcoll -Rcoll Rcoll $collimator inner
503 5 rcc -10 0 0 10 0 0 Raper $aperture inner
504 5 rpp 0 Raper -18 18 -18 18 $compensator outer
505 5 so Raper $spherical compensator inner
601 rpp -35 250 -20 20 -20 20 $steel shell inner
602 rpp -40 250 -25 25 -25 25 $steel shell outer
701 rpp 300 340 -40 40 -40 40 $water tank

```



```

mode h
imp:h 0 1 14r    $ 1,801
c material cards
m280 1000.    -0.111915 $Water (density of 1 assumed)
      8000.    -0.888085
m289 1000.    -0.055491 $Lexan
      6000.    -0.755751 8000.    -0.188758
m290 29000.   -0.67 $brass
      30000.   -0.33
m252 82000.   -1 $lead
m256 1000.    32 $ABS resin,(C3H3N)2:(C4H6)3:(C8H8)5
      6000.    29 7000.    1
m204 7000.    -0.755636 $air -0.001225 (US S. Atm at sea level)
      8000.    -0.231475 18000.   -0.012889
m244 26000.   -1 $iron (-7.86)
c transformation cards
c for s2 contoured second scatterer, 1st collimator
*tr1 50 0 0
c for range shifter, 2nd collimator
*tr2 100 0 0
*tr22 50 0 0 9j -1
c for 3rd collimator
*tr3 100 0 0 9j -1
c for 4th collimator
*tr4 150 0 0 9j -1
c for final aperture, patient specific compensator
*tr5 250 0 0
c void some cells
c void 303 401 501 502 503 504 505 601 801 $keep S1,S2
c void 401 502 $void shifter, compensator
c source definition
c sdef x -30 y d1 z d2 erg d3 par h vec 1 0 0 dir 1 ccc 4 wgt Step_Wgt $Gauss
sdef pos -30 0 0 erg H_Eng par h vec 1 0 0 dir 1 wgt Step_Wgt $point source
c sp1 -41 0.54 0    $Y position Gaussian distribution
c sp2 -41 1.22 0    $Z position Gaussian distribution
c sp3 -4 0.184 100  $Energy Gaussian distribution
c sp3 -4 0.292 180  $Energy Gaussian distribution
c sp3 -4 0.346 250  $Energy Gaussian distribution
c physics settings
cut:h j 20 $emin, range 4.3 mm in water
phys:h 300 $emax
c lca  2 1 0 5j 2 $ use incl4 model
lca  2 1 0 5j 1 $ use cem03 model, spend less time than incl4
lcb  6j 1 0 $ 1 0 stand for cutoff kinetic energy and no correction

```

```

lea 2j 0 $ use mass-energy balance in cascade phase
c tally cards
f1:h 701.2 $current on the front surface of water tank
c e1 1 248i 250 $energy bin
c f11:h 701.2 $current on the front surface of water tank
c *c11 175 34i 0 $angle bin
f6:h 701 $MeV/g, absorbed dose
fm1:h No_Step $for N steps of S1
c fm11:h No_Step $for N steps of S1
fm6:h No_Step $for N steps of S1
c mesh tally, flux, and deposited energy in unit volume of water
tmesh
rmesh21:h flux pedep
cora21 300 59i 330
corb21 -15 29i 15
corc21 -15 29i 15
endmd
fm21:h No_Step $for N steps of S1
c running card
nps 1e7
print
prdm 2j 1 1 0 $mctal file,1 dump,10 TFC

```

## B. 2 Command to Execute the “Mcnp\_pstudy” Script

The input file of MCNPX-Case 1 for “mcnp\_pstudy” was called “250\_large\_void.” The command is shown in Figure B-1.

```
perl mcnp_pstudy -i 250_large_void -setup -run -collect
```

Figure B-1. Command to execute the “mcnp\_pstudy” script

The options “-setup”, “-run” and “-collect” can be specified either in the input file or in the command line. In addition, they can also be specified step by step in

separate command lines. Notice that the format of the input file must be a “UNIX” format; while a “DOS” format file cannot be read in. After “-setup”, eleven sub-cases (representing eleven steps in S1) and a log file were created in current directory, shown in Figure B-2. After “-run”, eleven sub problems would be executed one after one, and the output files would be created.

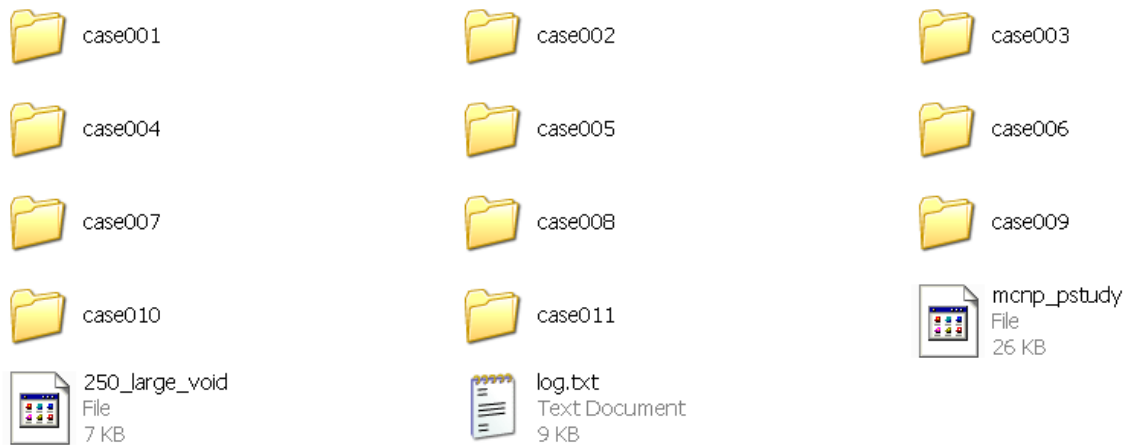


Figure B-2. Creation of sub-directories and files by the “mcpn\_pstudy” script

### B. 3 Command to Execute the “Merge\_mctal” Script

In this research, a script “mergesh” was edited to invoke the command to execute the “merge\_mctal” script. The content of “mergesh” is shown in Figure B-3.

```
perl merge_mctal -o 250_l_void mctal01 mctal02 mctal03 mctal04 mctal05 mctal06 \
mctal07 mctal08 mctal09 mctal10 mctal11
# this is a script to execute perl merge_mctal, Fada Guan 2009 08 04
```

Figure B-3. The content of “mergesh”

The modifications of the “mctal” files from MCNPX-Case 1 are shown in Figure B-4. The original mesh tally ‘21’ was changed to an F5 tally ‘25’. The number of meshes 6320 was kept after ‘f.’ ‘2’ after ‘s’ representing ‘flux’ and ‘pedep’ were used in a mesh tally was kept.

```

ntal      1
  25
tally     25      1      0
f         6320
d          0
u          0
s          2
m          0
c          0
e          0
t          0
vals

```

Figure B-4. The modifications of the “mctal” files from MCNPX-Case 1

After modification of all the mctal files, the command “./mergecsh” was executed on a UNIX platform; then the eleven “mctal” files were merged to one “mctal” file - “250\_1\_void” in the current directory. The procedure is shown in Figure B-5.

```
    :./mergescsh
...Reading MCTAL file:  mctal01
...Reading MCTAL file:  mctal02
...Merging  (except TFC)
...Reading MCTAL file:  mctal03
...Merging  (except TFC)
...Reading MCTAL file:  mctal04
...Merging  (except TFC)
...Reading MCTAL file:  mctal05
...Merging  (except TFC)
...Reading MCTAL file:  mctal06
...Merging  (except TFC)
...Reading MCTAL file:  mctal07
...Merging  (except TFC)
...Reading MCTAL file:  mctal08
...Merging  (except TFC)
...Reading MCTAL file:  mctal09
...Merging  (except TFC)
...Reading MCTAL file:  mctal10
...Merging  (except TFC)
...Reading MCTAL file:  mctal11
...Merging  (except TFC)

...Creating merged MCTAL file = 250_l_void
```

Figure B-5. The execution and print-out information from “mergescsh”

## APPENDIX C

## MCNPX PLOT COMMANDS USED IN THIS RESEARCH

## C.1 Commands for Energy and Angular Distribution Spectra in MCNPX-Case 1, 2 and 3

```

mcnpz z
rmctal mctal
tal 1 free e
nonorm
plinear
tal 11 free c

```

## C.2 Commands for Fluence and Dose Distributions in MCNPX-Case 1, 2 and 3

```

mcnpz z
rmctal mctal
tal 21 free i fixed j 1 fixed k 40 fixed s 1
nonorm
plinear
fixed s 2
linlin
plinear
tal 21 free ik fixed s 1
tal 21 free ik fixed s 2

```

## C.3 Commands for Fluence and Dose Distributions in MCNPX-Case 4 through 15

The commands for MCNPX-Case 4 are listed below. In other cases, only the indexes of i, j and k are different. The indexes of i can be found in Table C-1.

```

mcnpz z
rmctal mctal
tal 21 free ij fixed k 15 fixed s 1
fixed s 2
tal 21 free I fixed j 15 fixed k 15 fixed s 1
nonorm
plinear
fixed s 2
tal 21 free jk fixed i 1 fixed s 1
fixed i 30

```

fixed i 1 fixed s 2  
 fixed i 30  
 tal 21 free j fixed k 15 fixed i 30 fixed s 1  
 fixed i 40  
 fixed i 50  
 tal 21 free j fixed k 15 fixed s 2 fixed i 30  
 fixed i 40  
 fixed i 50

Table C-1. Coordinate or Index of i in the mesh tally for transverse dose distribution

MCNPX-Case No.	Coordinate or Index	Proximal point at SOBP	Middle point at SOBP	Distal point at SOBP
1	X (cm)	316	321	326
	Index of i	32	42	52
2	X (cm)	320	325	330
	Index of i	40	50	60
3	X (cm)	324	329	334
	Index of i	48	58	68
4	X (cm)	315	320	325
	Index of i	30	40	50
5	X (cm)	320	325	330
	Index of i	40	50	60
6	X (cm)	324	329	334
	Index of i	48	58	68
7	X (cm)	314	319	324
	Index of i	28	38	48
8	X (cm)	319	324	329
	Index of i	38	48	58
9	X (cm)	323	328	333
	Index of i	46	56	66
10	X (cm)	306	310	314
	Index of i	12	20	28
11	X (cm)	308	312	316
	Index of i	16	24	32
12	X (cm)	311	315	319
	Index of i	22	30	38
13	X (cm)	302.5	303.5	304.5
	Index of i	5	7	9
14	X (cm)	303	304	305
	Index of i	6	8	10
15	X (cm)	303.5	304.5	305.5
	Index of i	7	9	11

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