

## Analysis of NTCP Based Radiobiological Models: A Systematic Review of Literatures

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Submitted: 02 Oct 2019; Accepted: 12 Oct 2019; Published: 23 Oct 2019

### Abstract

To achieve the optimal treatment goal, radiobiological parameters have to evaluate and predict the outcome of this treatment plan in terms of both TCP and NTCP. Different types of radiobiological model were used to achieve prescribed treatment dose of radiation during the tumor control. Where TCP models play an important role in order to achieve desired dose to the tumor. A suitable NTCP model was theoretically found among different models that can be used in treatment plan evaluation. Theoretically, six different radiobiological dose response models were analyzed in this project. Lyman–Kutcher–Burman, Critical element, critical volume, Relative Seriality, Parallel architecture, Weibull distribution models were analyzed from the derivation. All models were discussed elaborately with its various parameters and were used in the calculation of normal tissue complication probability during the treatment in radiotherapy. Further, all models were compared with each other. The models denote the dose for 50% complication probability (D50) parameters is the most commonly used radiobiological models for the normal tissues. The functional subunit response models (critical element & Relative seriality, Critical Volume, parallel architecture) are used in the derivation of the formulae for the normal tissue. Since all complicated NTCP model predict same as the simple NTCP model that is Lyman–Kutcher–Burman model as well as it is computationally efficient. Also Lyman–Kutcher–Burman model can be used in different treatment planning system incorporating with other model. For this reason, our suggested model is Lyman–Kutcher–Burman NTCP model which can be used in treatment plan evaluation. After analyzing six different model of NTCP, finding of the study is the treatment plan evaluation in where Lyman–Kutcher–Burman model is the best model for biological plan evaluation.

**Keywords:** Radiobiological Model, TCP, NTCP

### Introduction

Radiobiological models is to predict the radiation response of biological systems While early approaches focused on modeling the radiation response for different fractionation schemes, newer developments attempt to model effect probabilities (TCP and NTCP), their volume dependence and the relative biological efficiency (RBE) of radiation with high linear energy transfer LET [1]. To reduce the suffering and death due to cancer is now one of the biggest challenges ever. Continuous research and technical developments give us hope and confidence about winning the battle against cancer. Radiotherapy has been a major weapon in this battle for a long time. About 60% of cancer patients receive radiotherapy as part of their disease management. The goal of radiation therapy is to deliver a therapeutic dose of radiation to target tissues while minimizing the risks of normal tissue complications. The radiotherapy has been based describe to the best possible tumor control for the case in which all regions of the tumor have exactly the same biological characteristics and sensitivities to radiation [2]. A biologically based model is developed for normal tissue complication probability as a function of dose and irradiated volume fraction for organs. The

organ is assumed to be composed of functional subunits which are arranged in a parallel architecture. The complication is produced only if a sufficiently large fraction of the FSUs is inactivated by radiation and an FSU is inactivated only when all the clonogenic cells within it are killed [3]. A model of normal tissue complication probability for tissues that may be represented by critical element architecture. We derive formulas for complication probability that apply to both a partial volume irradiation and to an arbitrary inhomogeneous dose distribution. The dose-volume iso-effect relationship which is a consequence of critical element architecture is discussed and compared to the empirical power law relationship. A dose-volume histogram reduction scheme for a “pure” critical element model is derived. The existing published dose-volume histogram reduction algorithms are analyzed. The study has shown that the existing algorithms, developed empirically without an explicit biophysical model, have a close relationship to the critical element model at low levels of complication probability [4]. The Empirical models have also found their niche. The Lyman, Kutcher and Burman Model, Relative Seriality Model, Critical Element Model, Critical Volume Model, Parallel architecture Model. For additional information regarding early application of biological modeling in treatment planning, the reader is referred to a review by Orton et al. (2008). In

this study, the LKB model has been used for NTCP Calculation [5]. The basic radiobiology in the mid of the last century has led to the formulation of first models of the cell kill in, eventually, to models that linked radiation sensitivity to cure rates for tumors. one of the first such formalisms was proposed by munro and gilbert (1961) [6].

### Method

The study had been followed AAPM Report no -166, the modern technology the radiation oncology, Basic Clinical Radiobiology, The Basic Science of Oncology, Comprehensive Biomedical Physics, Radiation Biology of Medical Imaging, Radiobiology for the Radiobiologists, 25 project paper, many national and international journals book references, internet and my practical work was done in that department.

### Radiobiology model

Dose response model divided into four types- Linear-Quadratic Model, Generalized Equivalent Uniform Dose, TCP Models, NTCP Models.

### Liner Quadratic model

The LQ formalism is most commonly used to model cell survival. A cell survival curve describes the relationship between the surviving fractions of cells. Radiation induced reproductive cell death has been conclusively linked to DNA damage, specifically to DNA double-strand breaks (DSB). The LQ model was introduced to relations for different fractionation regimes. The formula for cell survival is:  $P = \exp(-\alpha D - \beta D^2)$

### Generalized Equivalent Uniform Dose

The concept of gEUD proposed by Niemierko [4]. (Niemierk1997) provides a single metric for reporting non-uniform tumor dose distributions [5-8]. It is defined as the uniform dose that, if delivered over the same number of fractions as the non-uniform dose distribution of interest, yields the same radiobiological effect. This equation:

$$gEUD = (\sum_i v_i D_i^a)^{\frac{1}{a}}$$

### Tumor control probability

A majority of mechanistic TCP models are based on the assumption that the number of surviving clonogenic tumor cells, i.e., cells capable of regrowing the tumor, follows the Poisson distribution [9-11]. The Poisson assumption has limitations when clonogen repopulation occurs during treatment. One obvious problem is that simple application of exponential tumor growth predicts that all

tumors recur at sufficiently long times after external beam treatment or for permanent implants with exponentially decaying sources.

### Normal tissue complication probability

The NTCP models aim to describe the complication probability in normal tissues in terms of dose-response curves. As there is extensive evidence, that the radiation response of normal tissue depends on the amount of irradiated normal tissue the irradiated volume is included as an additional important parameter. The extent of the volume effect is dependent on the architecture of the respective tissue and several models have been proposed [12-14].

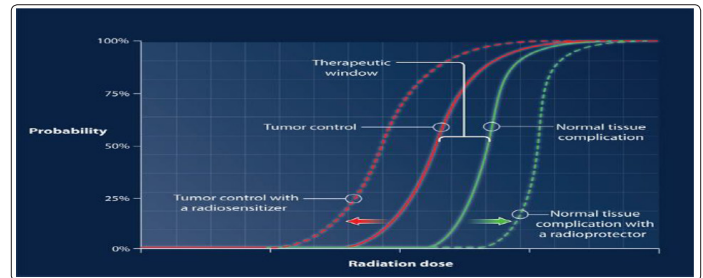


Figure 1: The normal tissue complication probability

### Results

Dose-response models can be categorized into several groups based on the statistical distribution. They use for describing the sigmoid shape of the dose-response curve (Figure 2). The five distributions used in the models Investigated in the present study are the poisson, binomial, probit, logit, and Weibull distributions.

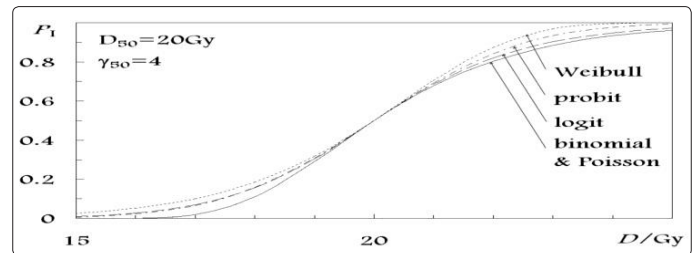


Figure 2: Statistical distribution used in NTCP models to describe the shape of the dose response curve

The Success of any Practical Implementation of the Concept of NTCP in Treatment Planning Depends:

1. Treatment planning & normal tissue specific parameters.
2. Extrapolate NTCP to other treatment modalities.

**Table 1: Overview of the dose-response models together with a summary of their inherent parameters**

Statistics	Model	Proposed	Equation	inherent model parameters
Probit	Lyman-Kutcher-Burman model	Kutcher-1985, Burman -1989	$NTCP = (2\pi)^{-\frac{1}{2}} \int_{-\infty}^t \exp(-x^2/2) dx$	M, N, T50, $D_{eff}$
Poisson	Critical Element Model	Schultheiss -1983	$P(D, v) = 1 - [1 - P(D, 1)]^v$	D50, $\gamma 50$
Binomial	Critical Volume Model	Niemierko- Goitein -1992	$P(D) = \sum_{K=M+1}^N \binom{N}{K} p^K (1-p)^{N-K}$	N0, M, N, $D_{50}$
Binomial	Relative seriality model	Kallman-1992	$P_1(D, V) =$	N0, M, N, $D_{50}$
Logit	parallel architecture model	Jackson-1995	$NTCP = \frac{1}{\sqrt{2\pi\sigma^2}} \int_0^f \exp(-v^2/2\sigma^2) dv$	N0, M, N, $D_{50}$
Weibull	Weibull distribution model	Jackson-1995	$P(D) = 1 - \exp[-(D/D_{50})^\gamma]$	N0, M, N, $D_{50}$

**Table 2: NTCP Model Used In Different Treatment Planning System**

Company name	Phillips Pinnacle	Varian Eclipse	CMS Monaco	RaySearch Laboratories
Company Product	Phillips medical system, Andover	Varian medical system, Paloalto	Maryland Heights	Raystation
NTCP	Lyman-Kutcher-Burman model, Kallman S model	Lyman-Kutcher-Burman model, Kallman S model	Parallel Architecture model	Lyman-Kutcher-Burman model, Kallman S model

**Table 3: Parameters Used In Derivation of Different NTCP Model**

Parameters Model	M	N	D50	$\gamma 50$	T50	$D_{eff}$	N0	S	K
Lyman-Kutcher-Burman model	√	√	X	X	√	√	X	X	X
Critical Element Model	X	X	√	√	X	X	X	X	X
Critical Volume Model	√	√	√	X	X	X	√	X	
Relative seriality model	X	X	√	√	X	X	X	√	X
parallel architecture model	X	X	√	X	X	X	X	X	√
Weibull distribution model	X	X	√	√	X	X	X	X	X

**Discussion**

Radiobiological modeling of radiotherapy plans are used for treatment plan comparisons. Radiobiological model plays a important role in plan optimization & evaluation in radiotherapy though aim of radiobiological model is to predict the radiation response in order to achieve radiotherapy goal like Tumor Control Probability & Normal Tissue Complication Probability. The goal of radiation therapy is to deliver a therapeutic dose of radiation to target tissues while minimizing the risks of normal tissue complications. The Lyman-Kutcher-Burman model proposed by Lyman. The Lyman-Kutcher-Burman model still suitable model for treatment plan evaluation & optimization. The most of treatment planning system like Eclipse, Monaco, Ray Station used Lyman-Kutcher-Burman based NTCP model including within the LQ formalism. The suitable model of NTCP calculation in treatment planning is Lyman-Kutcher-Burman model as it predicted almost same as the other complicated model & other model can be used when absolutely necessary. The five statistical parameters which were the poisson, binomial, probit, logit, and Weibull distributions have been used to investigate the different radiobiological models for NTCP in different articles.

**Conclusion**

From the above discussion evident that, the 15 literature reviews, most of the works have been used to the treatment plan evaluation. The treatment plans have been compared according to the biological evaluation. This paper has been proposed overall biological

evaluation tools, which give information both healthy tissue and target volume. In this study, there is a significant radiobiological modified treatment planning has been using the biological criteria. This study shows to make out the biological evaluation and make more accessible current radiobiological modeling knowledge, and may serve as a useful aid in the prospective and retrospective analysis of radiotherapy treatment plan.

In present study, Number of Models compared with six of these NTCP Models: The Critical, the Relative Seriality model & the Critical element model the Lyman-Kutcher -Burman model, the Parallel architecture, and the Weibull distribution model. In this study, Lyman-Kutcher-Burman NTCP model has been most used in treatment plan evaluation. The common parameters  $\gamma 50$ , M, N, D50, has been used in the treatment planning system. 3DCRT, IMRT, VMAT, and Helical Tomotherapy (HT), treatment planning system has been used to the radiobiological models. Radiotherapy planning software Philips, Pinnacle, Monaco, Eclipse, Raystation, has been used to calculate the biological responses in this study.

Still now, no standard protocol can be used for NTCP calculation due to problem of accurate NTCP model and absolute parameters. It should be more specific to maintain accurate use of published parameters values because the published data are available for many tumor sites and complication parameters in different models.

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