Deep BOO: Automating Beam Orientation Optimization in Intensity Modulated Radiation Therapy. Part I: Methodologies

Olalekan Ogunmolu*

Department of Radiation Oncology, UT Southwestern Medical Center, 2280 Inwood Road, Dallas, Texas , USA

olalekan.ogunmolu@utsouthwestern.edu

Abstract. Finding the optimal beam angle set from a robot's tool frame from which to aim radiation to cancer tumor(s) whilst simultaneously minimizing radiation intensity to organs-at-risks during intensity modulated radiation therapy is an exhaustive combinatorial search/human-based parameters-tuning process. So far, traditional optimization approaches for solving this problem have not yielded real-time feasible results. Aiming to automate the selection process, we introduce novel techniques from (i) pattern recognition, (ii) monte carlo evaluations, (iii) game theory, and (iv) neuro-dynamic programming. This is so as to find admissible policies for beamlets selection duringh IMRT treatment planning. In this sentiment, we optimize a deep neural network policy which guides monte carlo simulations and evaluations of promising beamlets within the robot gantry's phase space. Our goal is not to exclusively find the minimum of an objective as traditional approaches try to do, but rather to go uphill every so often so as to arrive at a pair of saddle point equilibrium strategy for the system. We let two fictitious neural players, within a zero-sum markov game framework alternatingly play a best response to their opponent's mixed strategy profile during epsidoes of a markov decision process. At inference time, we run the trained policy on test patients' target volumes to predict feasible beam angles. This part of our paper describes the problem and formally introduces our approach. The implementation details and results are presented in the addendum to this paper [1].

1 Introduction

Intensity Modulated Radiation Therapy (IMRT) is a state-of-the-art cancer treatment method that modulates the intensity of radiation delivered from a robot's beam eye-view and the geometric field-shape of the resulting beam in order to produce conformal dose distributions around a tumor in a patient lyingh in a supine position on a treatment table. Before actual treatment, a doctor contours the *critical structures* (or tumors) and *organs-at-risk* (or OARs) within a *target volume* (region of the patient's CT scan that contains the tumor and other organs), and then prescribes doses that must be delivered to the respective structures. Each beam is divided into beamlets, where each beamlet is aimed from the same angle; each beamlet in a beam may be of a different intensity from that of its neighbors – helping to achieve prescription dose for the target while sparing OARs. Generally, the radiation intensities are delivered from about 5-7 different beam

orientations with multiple collimator units. The process of determining what intensity (influence) to give each beamlet so as to meet the doctor's objective is termed *fluence map optimization*.

A key challenge in achieving the doctor's prescription request is finding (i) what beam angles and (ii) what beamlet intensities are best for delivering the required dose so that the tumor is eliminated while OARs are spared. This is the quintessential IMRT Treatment planning system (TPS) problem. A good "influence" radiation profile should maximize and minimize the dose delivered to tumor(s) and healthy organs respectively, and it should produce sharp dose gradients at the transition between tumors and normal tissues. Maximizing dose to tumors while simultaneously minimizing dose to critical structures poses a conflicted objective because it often occurs that the tumor(s) intersect(s) with some of the OARs. As such, finding good candidate beam angles in practice is often a trial-and-error process, with success depending on the dosimetrist/treatment planner's experience and the amount of time spent on tuning the angles. The process of selecting the best beam angles among all available beam orientations from which radiation is to be delivered in a treatment plan (TP), and optimizing the resulting chosen beams to fit a doctor's prescribed fluence profile is termed beam orientation optimization (BOO). Human intuition-based beam angle selection is laden with the subjectivity of the dosimetrist's judgment, not to mention the long time it takes in settling on a good set of candidate beam angles. Thus, having an automated framework for choosing beam angles is desirable since it will reduce the trial and error process, and save treatment time.

The second phase of the treatment planning process involves finding the curvature of beamlet intensities from which a beam can be shaped with respect to the geometrical shape and fluence across e.g. a high resolution multi-leaf collimator (MLC) field; this is so as to deliver a fixed or moving nonuniform photon beams or charged particles.

Solving for good candidate beam angles is a non-convex problem with many local minima (e.g. [2], [3]). It is at best a combinatorial optimization process – seeking to find an optimal set of beamlet candidates within the gantry space of a robot (see figure to the right), *i.e.* the workspace configuration and robot angle variables where the dynamics of the system is contained. When just the gantry (the robot's tool frame) of the linear accelerator (LINAC) machine is rotated in order to deliver beams to a patient, we have *coplanar beams*, *i.e.* all other joint angles of the robot are fixed while only the head rotates – resulting in a



IMRT TPS setup. ©Cyberknife.

set of coplanar beams that are swept out by the gantry. In this paper, we focus on finding good coplanar beams in BOO problems as more often than not, only coplanar beams are employed when delivering radiation.

Given the different ways a set of beamlets can be chosen and how a beam can be shaped with respect to the geometrical shape and fluence field, the dimensionality of the parameters to be optimized using traditional optimization techniques increase to the extent that such methods are not real-time feasible. This is coupled with the nonconvexity of the solution surface of the BOO process as the physics of dose deposition changes with each beam orientation. Therefore, we resort to an approximate dynamic programming (ADP) formulation to derive approximately optimal control laws/policies in high dimensional state spaces. Particularly, we leverage on recent machine learning breakthroughs [4] to guide a monte carlo tree search [5–7] of promising beam angle candidates by rapidly exploring different parts the beam space.

1.1 Contributions

Finding the right combination of beam angles within the configuration space of the gantry is a hard computational problem given the high branching factor during exploration, tree depth, and complexity of finding a good known heuristic value function for non-terminal beam angle positions. This is the first article, to the best of our knowledge, that provides a comprehensive description of Monte-Carlo tree search within the framework of BOO. It adds new pseudocode that transforms a BOO problem into a game planning strategy. Coupled with neural fictitious self-play, we continually refine the predictions from a neural network policy so as to drive the weights of the network to a **saddle equilibrium**. This is essentially a minimax zero-sum two player markov game that aims for robust policies (e.g. see [8]); To this end,

- we use a deep neural network to model the nonlinear dynamical system and generate policies that guide MCTS simulations for two players in a zero-sum Markov game
- we formulate an MCTS tree lookout strategy that guides transition from one beam angle set to another during each MDP episodic setting for either player
- each player seeks to find an alternating best response strategy to the current player's average strategy driving the weights of the network policy toward an approximate saddle equilibrium [9]

1.2 Related Work

There have been attempts at providing a holistic mathematical formulation for the choice of feasible beam angles from which a good clinical plan can be achieved. For example, Craft [2] locally tuned a set of beam angles within a beam's continuous state space using linear programming duality theory and gradient descent. They reported that the BOO problem for a simple 2D pancreatic case has many local minima (\approx 100) when the local optimization is warm-started from 100 different beam angles.

Building on Craft's work, Bertsimas et. al [10] resolved to a two meta-step algorithm: the BOO problem is solved by dynamically selecting beam angle candidates (based on the linear accelerator machine precision) within the continuous beams' phase space via a local minimum search with gradient descent, and then explores a different portion of the solution space by taking finite steps of simulated annealing. Bertsimas et. al [10] define a linear programming problem with constraints that capture a doctor's preference for dose delivery. While this approach takes advantage of global and local search with improvements in solutions obtained from equispaced angles, it has the drawback of presenting the objective function as convex and assuming the doctor's preferences can be represented as a set of linear constraints.

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Jia et. al [11] split the BOO problem into two consecutive phases: first, by progressively identifying non-consequential beam angles through the evaluation of a multi objective function, and secondly, by optimizing the fluence on beam angles that are assigned a high confidence by a dosimetric objective component. Heuristic search strategies have also previously developed e.g. [12–14]. Other lines of work have treated IMRT treatment planning as an inverse optimization problem, with techniques ranging from adaptive l_{21} optimization [11], mixed integer linear programming [15–17] and simulated annealing [18, 19].

1.3 Notations and Definitions

First, let us formalize definitions and notations used throughout the rest of this article. The state of the dynamical system, will be denoted by $s \in \mathcal{S}$, and it is to be controlled by a

Notation	Definition/Examples	Notation	Definition/Examples
m	dimensionality of a node's beam set, e.g. $m=5$	n	dimension of discretized beam angles, e.g. $n=180$ for 4° angular resoultion
Θ	discretized beam angle set e.g. equally spaced angles between 0° and 360° , spaced apart at 4°	$a_t \in \mathcal{A}$	control or action, $a_t \in \mathcal{A}$ at time step $t \in [1, T]$ used in determining the probability of transitioning from a beam angle subset to another within $\boldsymbol{\Theta}$
$ heta^j \subseteq oldsymbol{arTheta}$	beam angles selected from $oldsymbol{\Theta}$ e.g. $ heta_k \in \mathbb{R}^m$	$s_t \in \mathcal{S}$	markovian system state at time step, $t \in [1,T]$ e.g. patient contour, beam angle candidates; dimensionality $2,727,936$ to $3,563,520$
γ	discount factor e.g. 0.99	f_{ψ}	parametric function approximator (deep neural network policy) for state s_t
$v_{\psi}(s)$	value estimate of state, s_t , as predicted by $m{f}_{m{\psi}}$	p(s)	probability distribution over current state, <i>s</i> generated by neural network policy
Q(s,a)	action-state values that encode the "goodness" of a beam-angle set, $\theta_k \in \mathbb{R}^m$, where m is the number of beams considered for a fluence generation, e.g. $m=5$	B_{X_t}	a concatenation of beams in consideration at time step, t , as a block of beams being fed to the neural network policy
$\mathcal{D}_{ij}(heta_k)$	dose matrix containing dose influence to voxel i from beam angle, θ_k , $\forall k \in \{1, 2,, n\}$ where n is range of the beam set \mathcal{B}		dose mask for target volume in consideration at time step , t

Table 1: Table of notations commonly used in this article

discrete action $a \in A$. States evolve according to the (unknown) dynamics $p(s_{t+1}|s_t, a_t)$, which we want to learn. We pose the learning problem within a discrete-time finite-time horizon, T.

A beam angle combination search task in this setting can be defined by a reward function $r(s_t, a_t)$ which can be found by recovering a policy $p(a_t|s_t;\psi)$ which specifies a distribution over actions conditioned on the state and parameterized by some vector ψ . The parameters, ψ , might correspond to the weights of a neural network or parameters of a Gaussian process, for example. Recovering the optimal weights may consist of the maximization problem,

$$\psi^{\star} = \arg\max_{\psi} \sum_{t=1}^{T} \mathbb{E}_{(s_t, a_t) \sim p(s_t, a_t | \psi)} \left[r(s_t, a_t) \right],$$

where the optimization problem is to determine an optimal vector ψ^* that maximize the sum of the respective instantaneous rewards $\sum_t r(s_t, a_t)$ of the policy, $p(a_t|s_t, \psi)$. For clarity, we will denote the action conditional $p(a_t|s_t, \psi)$ as $\pi_{\psi}(a_t|s_t)$. Other notations used in the article are delineated in table 1.

2 Methods and Materials

For games with perfect information, there is an optimal value function, $v^*(s)$, that decide the game's outcome for every possible state under perfect play (*i.e.* a game with an optimal min-max state-action value function, Q(s, a), produced by a policy $\pi = \{\pi_{min}, \pi_{max}\}$. It follows that one can devise a planning strategy that guides the search for optimistic beam angle configurations within the beam phase space by using a probability distribution, p(s, a), over a set of deterministic *pure strategies* for the tree [9].

2.1 Overview

The search for an *approximately* optimal beam angle set is performed by optimizing the parameters of a function approximator, which in this case are the parameters of a deep neural network, ψ . The parameters approximate a policy, π which repeatedly guides episodic MCTS simulations of 'best-first' beam angle combinations. Essentially, the MCTS performs a sparse lookout simulation, recursively expanding the sub-nodes (consisting of children beam angle sets) under a tree's root node (the starting beam angle set), and selectively adjusting beam angle(s) that contribute the least to an optimal fluence within a beam angle set. Successor nodes beneath a terminal node are **approximated** with a value, v(s), to assure efficient selectivity.

We model the decision process for optimal beam selection by maintaining a probability distribution over the set of possible states, based on a set of observations and observation probabilities for the underlying Markov Decision Process. The state of the system, s_t , is markovian containing the concatenation of the patient ct mask as well as a subset of the beams in consideration at the current time step t. The policy that guides the search for the optimal beam angle selection is computed by a deep neural network policy, consisting of multiple residual blocks fashioned similar to modern neural

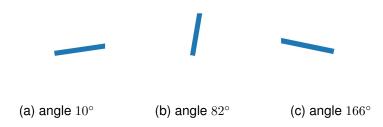


Fig. 1: Example 2D gantry angle representations

network architectures [20,21]. Each beam angle is represented as gantry angle images (see Fig. 1). Depending on the patient in question, the state at time step, t, varies between $2,700,000 \sim 3,600,000$ in number of parameters.

2.2 Data Preprocessing

This section describes the data preparation for the masks that we use in training the network.

Patient Mask. We obtained 77 anonymized patient CT scans and associated data from our clinic. The scans relate to prostate cases used in previous IMRT cancer treatment planning. The prostate data contains scans of six organs, namely the patients' body, bladder, left and right femoral heads, rectum, and planning target volume (PTV) or tumor. Each patient's scan is represented in 3D as $N \times W \times H$, where N is the total number of slices, W is the image width and H is the image height. We resized each slice within each patient's ct mask, D, to a square-shaped 2D matrix of size $N \times 64 \times 64$. **Beam Orientation Representation:** We discretized the gantry angle space at a 4° grid resolution from 0° through 360° . Since the beam angles and patient's geometry are spatially related, we use deep neural networks consisting of successive layers of convolution [22] and roly position for the proposition [23] to encode the

grid resolution from 0° through 360° . Since the beam angles and patient's geometry are spatially related, we use deep neural networks consisting of successive layers of convolution [4], batch normalization [22], and relu nonlinearities [23] to encode the spatial relationship between the pose of the patient from the gantry's eye view. We generate 3D images that represent the orientation of the gantry with respect to the patient for each discretized beam angle as shown in (Fig. 1).

2.3 Neural Network Architecture

In addition to the resized masks, D, we define five feature planes, X_t as a block of beam configurations: B_{X_t} , denotes the beam angles that generate the current fluence. For example, if we are considering five beams for the geometric shape for the fluence, B_{X_t} would contain the 3D RGB images of the beams being considered at time step t. We then augment the state with a memory of five previously used beam blocks, $\{B_{X_t}, \ldots, B_{X_{t-5}}\}$, during the search process; we make this augmentation due to the partial observability of the patient-gantry setup from a single set of features, in order to mitigate the uncertainty of decisions under our POMDP formulation.

The dose masks and beam blocks are as shown in Fig. 2 so that the input planes to the network are sized as $T \times N \times H \times W$; T is the total number of input planes

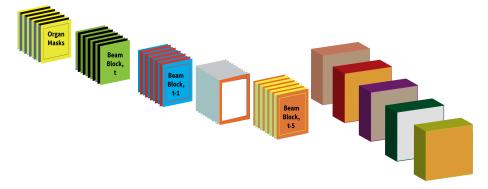


Fig. 2: [Left]: Concatenation of the target volume masks and the beam angles before feeding the input planes to the residual tower neural network. The first six planes (top-most mask of left figure) contain the delineated organs and the PTV. This is concatenated with a block of m beams from the current time step, regressed to the previous 5 time steps (here, 5 was heuristically chosen). [Right] Each beam angle in a beam block is represented as shown. Together with the target volume, these form an input plane of size $36 \times N \times W \times H$ to the policy/value neural network tower of residual blocks.

 $(T=6 \text{ structures} + 5 \text{ beams} + 5 \times 5 \text{ regressed beams})$. Thus, the input to the network are arranged as: $s_t = [D_t, B_{X_t}, B_{X_{t-1}}, B_{X_{t-2}}, B_{X_{t-3}}, B_{X_{t-4}}, B_{X_{t-5}}]$. To address robust learning of patient geometry and beam angle orientations, we use a modern neural network architecture with many residual blocks [21]; each layer of the network fits a residual nonlinear mapping to its input data; we end up with a deeply stacked network whose input features, s_t , are processed by 34 residual blocks described as follows: (i) a 3D convolution with $64 \times l$ filters, a square kernel of width 7, and double strided convolutions, where l is the depth of the stack in the network; (ii) a 3D batch normalization layer [22]; (iii) nonlinear rectifiers [23]; (iv) a 3D convolution of $64 \times l$ filters; (v) a 3D batch normalization layer; (vi) a skip connection from the input to the block, in order to facilitate efficient gradients' propagation; and (vii) nonlinear rectifiers.

We split the output of the network into two heads: (i) the first head is a probability distribution over which angle in the current beam block contributes the least to an optimal fluence cost at the current time step, while (ii) the second head estimates the *value* (in an *ADP sense*) of the subtree beneath the current node This probabilty head is comprised of two residual blocks, each containing: (i) a 3D convolution of $64 \times l$ filters, followed by a square kernel of size 1, and a single strided convolution; (ii) a 3D batch normalization layer; (iii) nonlinear rectifiers; (iv) a 3D convolution of $64 \times l$ filters (v) a 3D batch normalization layer; (vi) a skip connection from the input to the block (vii) nonlinear rectifiers (viii) a fully connected layer that maps the resulting output to the total number of discretized beam angle grids; and (ix) a softmax layer then maps the neuron units to logit probabilities $p_i(s|a)$ for all beam angles.

The value head applies the following transformation modules (i) a 3D convolution of $64 \times l$ filters, a square kernel of size 1, and a single strided convolution; (ii) a 3D batch normalization layer; (iii) nonlinear rectifiers; (iv) a second 3D convolution of $64 \times l$

filters; (v) a 3D batch normalization layer; (vi) a skip connection from the input to the block (vii) nonlinear rectifiers and a sequential layer consisting of:

- a linear layer that maps the output of the above connections to a 512-layer hidden unit
- a linear layer that maps the output of the above connections to a 256 hidden unit followed by rectified nonlinearities
- a linear layer that maps to a scalar value, and followed by rectified nonlinearities
- a tanh nonlinearity that maps the output to the closed interval [-1, 1].

The parameters of the neural network were initialized using the initialization method proposed in [24]. Since decisions about what beam angle combination is optimal at the current time step are made under uncertainty, we use a *probability model* to choose among **lotteries**, where the lotteries in this case are probability distributions over all discretized beam angles in our setting. Each state during our learning process is constructed by appending the beam block at the current time step to a history of beam blocks for the previous five time steps. We append a new beam block to the current set of beam blocks and target volume dose mask (see Fig. 2) with a FIFO policy so that we minimize the partial observability of our MDP system. When we transition to a new state, the beam block that has been in the state set for the longest time (*i.e.* at the **head** of the queue) is deleted first, and the new state's beam block is enqueued at the tail as in a **queue** data structure. Let us now introduce some definitions.

Definition 1. A beam block is a concatenation of beamlets, $\{\theta_1, \theta_2, \dots, \theta_m\}$ as a tensor of dimension $m \times C \times H \times W$ (see Fig. 2) that together with the patient's ct mask form the state s_t at time step t.

Definition 2. Every **node**, x has the following fields: (i) a pointer to the parent that led to it, x.p; (ii) the beamlets, x_b , stored at that node where $b = \{1, \ldots, m\}$; (iii) a set of move probabilities prior, p(s, a); (iv) a pointer, x.r, to the reward, r_t , for the state s_t ; (v) a pointer to the state-action value Q(s, a) and its upper confidence bound U(s, a) (6) (vi) a visit count, $\mathbb{N}(s, a)$, that indicates the number of times that node was visited; and (vii) a pointer x.child; to each of its children nodes.

In order for the state definition to capture as much information as possible under uncertainty, we define the state broadly enough to capture all subjective unknowns that might influence the payoff/reward to be received by a rational decision-making agent. We want an adaptive allocation rule for determining the transition between states such that as the search process progresses, and achieves convergence as the simulation grows, the worst possible bias is bounded by a quantity that converges to zero. We leverage on the *upper confidence bound* algorithm of Agrawal [25] that assures an asymptotic logarithmic regret behavior. Since we do not know what node may yield the best bandit, a player might be biased towards always selecting the beams set with the maximum value. Therefore, we attach a regret term, U(n(s)) to the Q(s,a)-value so as to ensure the optimal beam does not evade the simulation i.e. $Q(s,a) - U(n(s)) \le Q(s,a) \le Q(s,a) - U(n(s))$; the width of this confidence bound guides the exploration strategy for states that are momentarily unpromising in values but may later emerge as promising states as the number of games evolve in length.

2.4 Fluence Map Optimization

Suppose \mathcal{X} is the total number of discretized voxels of interest (VOI)'s) in a target volume, and $\mathcal{B}_1 \cup \mathcal{B}_2 \cup \ldots \cup \mathcal{B}_n \subseteq \mathcal{B}$ represents the partition subset of a beam \mathcal{B} , where n is the total number of beams from which radiation would be delivered. Suppose we let $\mathcal{D}_{ij}(\theta_k)$ be the matrix that describes each dose influence, d_i , delivered to a discretized voxel, i, in a volume of interest, VOI_h $(h=1,\ldots,\mathcal{X})$, from a beam angle, θ_k , which is within the beamlet $k \in \{1,\ldots,n\}$, then it follows that one can compute the matrix $\mathcal{D}_{ij}(\theta_k)$ by calculating each d_i for every bixel, j, at every φ° , $j=1,\ldots,k$ resolution. Doing this, we end up with a large but sparse matrix, $\mathcal{D}_{ij}(\theta_k)$, which consists of the dose to every voxel, i, incident from a beam angle, θ_k at every $360^\circ/\varphi^\circ$ (in our implementation, we set φ to 4°).

Let us introduce a decision variable, x_j , which represents the intensities of beamlets, $j \in \mathcal{B}$; then it is trivial to find the dose influence, d_i , that relates the bixel intensities, x_j , to the voxels of interest, VOI_h . The fluence problem aims to find the values of x_j for which the d_i to the tumor is maximized, while simultaneously minimizing the d_i to critical structures. These objectives are conflicted by definition, and it is typical for dosimetrists to spend hours before finding a good set of candidate beams. For the voxels in all OARs, a weighted quadratic objective minimizes the l_2 distance between a pre-calculated dose Ax, and a doctor's prescribed dose, b, while a weighted quadratic objective maximizes the l_2 distance between Ax and b. The pre-calculated dose term is given by $Ax = \{\sum_s \frac{w_s}{v_s} \mathcal{D}_{ij}^s x_s \, | \, \mathcal{D}_{ij} \in \mathbb{R}^{n \times l}, n > l \}$, which is a combination of the dose components that belong to OARs and those that belong to PTVs. $w_s = \{\underline{w}_s, \overline{w}_s\}$ are the respective underdosing and overdosing weights for the OARs and PTVs, and v_s represents the total number of voxels in each structure. The cost function is

$$\frac{1}{v_s} \sum_{s \in \text{OARs}} \|(b_s - \underline{w}_s \mathcal{D}_{ij}^s \mathbf{x}_s)_+\|_2^2 + \frac{1}{v_s} \sum_{s \in \text{PTVs}} \|(\bar{w}_s \mathcal{D}_{ij}^s \mathbf{x}_s - b_s)_+\|_2^2$$
(1)

where the underdosing weights are typically set as $\underline{w}_s=0$ since it is desirable to deliver as little dose as possible to critical structures, while the overdosing weights are chosen to deliver the prescribed dose to the tumor; $(\cdot)_+$ denotes a Euclidean projection onto the nonnegative orthant \mathbb{R}_+ . We can succintly rewrite the objective function above, subject to nonnegative bixel intensity constraints, as

$$\min \frac{1}{2} ||A\mathbf{x} - \mathbf{b}||_2^2 \quad \text{ subject to } x \ge 0,$$

with the attraction of being differentiable and convex, penalizing dose volumes that exceed a doctor's prescribed constraint. The Lagrangian becomes

$$L(\boldsymbol{x}, \boldsymbol{\lambda}) = \min \frac{1}{2} \|A\boldsymbol{x} - \boldsymbol{b}\|_2^2 - \boldsymbol{\lambda}^T \boldsymbol{x}.$$

The above problem can be solved with dual gradient descent (DGD) but DGD has the drawback that the primal and dual updates are not robust to objective's constraints [26]. The alternating direction method of multipliers (ADMM) [26] tackles the robustness problem by adding a quadratic penalty term to the Lagrangian and alternatingly updating

the x and λ variables in a "broadcast and gather" process. This turns out to be attractive since we will be solving a large-scale learning problem for the optimal beam angle set combination. Introducing an auxiliary variable z, the above is transformed into the following

$$\min_{\mathbf{x}} \frac{1}{2} \|\mathbf{A}\mathbf{x} - \mathbf{b}\|_{2}^{2}, \text{ subject to } \mathbf{z} = \mathbf{x}, \ \mathbf{z} \ge 0.$$

so that the Lagrangian can be written as,

$$\min_{\mathbf{x}, \mathbf{z}} \frac{1}{2} \| \mathbf{A} \mathbf{x} - \mathbf{b} \|_{2}^{2} - \lambda^{T} (\mathbf{z} - \mathbf{x}) + \frac{\rho}{2} \| \mathbf{z} - \mathbf{x} \|_{2}^{2}, \tag{2}$$

where $\lambda \in \mathbb{R}^n$ is a multiplier and $\rho > 0$ is an ADMM penalty parameter. The x subproblem to (2) yields

$$\min_{\mathbf{x}} \quad \frac{1}{2} \mathbf{x}^T (\mathbf{A}^T \mathbf{A} + \rho \mathbf{I}) \mathbf{x} + (\mathbf{\lambda}^T - \mathbf{A}^T \mathbf{b} - \rho \mathbf{z}^T) \mathbf{x},$$

so that the x-update (due to the convex quadratic nature of the problem) becomes,

$$\mathbf{x}^{k+1} = \left(\mathbf{A}^T \mathbf{A} + \rho \mathbf{I}\right)^{-1} \left(\mathbf{A}^T \mathbf{b} + \rho \mathbf{z}^k - \boldsymbol{\lambda}^k\right). \tag{3}$$

Similarly, the z-update for (2) can be found by the z-minimization subproblem

$$\min_{\mathbf{z}} - \boldsymbol{\lambda}^T \mathbf{z} + \frac{\rho}{2} \|\mathbf{z} - \mathbf{x}\|_2^2 := \min_{\mathbf{z}} \frac{\rho}{2} \|\mathbf{z} - \mathbf{x} - \frac{1}{\rho}(\boldsymbol{\lambda})\|_2^2.$$

Using the soft-thresholding operator, $S_{\lambda/\rho}$, we find that

$$\boldsymbol{z}^{k+1} = S_{\boldsymbol{\lambda}/\rho} \left(\boldsymbol{x}^{k+1} + \boldsymbol{\lambda}^k \right), \tag{4}$$

where $S_{\lambda/\rho}(\tau) = (x - \lambda/\rho)_+ - (-\tau - \lambda/\rho)_+$. λ is updated as

$$\lambda^{k+1} = \lambda^k - \gamma(z^{k+1} - x^{k+1}),\tag{5}$$

where γ is a parameter that controls the step length. The inverse operation in (3) can be carried out with any iterative solver e.g. conjugate gradient. We use an over-relaxation parameter, $\alpha^k=1.5$, and set the quadratic penalty to $\rho=1.5$, in the z and λ updates: $\alpha^k A \, x^{k+1} - (1-\alpha^k) z^k$. The stopping criterion is met when the primal and dual residuals are sufficiently small, i.e.,

$$r^k = \| \mathbf{x}^k - \mathbf{z}^k \|_2 \leq \epsilon^{\mathrm{pri}} \ \ \text{and} \ \ s^k = \| - \rho \left(\mathbf{z}^{k+1} - \mathbf{z}^k \right) \|_2 \leq \epsilon^{\mathrm{dual}},$$

with,

$$\epsilon^{
m pri} = \sqrt{
ho}\epsilon^{
m abs} + \epsilon^{
m rel} \max\{\|m{x}^k\|_2, \|-m{z}\|_2\}, \ ext{and}$$

$$\epsilon^{
m dual} = \sqrt{n}\epsilon^{
m abs} + \epsilon^{
m rel}(
hom{\lambda}^{m{k}}),$$

where $\epsilon^{\rm pri} > 0$, $\epsilon^{\rm dual} > 0$ are the primal and dual feasibility tolerances for the primal and dual feasibility conditions (see [26, §3.3]). We set $\epsilon^{\rm abs} = 10^{-4}$ and $\epsilon^{\rm rel} = 10^{-2}$.

2.5 Game Tree Simulation

Our tree-search policy is based on a sparse look-ahead search strategy for large state-space Markov Decision Problems (MDPs). The game simulation is one of **perfect recall** *i.e.* each node's current information state implies knowledge of the ancestral nodes that led to it.

The algorithm can be seen as a set of sequential simulation of different beam angle combinations guided by probabilities obtained from a two-player zero-sum game of neural fictitious self-play (FSP); the network roll-out policy is used to guide the search for a *best-first* set of beam angle candidates. The best-first leaf node encountered is given by the child node with the highest reward along a tree that is expanded. What follows is a set of sampling-based lookout algorithm that focuses learning on regions of the state space that are likely to have a good fluence profile and computes action-value pairs based on selectively sampled angles.

We randomly sample beam angles within the gantry angle space by carrying out a lookahead search at fixed depth; we restrict sampling to 90 beam grid in Θ at every stage of our simulation. A partial tree of beam angle candidates is randomly constructed, starting from a root node, and we progressively add child nodes to the current node in a **selection** process guided by move probabilities, p(s,a). At the first iteration, the subset of beam angles are randomly sampled from Θ . In our expansion phase (see algorithm 1), to prevent angles from being too close to one another, we define a minimum pairwise distancem, $\bar{d}_i \in \mathbb{R}^+$ between the beamlets in a beam block, given by $\|\theta_i - \theta_j\| \ge \bar{d}_i$, $\forall \{j \in m \setminus i\}$ where we set $\bar{d}_i = 20$. By repeatedly performing roll-outs, we maintain a history of state-action value pairs, stored along the edges of the tree. This aids faster convergence if the same state is encountered more than once: we can bias an action selection based on old actions that were chosen.

When we finish each simulation, a 'best move' for the current beam block is selected, as computed by the tree search; we exponentiate the move probabilities by a temperature slightly larger than unity to encourage diversity in early play; specifically, we compute

$$p(a|s_0;\psi) = \frac{N(s_0,a)^{1/\tau}}{\sum_b N(s,b)^{1/\tau}},$$

where τ is the temperature factor that diversifies the move probabilities. The UCT [7] algorithm applied to optimal beam angle selection is presented in algorithm 1.

Definition 3. A minimizer player is at a **terminal state** if the FMO cost for the beams at a leaf node is greater than the cost of the beams at its direct ancestor node.

Definition 4. A maximizer player has reached a **terminal state** if the FMO cost for the beams at a leaf node is less than the FMO cost for the beams at its direct ancestor node.

Definitions § 3 and § 4 help preserve the neighborhood search for the beam improvement tuning as specified earlier. When angles are at the edges i.e. 0° or 360° and an angle change outside the range $0 \le \theta \le 360$ is recommended, we "wrap" around to enforce cyclicity. The index of the angle to select is determined by taking an argmax of the mixed strategy produced by the tree policy $\pi_{\psi_{t-1}}$; this policy is constantly being

Algorithm 1 Deep BOO Monte Carlo Tree Search

```
return False
function MCTS(s_0)
                                                               end if
    s_0 \leftarrow \boldsymbol{x}_0(s_0)
                                                          end function
    while search time < budget do
         \bar{x} \leftarrow \text{EXPAND\_POLICY}(x_0)
         \bar{x}.r \leftarrow \text{FMO\_POLICY}
                                                           function FMO\_POLICY(x)
         BACKUP(\bar{x}, \bar{x}, r)
                                                               find optimal fluence objective,
    end while
                                                           h^{\star}(\boldsymbol{x}(s)|\cdot)
    return BEST_CHILD(x_0, c)
                                                               return r = -h^*(\mathbf{x}(s)|\cdot)
end function
                                                          end function
function SELECT_MOVE(x, c)
                                                           function EXPAND(x)
    if p_1 to play then
                                                               sample a \bar{\theta} using prior \boldsymbol{x}.p(s,a)
         return \operatorname{argmax}_{\bar{x} \in \operatorname{children of } x} Q(\bar{x}) +
                                                               \bar{a} = SELECT\_MOVE(x, c)
                                                               add \bar{a} to tried actions \bar{\mathcal{A}}
     2 \ln n(\bar{\boldsymbol{x}}.s)
                                                               form new node \bar{x}.\bar{s}
    \overline{N(\bar{x}.s,a)}
                                                                  with \bar{\theta} \leftarrow \bar{\theta} + \bar{a}
                                                               with \pi_{t-1}, create \bar{x}.p(\bar{s},\bar{a}), \bar{x}.\bar{v}(\cdot)
         return arg \min_{\bar{x} \in \text{ children of } x} Q(\bar{x}) –
                                                               while not \bar{x} in children of x_0 do
    2 \ln n(\bar{x}.s)
                                                                     add child \bar{x} to x
                                                               end while
    end if
                                                               return \bar{x}
end function
                                                           end function
function EXPAND_POLICY(x)
                                                          function BACK_UP(x, c)
    while x is nonterminal do
                                                               while \bar{x} is not null do
         if x is not fully expanded then
                                                                     N(\bar{x}) \leftarrow \bar{x} + 1
              return EXPAND (x)
                                                                     Q(\bar{x}) \leftarrow Q(\bar{x}) + \bar{x}.r
         else
                                                                    \bar{x} = \text{ parent of } \bar{x}
              x \leftarrow \text{BEST\_CHILD}(x, c)
                                                               end while
         end if
                                                           end function
    end while
    return x
                                                          function BEST_CHILD(x_0)
end function
                                                               if p_1 to play then
                                                                     return x_0[\arg\min \text{ children of } x_0.r]
function FULLY_EXPANDED(x, d)
    d_i \leftarrow \text{pairwise\_distance}(\boldsymbol{x}.s)
                                                                     return x_0[\arg\max \text{ children of } x_0.r]
    \min_{e} = \min(d)
                                                               end if
    if min_elem < d then
                                                          end function
         return True
    else
```

optimized in a separate thread. Similarly, the direction in which to modify the angle is determined by an argmax of the third head of the tree policy $\pi_{\psi_{t-1}}$ as discussed in § 2.3. The **EXPAND_POLICY** and **FMO_POLICY** procedures of the MCTS simulation in algorithm 1 can be seen as a form of Add/Drop simulated annealing as described in [19]. The **BEST_CHILD** procedure compares the quality of all beam angle sets traversed as the root node is expanded to one or more child nodes; the greedy approach of selecting the beam angle set with the highest reward results in a locally optimal set of beam angles candidates at each iteration of running a game.

Definition 5. We define an **upper confidence bound**, U(s,a), on Q(s,a) that adds an exploration bonus that is highest for seldomly visited state-action pairs so that the tree expansion policy selects the action a^* that maximizes the augmented value:

$$\bar{Q}(s,a) = Q_j(s,a) + c\sqrt{\frac{2 \ln n(s)}{\mathbb{N}(s,a)}}, \quad \text{where } a^* = \arg\max_a \bar{Q}(s,a). \tag{6}$$

Q(s,a) is the highest average observed reward from node j – encouraging exploitation of the current node, and $\ln n(s)$ is the natural logarithm of the total number of rollouts through state s. The second term in the (6) encourages exploration of other beam angles and c is a scalar exploration constant. Note that (6) is a a version of the UCB1 algorithm [7]. This estimates the expected reward of a set of beam canditates once we compute its upper confidence.

2.6 Self-Play Neuro-Dynamic Programming

This section describes the approximate dynamic programming scheme and self-play formulation for generating saddle point equilibrium strategies for an approximately optimal outcome (value) by playing a weakened fictitious self-play (FSP) between two neural network behavior strategies. We consider a trial-and-error search problem via interaction with the dynamical system so as to discover a mapping between poatient geometry and beam angles in order to maximize an extrinsic reward signal; we approach this with optimal control of incompletely known Markov decision processes (MDP). The MDP consists of states S, actions, A, a transition probability, $\mathcal{P}_{ss'}^a$ and a reward function \mathcal{R}^a_s . $\mathcal{P}^a_{ss'}$ is a probability distribution that governs the evolution of states e.g. from $s \to s'$, while \mathcal{R}_s^a determines a reward after the transitions occur. In **Q-learning**, an agent's experience is made up of distinct episodes: at the t'th episode, the agent observes its current state, s_t , selects and executes an action a_t , which transitions the agent to a new state s_{t+1} ; an immediate reward, r_t , is subsequently given to the agent, by which the previous Q-values are adjusted. Q-learning relies on *value iteration* to reach convergence. In practice, this takes a long time to complete. **Batch Q-learning** algorithms e.g. [27] attempt to learn a stochastic policy $\pi_{\psi}: \mathcal{S} imes \mathcal{A} o \mathcal{R}$ that maximizes the cumulative discounted reward $\sum_{t=0}^{T-1} \gamma^t r(s_t, a_t)$, where ψ are the parameters of a policy.

Weakened fictitious self-play applies machine learning to determine an *approximate* best response strategy to an opponent's mixed strategy in games of self-play. An agent continually plays a game against an opponent and the outcome of the game is given by averaging the outcome of all individual plays.

We apply FSP in a machine learning framework to a game's strategy profile so that each player plays an *approximate* best response strategy to their opponent's mixed strategy at each iteration. In our formulation, each player does not necessarily know the strategy of its opponent ahead of time *i.e.* their security levels do not necessarily coincide. To ensure that equilibrium can be found within pure strategies, we let one player act after observing the decision outcome of the other player. The game thus has a **dynamic** property since the strategy of the successive acting player explicitly depends on that of the first player [28, pp.22].

Each player's action is governed by a mixed strategy – obtained by adding a Gaussian random walk sequence with standard deviation 2 to the prior probability distribution predicted by the neural network policy or computed by the tree policy; this is then normalized by the sum of the resulting noised distribution. We do this to encourage further exploration and to transform each player's pure strategy to a mixed strategy. As such, a player's mixed strategy is simply a random variable whose values are the player's pure strategies. Players p_1 , and p_2 's mixed strategies are independent random variables, repeatedly implemented during game simulations such that their actual decisions are based on chance events. However, as the number of times the game is played gets larger, the frequency with which different actions for p_1 and p_2 are chosen will converge to the probability distribution that characterize their random strategies [28, pp.24]. For a game Γ , suppose that $y = \{y_1, \dots, y_m | \sum_{i=1}^m y_i = 1\}$ and $z = \{z_1, \dots, z_n | \sum_{i=1}^n z_i = 1\}$ are the respective probability distributions for players p_1 and p_2 , defined on the n and m—dimensional simplices respectively, the average value of the game will correspond to player p_1 minimizing a cost $\mathcal{J}(y,z) = y^T \Gamma z$ and player p_2 maximizing $\mathcal{J}(y,z)$.

The fictitious self-player samples from previous episodes of the game of self-play at each iteration k. For each *pure strategy*, p(s,a), predicted by the neural network policy, f_{ψ} , each player makes a decision governed by a probability distribution on the space of its pure strategies. The opposing agent's policy is randomly sampled from a previous checkpoint of the training scheme. The goal of the two-player zero-sum neural fictitious self-play is to drive the network to an approximate saddle equilibrium, via experiential learning, thus guaranteeing that the realization of an approximate optimal value function for the optimal beam angle set that generates a desired fluence/DVH profile.

3 Conclusion

Beam orientation optimization is a key component of conformal IMRT radiotherapy TPS. It has a nonconvex solution surface given the way the dose coefficients can significantly alter based on the gantry angle from which beams are aimed to a target volume. As such, in modern clinics, this problem is typically solved with many hours of planning, hand-tuning and refinement – usually by experienced dosimetrists and radiation therapists.

Given the long time traditional optimization approaches take in solving this problem, we adopt a hybrid approach, leveraging on monte carlo simulation of games in high dimensional state-spaces, neuro-dynamic programming, convex optimization of fluence profiles of a target radiation, and game theory to arrive at good candidate beamlets. Our monte carlo tree search formulation is the first, to the best of our knowledge, that transforms the BOO problem into a monte carlo tree search strategy; and provides a

pseudocode that shows how we implement the tree search to navigate the complex state space of respective beamlets. In the part II of this paper, we elucidate on the end-to-end optimization pipeline and provide our initial results based on a set of numerical simulations for select anonymized patients data in our clinic.

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