```
import pandas as pd
import numpy as np
import regex as re
from matplotlib import pyplot as plt
from docplex.mp.model import Model
import importlib

import load_data as ld
import analytics as an
import models as md

# Reloads all local modules
def reload_all():
    for _ in [ld, an, md]:
        importlib.reload(_)
```

Load in Map Data

Small Data

```
In [2]: specs_small = ld.get_specs('smallexample')
    s_c, s_t, s_b = ld.load_data('smallexample', specs_small)

Number of beams: 5
    Vertical pixel resolution: 8
    Horizontal pixel resolution: 8
    Maximum Dose Allowed Over Critical Area: 2
    Minimum Dose Required Over Tumor Area: 10

Map Loaded: Critical
    Map Loaded: Tumor
    Map Loaded: Beams
```

Actual Data

```
In [141... specs_actual = ld.get_specs('actualexample')
    a_c, a_t, a_b = ld.load_data('actualexample', specs_actual)

Number of beams: 126
    Vertical pixel resolution: 60
    Horizontal pixel resolution: 80
    Maximum Dose Allowed Over Critical Area: 2
    Minimum Dose Required Over Tumor Area: 10

Map Loaded: Critical
    Map Loaded: Tumor
    Map Loaded: Beams
```

Model 1

```
\min \sum_{i} \sum_{i} \sum_{k} x(i) b(ijk) (t(jk))
        st
        \sum_{i} x(i) b(ijk) <= 2 for every critical cell (jk)
        \sum_{i} x(i) b(ijk) >= 10 for every tumor cell (jk)
        x(i) >= 0 for every beam (i)
          if False:
In [4]:
              sol = md.build_model_1(specs_actual, a_c, a_t, a_b)
              an.plot_beams(sol, a_b, a_c, a_t, print_vars = True)
          else:
              sol = md.build_model_1(specs_small, s_c, s_t, s_b)
              an.plot beams(sol, s b, s c, s t, print vars = True)
         Intensity variables added.
         CRITICAL ERROR FOR 0 1
         CRITICAL ERROR FOR 0 2
         CRITICAL ERROR FOR 1 2
         CRITICAL ERROR FOR 2 0
         Intensity constraints added.
         Object Function Constructed.
         -5.200x1-4.200x2-3.600x3+1.200x4-3.100x5
         Model Exported.
         Model: m1
          - number of variables: 5
            - binary=0, integer=0, continuous=5
          - number of constraints: 25
            - linear=25
          - parameters: defaults
          - objective: minimize
          - problem type is: LP
         ERROR: NO SOLUTION
         AttributeError
                                                     Traceback (most recent call last)
         <ipython-input-4-6a4c505c8710> in <module>
               4 else:
                      sol = md.build_model_1(specs_small, s_c, s_t, s_b)
                      an.plot_beams(sol, s_b, s_c, s_t, print_vars = True)
         ---> 6
         ~\Desktop\Homework\Radiation-Therapy-Optimization\analytics.py in plot_beams(sol, b, c, t, cmap_choice, print_vars)
                      """Plot the path of the beams in python"""
              28
              29
         ---> 30
                     m = calc_m(sol, b, print_vars)
              31
```

```
~\Desktop\Homework\Radiation-Therapy-Optimization\analytics.py in calc_m(sol, b, print_vars)
             sol 1 = []
 ---> 9
             for var, value in sol.iter_var_values():
      10
                  if str(var)[0] == 'x':
      11
                      if print vars == True:
 AttributeError: 'NoneType' object has no attribute 'iter_var_values'
Model 2
\min \sum_{i} \sum_{j} \sum_{k} x(i) b(ijk) (c(jk))
st
\sum_{i} x(i) b(ijk) - s(jk) <= 2 for every critical cell (jk)
\sum_{i} x(i) b(ijk) + s(jk) > = 10 for every tumor cell (jk)
x(i) >= 0 for every beam (i)
0 \le s(jk) \le 10 for every slack/surplus variable
 if True:
      sol = md.build_model_2(specs_actual, a_c, a_t, a_b)
      an.plot_beams(sol, a_b, a_c, a_t, print_vars = True)
      an.report_effectiveness(sol, a_b, a_c, a_t, plot=False)
  else:
      sol = md.build_model_2(specs_small, s_c, s_t, s_b)
      an.plot_beams(sol, s_b, s_c, s_t, print_vars = True)
      an.report effectiveness(sol, s b, s c, s t, plot=False)
 Intensity variables added.
 Intensity constraints added.
 Object Function Constructed.
 Model Exported.
 Model: m2
  - number of variables: 9726
    - binary=0, integer=0, continuous=9726
  - number of constraints: 20026
    - linear=20026
  - parameters: defaults
  - objective: minimize
  - problem type is: LP
 Version identifier: 20.1.0.0 | 2020-11-10 | 9bedb6d68
 CPXPARAM Read DataCheck
 Parallel mode: deterministic, using up to 4 threads for concurrent optimization:
```

32

In [5]:

from matplotlib.colors import colorConverter

- * Starting dual Simplex on 1 thread...
- * Starting Barrier on 3 threads...

Tried aggregator 1 time.

LP Presolve eliminated 20026 rows and 9726 columns.

All rows and columns eliminated.

Presolve time = 0.03 sec. (5.55 ticks)

Dual simplex solved model.

Model Solved.

x3 15.625

x5 15.625

x9 14.492753623188408

x11 1.93661971830986

x15 13.513513513513512

x17 13.513513513513512

x21 13.513513513513512

x27 12.5

x29 12.345679012345679

x33 12.82051282051282

x35 13.157894736842106

x66 0.3523608174770967

x77 15.625

x87 11.904761904761905

x93 12.82051282051282

x98 12.987012987012987

x104 11.627906976744187

MODEL REPORT

No units of radiation were delivered to any critical cells.

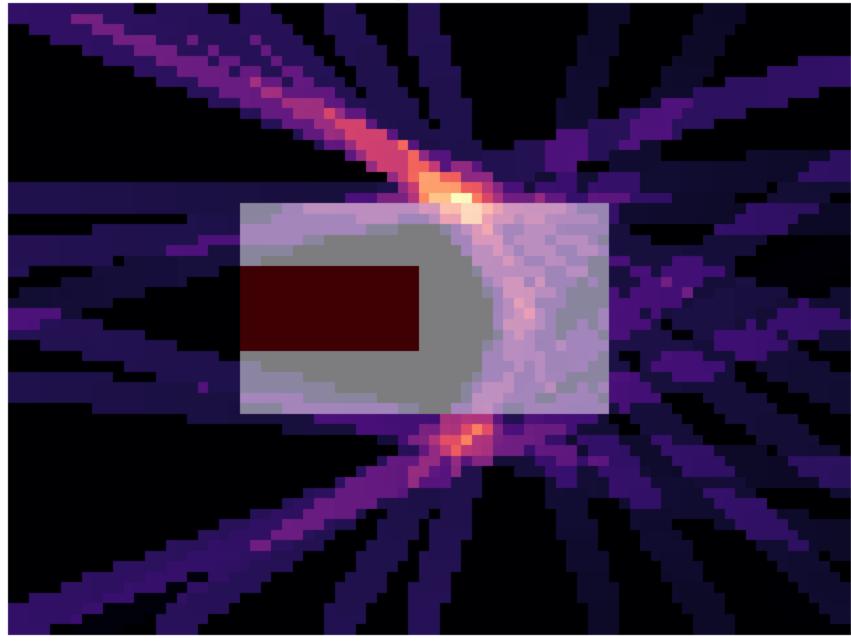
8625.8 units of radiation were delivered to tumor cells.

22.8 units were delivered to each cell, on average.

368 cells were found to have acceptable levels of radiation, out of 564.

65.25% of cells were found to have acceptable levels of radiation.





Model 2.1

 $\min \sum_i \sum_j \sum_k \mathsf{x(i)} \; \mathsf{b(ijk)} \; (\mathsf{c(jk)}) + \mathsf{s1(jk)} \; \mathsf{t(jk)} + \mathsf{s2(jk)} \; \mathsf{c(jk)}$

```
\sum_{i} x(i) b(ijk) + s2(jk) > = 10 for every tumor cell (jk)
0 <= x(i) <= 100 \text{ for every beam (i)}
0 \le s(jk) \le 10 for every slack/surplus variable (jk)
 if True:
     sol = md.build_model_2_1(specs_actual, a_c, a_t, a_b)
     an.plot_beams(sol, a_b, a_c, a_t, print_vars = True)
     an.report effectiveness(sol, a b, a c, a t, plot=False)
 else:
     sol = md.build_model_2_1(specs_small, s_c, s_t, s_b)
     an.plot_beams(sol, s_b, s_c, s_t, print_vars = True)
     an.report_effectiveness(sol, s_b, s_c, s_t, plot=False)
Intensity variables added.
Intensity constraints added.
Object Function Constructed.
Model Exported.
Model: m1
 - number of variables: 9726
    - binary=0, integer=0, continuous=9726
 - number of constraints: 20026
    - linear=20026
 - parameters: defaults
 - objective: minimize
  - problem type is: LP
Version identifier: 20.1.0.0 | 2020-11-10 | 9bedb6d68
CPXPARAM_Read_DataCheck
Parallel mode: deterministic, using up to 4 threads for concurrent optimization:
 * Starting dual Simplex on 1 thread...
 * Starting Barrier on 3 threads...
Tried aggregator 1 time.
LP Presolve eliminated 19704 rows and 9331 columns.
Reduced LP has 322 rows, 395 columns, and 4393 nonzeros.
Presolve time = 0.06 sec. (7.12 ticks)
Iteration log . . .
Iteration:
               1 Dual objective
                                                     10.000000
Iteration: 62 Dual objective
                                                    550.355840
Iteration: 124 Dual objective
                                                    881.711126
Iteration:
            186 Dual objective
                                                   1078.154005
Iteration:
             248 Dual objective
                                                   1128.821369
Dual simplex solved model.
Model Solved.
x3 15.625
x5 15.625
```

 $\sum_{i} x(i) b(ijk) - s1(jk) <= 2$ for every critical cell (jk)

x9 14.492753623188408

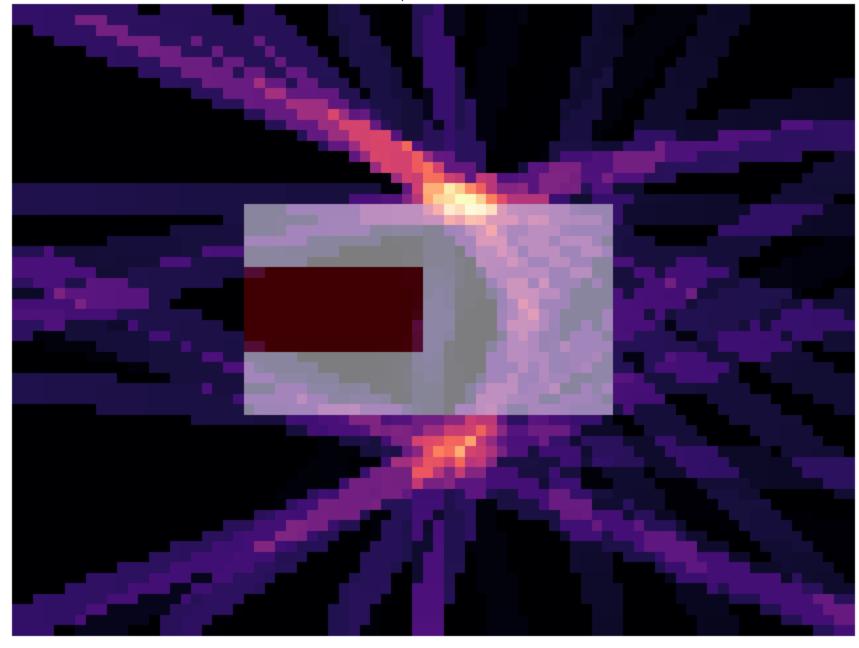
In [6]:

x15 13.513513513513512 x17 13.513513513513512 x21 13.513513513513512 x27 9.720877062887908 x29 12.345679012345679 x33 12.82051282051282 x34 7.97180298237686 x35 13.157894736842106 x37 2.6655620826429707 x40 2.388190992619371 x43 0.5918793763348669 x49 2.2433267290022836 x66 12.778087586210042 x83 14.994361348528017 x87 11.904761904761905 x92 12.195121951219512 x93 0.4748338081671406 x98 2.143557582147733 x99 12.345679012345679 x104 11.627906976744187 MODEL REPORT

x11 4.637457502263864

- 93.8 units of radiation were delivered to critical cells.
 5.5 units were delivered to each cell, on average.
 8 cells were found to have acceptable levels of radiation, out of 564.
 5.88% of cells were found to have acceptable levels of radiation.
- 9612.7 units of radiation were delivered to tumor cells.
 18.9 units were delivered to each cell, on average.
 418 cells were found to have acceptable levels of radiation, out of 564.
 74.11% of cells were found to have acceptable levels of radiation.

Map of Radiation



Model 3

 $\min \sum_i \sum_j \sum_k \mathsf{x(i)} \; \mathsf{b(ijk)} \; (\mathsf{c(jk)} + \mathsf{p(critical)} \; \mathsf{c_neighbor(jk))} \; + \; \mathsf{s1(jk)} \; \mathsf{t(jk)} \; + \; \mathsf{s2(jk)} \; \mathsf{c(jk)}$

```
\sum_{i} x(i) b(ijk) + s2(jk) > = 10 for every tumor cell (jk)
0 <= x(i) <= 100 \text{ for every beam (i)}
0 \le s(jk) \le 2 for every slack variable (jk)
0 \le s(jk) \le 10 for every surplus variable (jk)
 reload_all()
 if True:
      sol = md.build_model_3(specs_actual, a_c, a_t, a_b)
      an.plot_beams(sol, a_b, a_c, a_t, print_vars = True, cmap_choice = 'magma')
      an.report_effectiveness(sol, a_b, a_c, a_t, plot=False)
 else:
      sol = build_model_3(specs_small, s_c, s_t, s_b)
     an.plot_beams(sol, s_b, s_c, s_t, print_vars = True)
     an.report_effectiveness(sol, s_b, s_c, s_t, plot=False)
Intensity variables added.
Intensity constraints added.
Object Function Constructed.
Model Exported.
Model: m1
 - number of variables: 9726
    - binary=0, integer=0, continuous=9726
 - number of constraints: 20026
    - linear=20026
  - parameters: defaults
 - objective: minimize
  - problem type is: LP
Version identifier: 20.1.0.0 | 2020-11-10 | 9bedb6d68
CPXPARAM_Read_DataCheck
Parallel mode: deterministic, using up to 4 threads for concurrent optimization:
 * Starting dual Simplex on 1 thread...
 * Starting Barrier on 3 threads...
Tried aggregator 1 time.
LP Presolve eliminated 19695 rows and 9321 columns.
Reduced LP has 331 rows, 405 columns, and 4494 nonzeros.
Presolve time = 0.05 sec. (7.16 ticks)
Iteration log . . .
Iteration:
                1 Dual objective
                                                      10.000000
Iteration:
               71 Dual objective
                                                     639.638573
Iteration:
            137
                    Dual objective
                                                    1100.107912
Dual simplex solved model.
Model Solved.
x3 15.625
```

 $\sum_{i} x(i) b(ijk) - s1(jk) <= 2$ for every critical cell (jk)

In [178...

x5 15.625 x9 14.492753623188408 x10 0.39617920816496266 x11 13.23969367749179 x15 12.802493819087317 x17 13.513513513513512 x21 12.78449332049425 x23 0.6660184479435245 x27 1.0460206082224968 x28 0.2582340885105248 x29 12.345679012345679 x33 12.08434727050849 x34 11.031162575675468 x35 13.157894736842106 x37 1.4035980705089934 x40 3.254551710038818 x43 1.140669068501193 x49 11.288743764464163 x66 15.248547335600907 x76 0.3450655624568668 x83 14.794402751719826 x87 11.904761904761905 x92 11.904761904761905 x93 0.625390869293307 x98 0.2834467120181401 x99 12.195121951219512 x104 11.627906976744187

MODEL REPORT

150.0 units of radiation were delivered to critical cells.
6.3 units were delivered to each cell, on average.
119 cells were found to have acceptable levels of radiation, out of 136.

87.5% of cells were found to have acceptable levels of radiation.

9953.2 units of radiation were delivered to tumor cells.

18.9 units were delivered to each cell, on average.

485 cells were found to have acceptable levels of radiation, out of 564.

85.99% of cells were found to have acceptable levels of radiation.

Map of Radiation

Model 4: Surgical Removal

Model

 $\min \sum_i \sum_j \sum_k \mathsf{x(i)} \ \mathsf{b(ijk)} \ (\mathsf{c(jk)} + \alpha \ \mathsf{c_n(jk)}) \ + \ \mathsf{s1(jk)} \ (\mathsf{t(jk)} - \beta \ \mathsf{tr(jk)}) \ + \ \mathsf{s2(jk)} \ (\mathsf{c(jk)} + \alpha \ \mathsf{c_n(jk)})$

```
\sum_{i} x(i) b(ijk) - s1(jk) <= 2 \ \forall (j,k) \ if \ c(j,k) = 1 \in (J, K)
         \sum_{i} x(i) b(ijk) tr(jk) + s2(jk) > = 10 \forall (j,k) if t(j,k) = 1 \in (J, K)
         0 <= x(i) <= 100 \ \forall \ i \in (I)
         0 \le s(jk) \le 2 \forall (j,k) \text{ if } c(j,k) = 1 \in (J, K)
         0 \le s(jk) \le 10 \ \forall (j,k) \ \text{if} \ t(j,k) = 1 \in (J, K)
         New Variables
         \alpha = P(Critical Region Not Identified Properly)
         \beta = P(Tumor Regrew After Removal)
         tr(jk) = Tumor Removed from Cell (jk)
           reload_all()
In [181...
           if True:
                sol, t removed = md.build_model_4(specs_actual, a_c, a_t, a_b)
                an.plot_beams(sol, a_b, a_c, t_removed, print_vars = True, cmap_choice = 'binary')
                an.report effectiveness(sol, a b, a c, t removed, plot=False)
            else:
                sol = md.build_model_4(specs_small, s_c, s_t, s_b)
                an.plot beams(sol, s b, s c, t removed, print vars = True)
                an.report_effectiveness(sol, s_b, s_c, t_removed, plot=False)
          Intensity variables added.
          Intensity constraints added.
          Object Function Constructed.
          Model Exported.
          Model: m1
           - number of variables: 9726
              - binary=0, integer=0, continuous=9726
            - number of constraints: 15127
              - linear=15127
            - parameters: defaults
            - objective: minimize
            - problem type is: LP
          Version identifier: 20.1.0.0 | 2020-11-10 | 9bedb6d68
          CPXPARAM Read DataCheck
          Parallel mode: deterministic, using up to 4 threads for concurrent optimization:
            * Starting dual Simplex on 1 thread...
            * Starting Barrier on 3 threads...
          Tried aggregator 1 time.
          LP Presolve eliminated 14809 rows and 9334 columns.
```

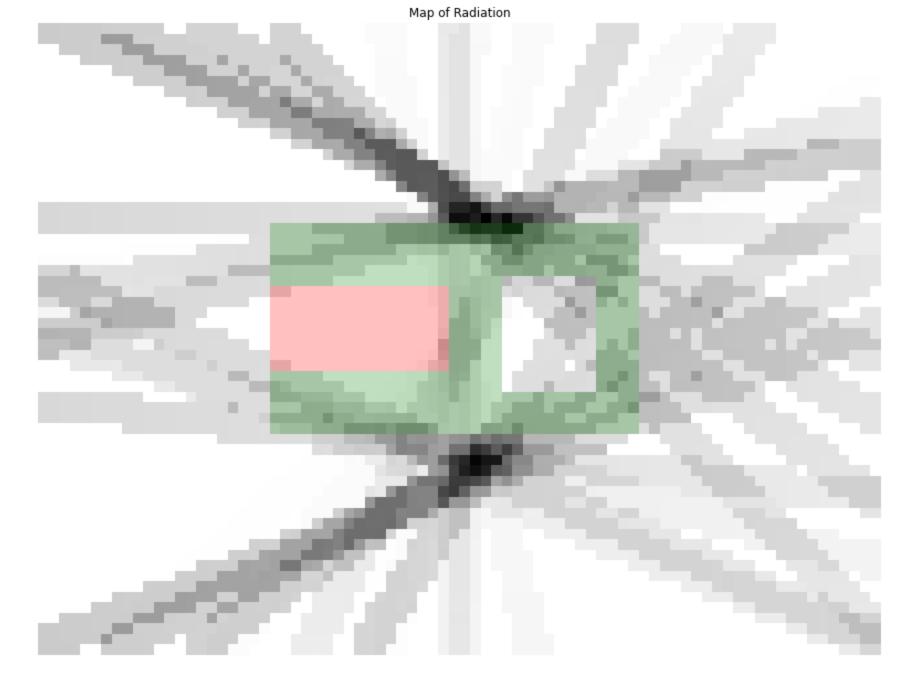
Reduced LP has 318 rows, 392 columns, and 4328 nonzeros. Presolve time = 0.03 sec. (6.01 ticks) Iteration log . . . Iteration: 1 Dual objective 10.000000 Iteration: 85 Dual objective 770.005784 Iteration: Dual objective 147 1064.564531 Dual simplex solved model. Model Solved.

x3 15.625 x5 15.625 x9 14.492753623188408 x10 0.4358136782146525 x11 14.285714285714286 x15 13.20339947790928 x17 13.513513513513512 x21 13.195548489666136 x23 0.29048656499636927 x28 0.30239095321973153 x29 12.345679012345679 x31 6.58537353877178 x34 4.0032139143586365 x37 1.5237266885792256 x40 2.8169014084507045 x43 10.260094802346735 x49 2.1301916948003385 x66 6.26584079381505 x76 0.15729569498595464 x83 14.794402751719826 x87 11.904761904761905 x92 11.904761904761905 x93 0.625390869293307 x98 7.0561946238335915 x99 12.195121951219512 x104 11.627906976744187

MODEL REPORT

151.9 units of radiation were delivered to critical cells.
6.3 units were delivered to each cell, on average.
121 cells were found to have acceptable levels of radiation, out of 136.
88.97% of cells were found to have acceptable levels of radiation.

7460.9 units of radiation were delivered to tumor cells.
17.2 units were delivered to each cell, on average.
389 cells were found to have acceptable levels of radiation, out of 465.
83.66% of cells were found to have acceptable levels of radiation.



Model 5: Magnetic Fields

Model

 $\min \sum_{m} \sum_{i} \sum_{j} \sum_{k} \mathsf{x}(\mathsf{mi}) \; \mathsf{b}(\mathsf{mijk}) \; (\mathsf{c}(\mathsf{jk}) + \alpha \; \mathsf{c}_{-}\mathsf{n}(\mathsf{jk})) \; + \; \mathsf{s1}(\mathsf{jk}) \; (\mathsf{t}(\mathsf{jk}) - \beta \; \mathsf{tr}(\mathsf{jk})) \; + \; \mathsf{s2}(\mathsf{jk}) \; (\mathsf{c}(\mathsf{jk}) + \alpha \; \mathsf{c}_{-}\mathsf{n}(\mathsf{jk}))$

$$\sum_{m}\sum_{i} x(mi) b(mijk) - s1(jk) <= 2 \ \forall \ (j,k) \ if \ c(j,k) = 1 \in (J, K)$$

$$\sum_{m}\sum_{i}$$
 x(mi) b(mijk) + s2(jk) >= 10 \forall (j,k) if t(j,k)-tr(j,k) = 1 \in (J, K)

$$0 \le x(mi) \le 100 \ \forall \ (m,i) \in (M, I)$$

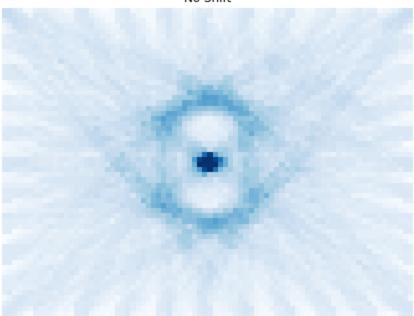
$$0 \le s(jk) \le 2 \ \forall \ (j,k) \ \text{if} \ c(j,k) = 1 \in (J,K)$$

$$0 \le s(jk) \le 10 \ \forall \ (j,k) \ \text{if} \ t(j,k) = 1 \in (J,\ K).$$

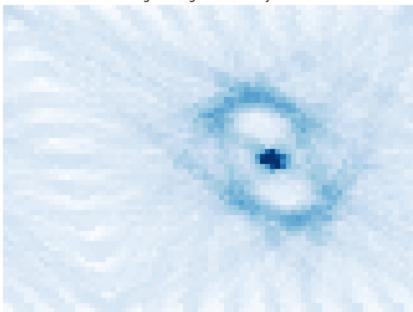
New Variables

 $m = Choice of magnetic field \in M = \{'None', 'Left', 'Right'\}$

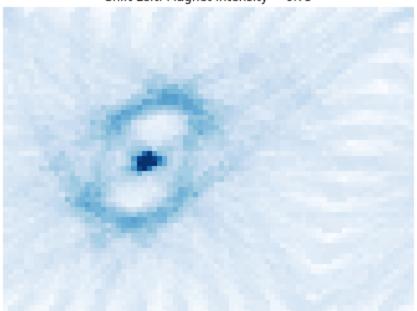
No Shift



Shift Right: Magnet Intensity = 0.75



Shift Left: Magnet Intensity = 0.75



```
In [180... reload_all()
    if True:
        sol, t_removed = md.build_model_5(specs_actual, a_c, a_t, a_b)
        an.plot_beams(sol, a_b, a_c, t_removed, print_vars = True, magnetic=True)
        an.report_effectiveness(sol, a_b, a_c, t_removed, plot=False, magnetic=True)
    else:
        sol = md.build_model_5(specs_small, s_c, s_t, s_b)
```

```
an.plot_beams(sol, s_b, s_c, t_removed, print_vars = True)
     an.report effectiveness(sol, s b, s c, t removed, plot=False)
Intensity variables added.
Intensity constraints added.
Object Function Constructed.
Model Exported.
Model: m1
 - number of variables: 9978
   - binary=0, integer=0, continuous=9978
 - number of constraints: 20179
   - linear=20179
 - parameters: defaults
 - objective: minimize
 - problem type is: LP
Version identifier: 20.1.0.0 | 2020-11-10 | 9bedb6d68
CPXPARAM Read DataCheck
Parallel mode: deterministic, using up to 4 threads for concurrent optimization:
 * Starting dual Simplex on 1 thread...
* Starting Barrier on 3 threads...
Tried aggregator 1 time.
LP Presolve eliminated 19965 rows and 9587 columns.
Reduced LP has 214 rows, 391 columns, and 7934 nonzeros.
Presolve time = 0.03 sec. (8.67 ticks)
Iteration log . . .
Iteration:
               1 Dual objective
                                                   10.000000
Iteration:
             77 Dual objective
                                                  480.231748
Dual simplex solved model.
Model Solved.
x3 15.625
x5 15.625
x9 14.492753623188408
x11 14.285714285714286
x15 12.990040657987992
x17 5.693896264972257
x21 12.5
x29 8.746261238224246
x66 10.105269170418492
x77 0.2648305084745779
x83 15.367685633575466
x87 6.6054180980812465
x92 0.08159934720522166
x98 4.156032624626077
x131 9.421628666646846
x137 2.4052849885479368e-05
x138 0.2421307506053251
x182 0.003228410008070698
x191 8.147936445559852
x216 0.1417434443656962
x222 0.001707752341755353
x270 13.157894736842106
```

x272 0.33738191632928505 x276 5.023635936538845 x325 7.228915662650603 x331 4.819277108433735 x335 16.949152542372882 x361 11.764705882352942 x372 10.7171999766859 x378 7.017543859649123

MODEL REPORT

27.9 units of radiation were delivered to critical cells.4.0 units were delivered to each cell, on average.129 cells were found to have acceptable levels of radiation, out of 136.94.85% of cells were found to have acceptable levels of radiation.

9087.4 units of radiation were delivered to tumor cells.
20.2 units were delivered to each cell, on average.
432 cells were found to have acceptable levels of radiation, out of 465.
92.9% of cells were found to have acceptable levels of radiation.

Map of Radiation

