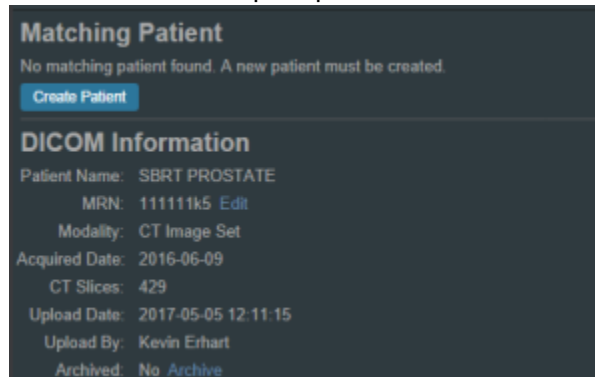


# Prostate Plan Walkthrough

Undergoing maintenance, please check back soon.

## Patient Creation

1. Open the Astroid Launcher and login
2. Select your realm
3. A list of available apps will be listed on the right, select *Planning* from this
4. If you see an Install button click it and wait for the version to install. You will know it is installed when you see a LAUNCH button
5. Click the LAUNCH button. The version of planning that is installed in your realm will now open
6. Click on the Imports
7. Select the CT image set (SBRT Prostate) from the list of available files for import
8. Ensure that the MR is correct
  1. If MR needs to be changed you may edit it by choosing the *Edit* button to the left
9. Click the *Create Patient* button to start the import process



**Matching Patient**

No matching patient found. A new patient must be created.

[Create Patient](#)

**DICOM Information**

Patient Name: SBRT PROSTATE

MRN: 111111k5 [Edit](#)

Modality: CT Image Set

Acquired Date: 2016-06-09

CT Slices: 429

Upload Date: 2017-05-05 12:11:15

Upload By: Kevin Erhart

Archived: No [Archive](#)

10. In the control pane on the left hand side, the image snapshot will be automatically selected based on the structure set DICOM UID information.
11. Fill in the Patient Intent information and select the appropriate HU to RSP curve (as shown below)

**Intent Data**  
Type \*: Curative  
Label: intent\_curative\_2017Jan4  
Treatment Site \*: Prostate\_Protocol  
Narrative:  
Protocol: 123456 New Protocol  
Body System \*: Male Reproductive System  
Body Part \*: Prostate  
ICD-10 Code: DV004ZZ

**RSP Data**  
HU to RSP Curve\*: Scanner1; 120 kVp; FOV [0, 1000]

**Import Structure Set**  
Structure Set: Date: 2016-06-09

**Import Structures**  
Treatment Site: Prostate\_Protocol  
Patient Structure: Skin  
Variant Label: variant\_1\_2017Jan4  

<input checked="" type="checkbox"/> Bladder neck (custom)	Importing new structure
<input checked="" type="checkbox"/> Testes (custom)	Importing new structure
<input checked="" type="checkbox"/> 2cm (custom)	Importing new structure
<input checked="" type="checkbox"/> Skin (matched)	Importing new structure
<input checked="" type="checkbox"/> penile bulb (custom)	Importing new structure
<input checked="" type="checkbox"/> Neurovascular Bu (custom)	Importing new structure
<input checked="" type="checkbox"/> Prostate (matched)	Importing new structure
<input checked="" type="checkbox"/> PTV (custom)	Importing new structure
<input checked="" type="checkbox"/> Seminal Vesicles (custom)	Importing new structure
<input checked="" type="checkbox"/> Bladder (matched)	Importing new structure
<input checked="" type="checkbox"/> Rectum (matched)	Importing new structure
<input checked="" type="checkbox"/> Urethra (custom)	Importing new structure
<input checked="" type="checkbox"/> Bowel (custom)	Importing new structure
<input checked="" type="checkbox"/> Left Femoral Hea (custom)	Importing new structure
<input checked="" type="checkbox"/> Right Femoral He (custom)	Importing new structure

☐ Archive DICOM on import

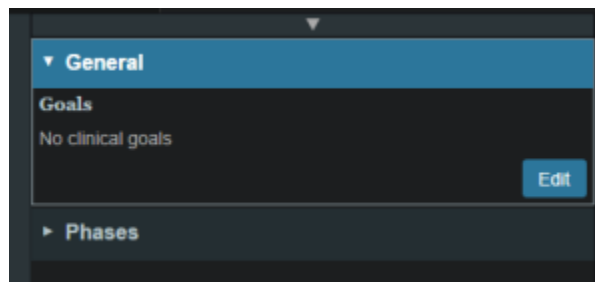
Create PatientCancel

12. You will see a list of the imported structures. Here you may choose whether or not to import each structure by checking or unchecking the box beside each structure name.

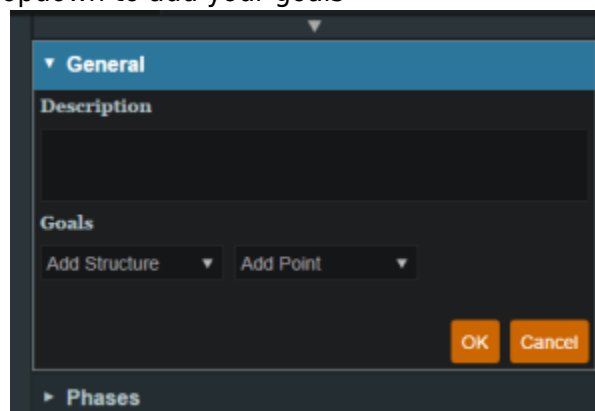
1. For this case, we will import all available structures and there is nothing to change on this page
13. You also have the ability to assign or edit any structures that are shown as *custom*, which indicates the name did not exactly match a directive structure from the treatment site template list you specified during patient creation.
  1. Matched, Assigned, and Custom structures are designated with corresponding tags at the end of the structure name in the structure list.
  2. Assigning a custom structure to a defined directive structure will result in the imported structure inheriting all the predefined structure properties (e.g. name, type, color)
  3. For all custom structures the type is by default set to "Other" unless it contains the letters "TV" (as in PTV or CTV), in which case it is assigned the type of "Target"; this may be changed here if needed
14. Click the "Create Patient" button to create the patient and import the CT Images and Structures into it.
15. The patient is now created and all available data has been imported, so it is time to proceed with entering the prescription information
16. Click on the *Back to Import* button

## Prescription

1. Click the *Patient Search* task
2. Click the entry from the table of patients that corresponds to your newly imported patient
3. Prescription information is stored in the *Directive*
4. Click *Open* under the *Directive* entry in the patient overview tree
5. Click *General* then *Edit*

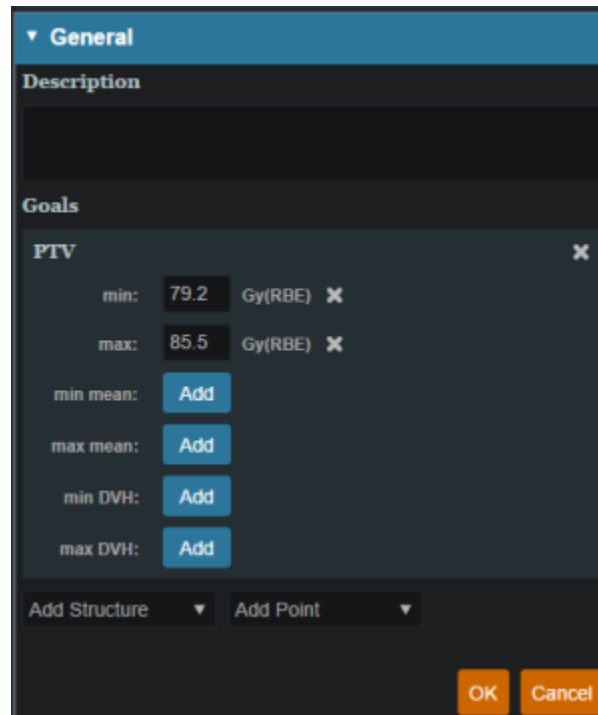


6. Under *General* fill in *Goals*
  1. Choose the structure dropdown to add your goals



2. In this example we will set a minimum dose to the PTV at 79.2Gy and the maximum dose at

85.5Gy as shown below



▼ General

Description

Goals

PTV

min: 79.2 Gy(RBE) ✕

max: 85.5 Gy(RBE) ✕

min mean: Add

max mean: Add

min DVH: Add

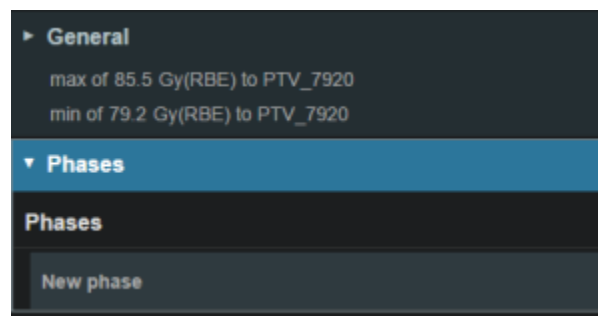
max DVH: Add

Add Structure ▼ Add Point ▼

OK Cancel

7. Next proceed to *Phases* and fill in *Fractions* and *Prescription*

1. Select *New Phase*



► General

max of 85.5 Gy(RBE) to PTV\_7920

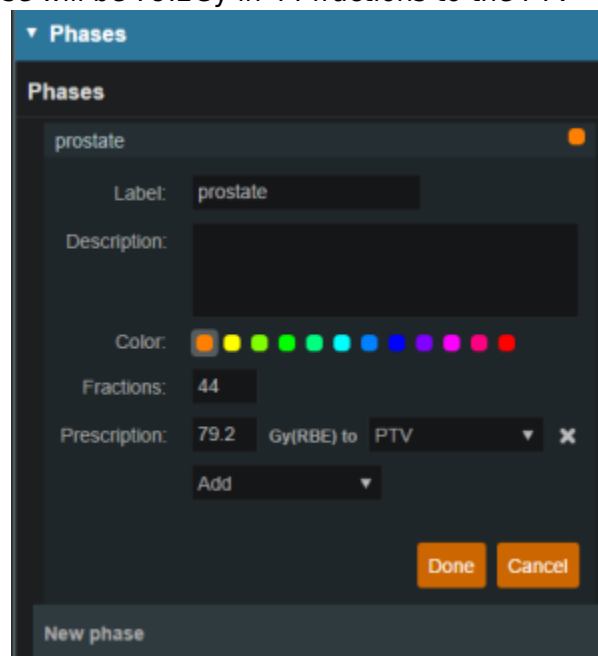
min of 79.2 Gy(RBE) to PTV\_7920

▼ Phases

Phases

New phase

2. For this example the dose will be 79.2Gy in 44 fractions to the PTV



▼ Phases

Phases

prostate

Label: prostate

Description:

Color: ● ● ● ● ● ● ●

Fractions: 44

Prescription: 79.2 Gy(RBE) to PTV ✕

Add ▼

Done Cancel

New phase

3. *Label* and *Description* are free text fields that allow you to “name” each Phase and provide any description or notes necessary to distinguish between each phase in the case of multiple phases. You must enter a “name” in the *Label* in order to proceed. You are free to enter your own values in these fields for this walkthrough
8. Click *Done* to complete the *Directive* editing
9. You can now click on the patient's name, “SBRT PROSTATE”, on the top line to return to the Patient Overview tree

## Plan Creation

1. The required *Intent*, *Course*, and *Snapshot* information was selected during import, so now we can move to creation of a *Request* and *Plan*
2. From the Patient Overview tree, click the *Add Request* button
  1. This creates a default request that fulfills all phases of the prescription
  2. If a plan will only fulfill a portion of the prescription, then the *Request* should be edited to reflect this, otherwise just proceed to adding a plan
  3. In this case, we will be fulfilling the entire prescription with a single plan, so no editing of the Request is needed
3. Now click the *Add Plan* button under the newly created Request entry
  1. Name the plan and click *OK*

Name: SBRT PROSTATE

MRN: 111112

Sex: Other

Date of Birth: 1946-01-01

Archived: No Archive

▼ course\_2017Jan4 Show Info edit

Add Intent

▼ intent\_curative\_2017Jan4 Show Info open

Add Directive

▼ directive\_2017Jan4 Show Info open

▼ snapshot\_2017Jan4 Show Info open

Add Request

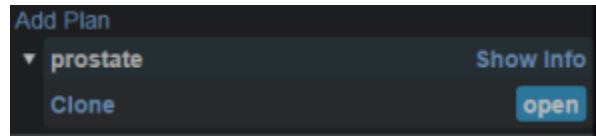
▼ request 1 Show Info open

Name: prostate

Description:

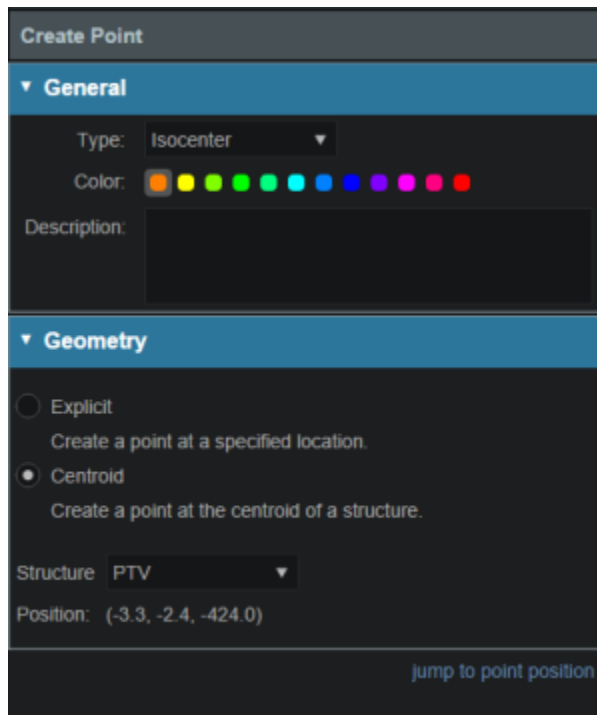
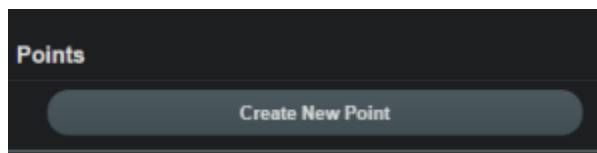
OK Cancel

4. Now open the new plan by clicking the *Open* button under the new plan entry in the Patient Overview tree



## General Plan Parameters

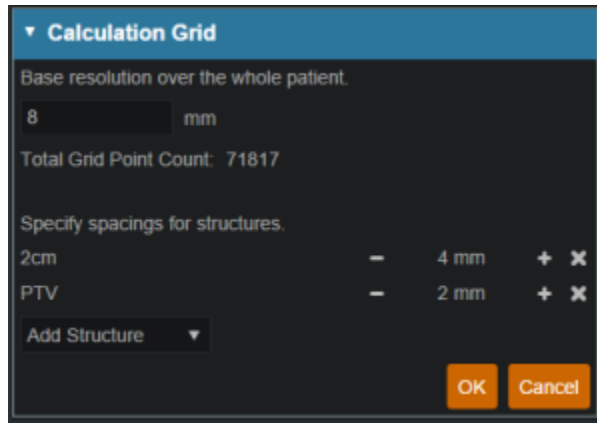
1. Creating our plan will begin in the *Patient Geometry* block, where we will create any points or planning structures that are needed (if not created in contouring software), such as a combination of the left and right femoral heads (see [Structure Geometry](#) for details on how to Create structures)
  1. We will create a point to use for the isocenter (note this step is done for illustration purposes and is not necessary for most plans)
    1. Select a type of isocenter and create the point at the centroid of the PTV



2. Click *DONE* at the bottom of the page
2. Next we will move on to defining the HU to Proton Relative Stopping Power (RSP) conversion
3. Click to open the *RSP Image* block
  1. If any density overrides are needed they are entered here in the *RSP Image* block
  2. If the HU to RSP curve was not chosen upon patient import the user may do so at this point
  3. We do have any density overrides to apply for this case and the proper HU curve has already been selected, so we can move past this block for this example, so click *Cancel* to close this block
4. We are now ready to define the dose calculation grid (see [Defining the Dose Grid](#) for a detailed

explanation)

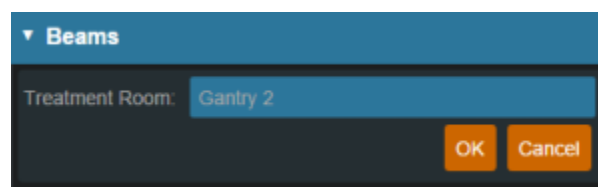
5. Click to open the *Calculation Grid* block
  1. Create a grid by performing the following steps:
    1. Set the base grid resolution to 8 mm
    2. Click *Add Structure* and select the “2cm” structure
      1. Set the size to 4 mm
    3. Add the PTV in the same manner and the sizes to 2 mm



4. The Prostate is included in the PTV volume so there is no need to set a separate grid size
5. Click OK

## Beams

1. We are now ready to begin defining the treatment beams
2. Click to expand the *Beams* block (see [Creating a New Beam](#) for detailed explanation)
  1. Before defining individual beams, we will first set some shared beam properties
  2. Set the *Treatment Room* by choosing “Gantry 2” from the dropdown



3. Click OK
4. Expand the *Spot Placement* sub-block and set the following values (note these are the plan level spot placement parameters, so it will apply to all our beams, unless we specifically override the values within an individual beam)
  1. Lateral margin- 15 mm
  2. Distal margin- 10 mm
  3. Spot spacing- 1 sigma
  4. Layer spacing- 0.8 Distal W80

5. Click OK
6. Now we are ready to make our beams
7. Start by clicking *Create New Beam*

1. Enter the following for each beam
  1. General
    1. Select the PTV as the target and if desired, enter a description for the beam, for other selections leave the defaults (see [Structure Geometry](#))

2. Approach
  1. At this point the user will specify the geometry of the beam by setting the isocenter as well as the gantry and couch angles
  2. Two beams at gantry angles of 90 and 270, respectively, will be used here, which is typical ( $\pm 5$  degrees) for standard prostate plans such as this example
  3. Couch angle of 0 is used here for each beam
    1. You may use the sliders, the  $\pm$  buttons, or simply type in the values to adjust these angles



**Approach**

Isocenter: p1 - Centroid of Prostate

Gantry Angle: 90 deg

Couch Angle: 0 deg

### 3. Snout

1. Depending on the size of the prostate a 10-12cm snout should be adequate, in this example select the 12cm Snout option
2. If the pelvic nodes need to be included, the user may need to increase the snout size to the 18 or 25cm snout

**Snout**

Snout: 12 cm

18 cm

25 cm

30x40 cm

### 4. Aperture (if desired)

1. If there is the need to use an Aperture for sharper penumbra the user can do so at this time (see [Creating an Aperture](#))
2. Note that Astroid has the ability to utilize an Aperture with pencil beam scanning for both SFO and IMPT plans
3. For this plan, no apertures are necessary

### 5. Shifter

1. Simply choose a shifter from the list if one is needed
2. For this plan, no shifters are necessary

**Shifter**

Shifter: none

30 mm

47 mm

80 mm

### 6. Air gap

1. The air gap is specified as the distance from the final beamline device (aperture or shifter) or the snout end if no devices are present
2. The view automatically shifts to the 3D view when entering this sub-block and you can adjust the air gap to ensure patient collisions are avoided
3. The default air gap of 30 mm is sufficient for both beams in this example

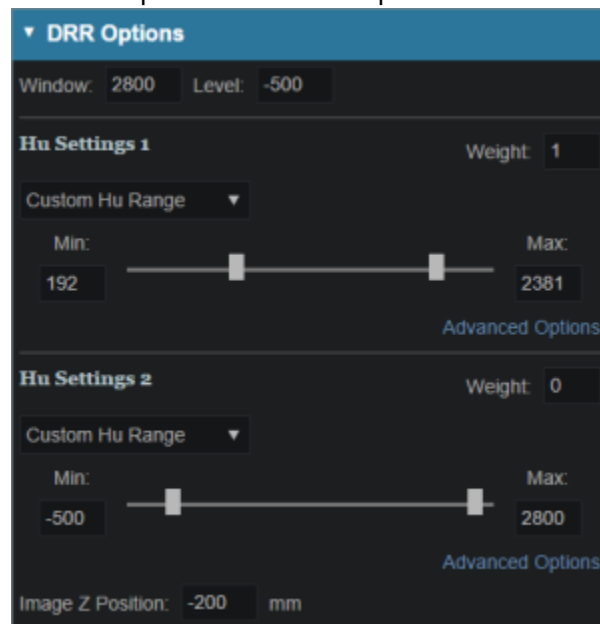
### 7. Spot placement

1. Since we want both beams to use the same spot settings, we have those at the plan level and there is no need to edit the spot placements here for the

beams

## 8. DRR options

1. The Proton DRRs shown in the *Beam* task are generally used to help determine appropriate treatment angles
2. In cases such as these, appropriate angles are readily determined and there is no need to utilize the DRR sub-task
  1. If you wish to explore this feature however, you may choose one of the preset *Density Presets* from the drop down or manually set the *level*, *window*, *min HU* and *max HU* or go into *Advanced Options* for additional control of the DRR's
  2. These proton DRRs are used only for visualization purposes and will have no effect on the plan construction process



2. Click *Done* to complete the first beam
3. After creating the first beam (at gantry angle 90) it is recommended that you *Clone* it and simply change new beam's gantry angle to save time in creating the second beam (the *Clone* button is available under a beam when it is selected for viewing from the list of beams)

**Beams**

Treatment Room: Gantry 2

Spot Placement Parameters: using user values

**PBS Beams**

b1 - G270 C0; 12 cm; no shifter; ●

**Beam Information**

Description:

Isocenter: p1 - Centroid of PTV

Gantry Angle: 270 deg

Couch Angle: 0 deg

Target: PTV

Air Gap: 30 mm

Entrance Z: 210 mm

Snout: 12 cm

Aperture: none

Shifter: none

**PBS Spot Placement**

using plan settings:

Clone Edit Delete

Create New PBS Beam

4. Once you have *Cloned* the beam choose the *Edit* button to make changes to the gantry angle

**PBS Beams**

b1 - G270 C0; 12 cm; no shifter; ●

b2 - G270 C0; 12 cm; no shifter; ●

**Beam Information**

Description:

Isocenter: p1 - Centroid of PTV

Gantry Angle: 270 deg

Couch Angle: 0 deg

Target: PTV

Air Gap: 30 mm

Entrance Z: 210 mm

Snout: 12 cm

Aperture: none

Shifter: none

**PBS Spot Placement**

using plan settings:

Clone Edit Delete

Create New PBS Beam

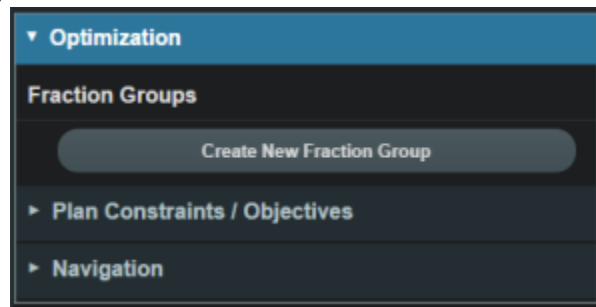
1. Note for more complex plans, there are often additional parameters to change (such as the aperture and the air gap) when creating additional beams via the *Clone* option

## Optimization

1. With the two beams completely defined, we can now move on organizing our treatment fraction groups and specifying the constraints and objectives for the treatment
2. Open the *Optimization* block (see [Astroid Optimization](#) for a detailed explanation of *Optimization*, *Feasibility* and *Constraints*, as well as *Running the Optimizer*)

## Fraction Group

1. Choose *Create a New Fraction Group* under the *Fraction Group* block (See [Fraction Groups](#) for a more detailed explanation)



1. Enter a Fraction Count of 44
2. Add a new Target to the *Target List* by clicking *Add Target*
  1. Set the Target to the PTV
  2. Add the following *Beam Set*
    1. In this case we will use an IMPT approach so add both the G90 and G270 beams in this same beam set
  3. Constraints associated with this target
    1. Click the *Add Structure* drop down and choose the PTV
    2. Add a min dose constraint of 79.2 Gy (1.8 Gy per fx)
    3. Add a max dose constraint of 85.5 Gy
4. At this point your *Fraction Group* should appear as shown below

3. Click *Done* to complete the Target entry and then *Done* again to complete the *Fraction Group*

## Constraints and Objectives

1. This is our only Fraction Group for this plan, so we'll move on to the plan level constraints and objectives
2. Open the *Plan Constraints/Objectives* block
  1. First we'll enter the *Constraints* for the planning (target) volumes and a global maximum constraint
    1. Remember these are the “hard stops/non negotiables” for the plan
    2. These doses will be decided per the users' department protocols
    3. For this case do the following:
      1. Click *Add Structure* and select Skin
        1. Add a max set to 85.5
      2. Click *OK*
      3. Note that we already have PTV dose constraints at the Fraction Group Target level, so no additional one are needed here



After the target constraints have been entered, it's generally good practice to test the plan feasibility. The Feasibility calculation is run by clicking *calculate* in the Feasibility block at the bottom of the *Plan Constraints/Objectives* sub-block. The Feasibility calculation is based solely on the constraints and it should be used to ensure there is a possible solution (plan) that can meet the specified constraints. The Feasibility calculation may be an iterative processes in order to get appropriate constraints established for a particular plan. In other words, the user may need to enter a constraint, check the feasibility, then progressively tighten the constraint and check the feasibility until the plan is no longer feasible. It is recommended practice to start by obtaining a feasible plan utilizing only target constraints then add OAR constraints as desired. Remember, using a narrow range of constraints can improve the optimizer performance and improve the resolution of the Pareto surface navigation.

1. Run the *Feasibility* by clicking the *calculate* link



1. The feasibility should return as passing, if not, please confirm your plan parameters match those shown throughout this walkthrough and try again
2. Now with the target constraints set, we'll move on to our OAR's
  1. Again these will be set per department protocol
  2. For this example create the following OAR constraints:
    1. Rectum max mean 50 Gy
    2. Bladder max mean 60 Gy
3. Run *Feasibility* again
  1. The feasibility should return as passing again, if not, please confirm your plan parameters match those shown throughout this walkthrough and try again
4. With narrow constraints set and feasibility established, we will now turn our attention to *Objectives*
  1. *Objectives* are the goals that you would like to achieve and these will be the driving forces for the optimizer
  2. Click *Edit* under the *Objectives* section
  3. For this example we will use the following objectives:
    1. Rectum min\_mean (Minimize the mean dose to the Rectum)
    2. Testes min\_max (Minimize the maximum dose to the Testes)
    3. Left & Right Femur min\_overdose 45 Gy (Minimize the amount of the structure the receives dose above 45 Gy)
    4. PTV min\_max (Minimize the maximum dose within the PTV)
    5. PTV max\_min (Maximize the minimum dose within the PTV)

**Objectives**

- Rectum**
  - ☐ min\_max ☒ min\_mean ☐ min\_overdose
- Testes**
  - ☒ min\_max ☐ max\_min
  - ☐ min\_mean ☐ max\_mean
  - ☐ min\_overdose ☐ min\_underdose
- Left Femoral Hea**
  - ☐ min\_max ☐ max\_min
  - ☐ min\_mean ☐ max\_mean
  - ☒ min\_overdose ☐ min\_underdose
  - 45 Gy(RBE)
- Right Femoral He**
  - ☐ min\_max ☐ max\_min
  - ☐ min\_mean ☐ max\_mean
  - ☒ min\_overdose ☐ min\_underdose
  - 45 Gy(RBE)
- PTV**
  - ☒ min\_max ☒ max\_min
  - ☐ min\_mean ☐ max\_mean
  - ☐ min\_overdose ☐ min\_underdose

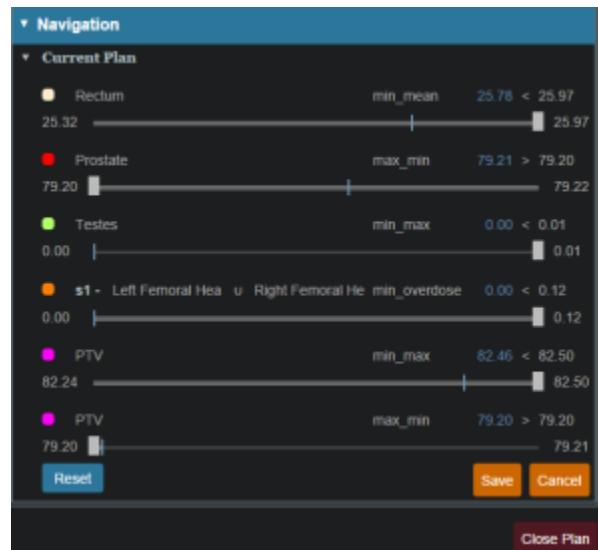
Add Structure ▼

OK Cancel

5. Click **OK** to complete the *Objectives*
6. We are now ready to run the MCO, by opening the *Navigation* sub-block and clicking the *calculate* link
  1. This may take up to 20 minutes for this plan depending upon your available cloud services resources
  2. Keep in mind that even though this process may take some time the planner will be able to make adjustments easily in the *Navigation* without running a new plan

## Navigation

1. Once the MCO has been completed, you can adjust the relative importance of the objectives using the slider bars in the *Navigation* block (see [Astroid Optimization Navigating the Solutions](#) for a detailed explanation of using the slider bars)
  1. Adjust the slider bars in the *Navigation* block to find what you feel is the optimal plan for this case
    1. Note: All of these adjustments are made without having to run a new plan
  2. If you find a plan that you like, but wish to continue exploring further, you can click the *Save* button to save the current slider state and then continue exploring
  3. You may return to the last saved state at any time by clicking the *Reset* button
  4. The *Cancel* button will close the navigation block, reverting back to the last *Saved* state



2. The plan is ready for physician review at this point
  1. Physicians often find it useful to Navigate the solution and explore trade-offs in real time, which can often alter the traditional physician approval process into an interactive, real-time trade-off analysis and on the spot plan approval
3. After the physician has decided on a treatment plan the user may then proceed to export the chosen plan to the EMR
  1. Note that these standard (non-plan specific) features are covered in other guides and not repeated herein

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[https://apps.dotdecimal.com/doku.php?id=planning:userguide:walkthroughs:prostate\\_plan&rev=1483807814](https://apps.dotdecimal.com/doku.php?id=planning:userguide:walkthroughs:prostate_plan&rev=1483807814)

Last update: **2021/07/29 18:25**