

# Shadie - A Domain Specific Language for Radiation Oncology

Hanspeter Pfister  
[pfister@seas.harvard.edu](mailto:pfister@seas.harvard.edu)



# The Team



- MGH / HMS Radiation Oncology Group

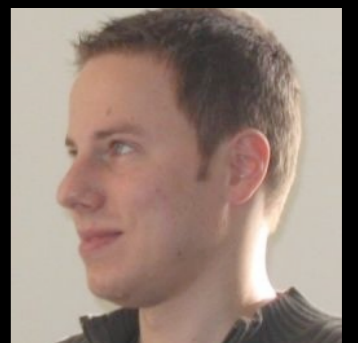
- George Chen



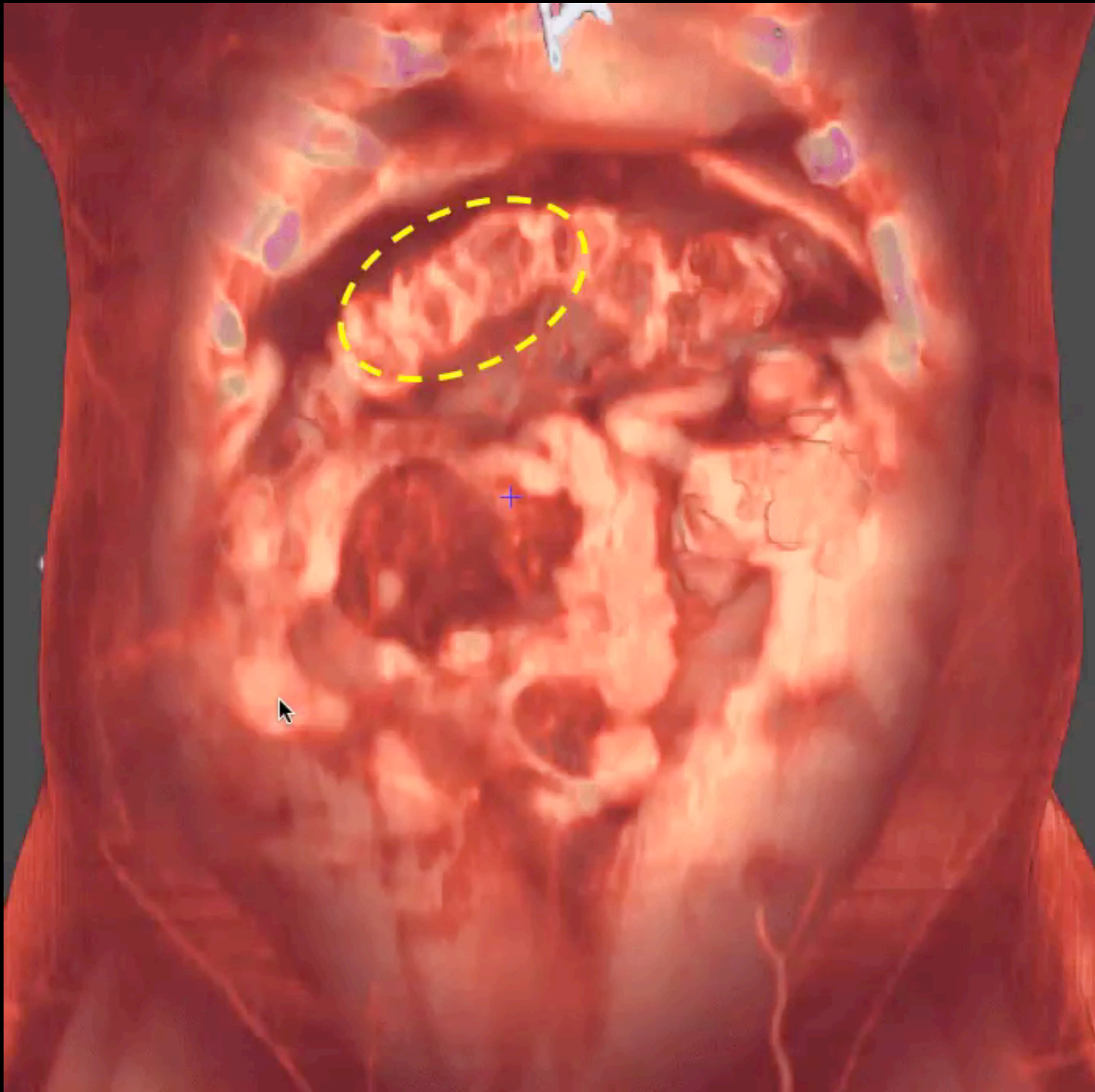
- John Wolfgang

- School of Engineering and Applied Sciences

- Milos Hasan (UC Berkeley)

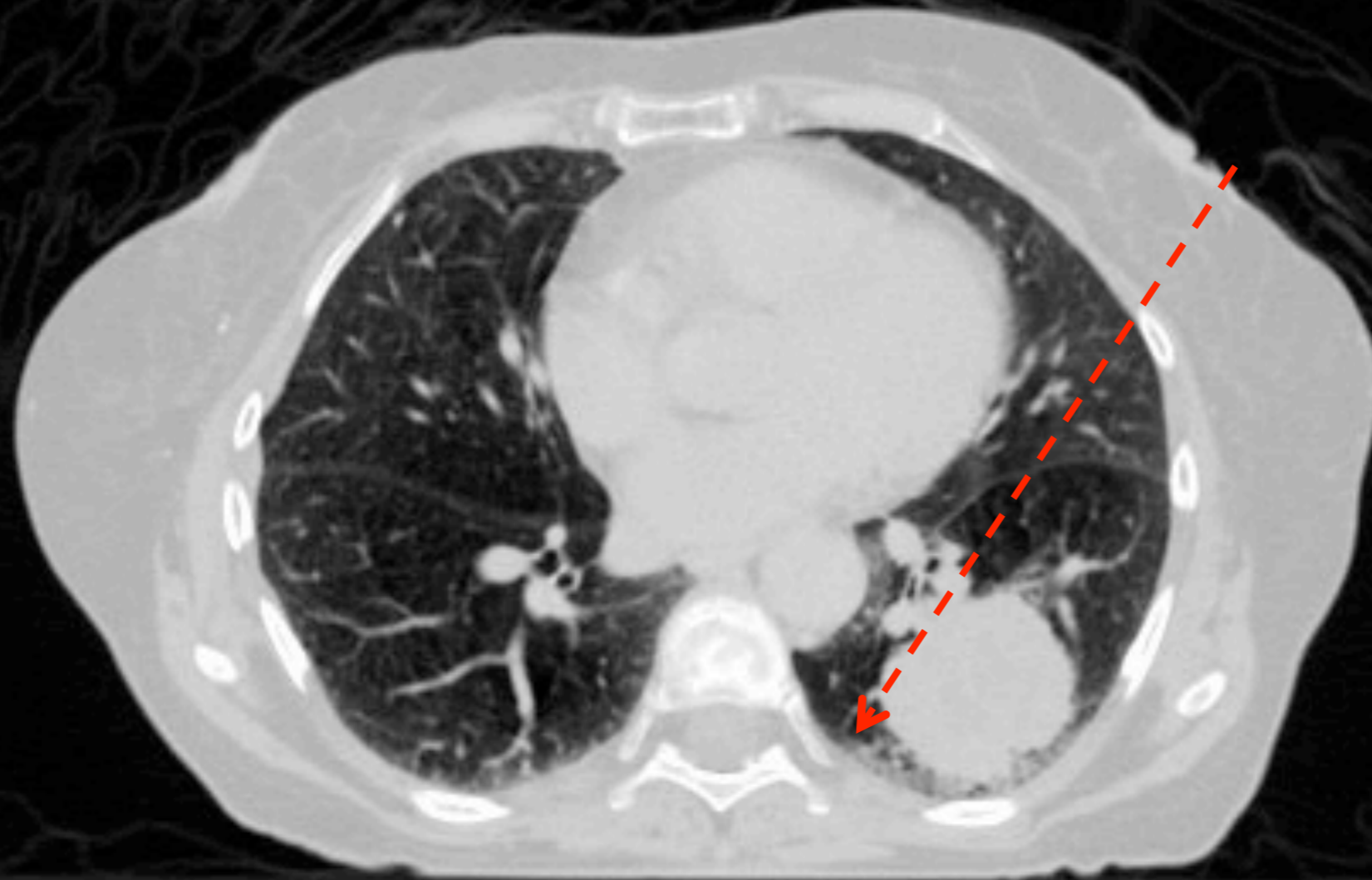


- Hanspeter Pfister



FOVIA

# Quantitative 4D Image Understanding



GTV: 4cm dia,  
5cm long

S/I Motion:  
1.25cm

- Shader to display range from BEV source to distal ITV
- rpl.mov
- Green less range, brown greater

# Applications

- Multiple data sets (serial CT)
- Visualize multimodality data sets (PET, functional imaging)
- Calculate dose on the fly
- Calculate distances, volumes as a fcn of BEV
- VISUALIZE UNCERTAINTY!!!

# What Scientists Want

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- Focus on the code that solves their problem
- Being able to tweak and change code
- Getting good speedup from GPUs

# The Challenge

- Scientists are not CUDA hackers ...

```
// Determinte the size of each block (of threads)
dim3 dimBlock(BLOCKDIM_X,
             BLOCKDIM_Y,
             1);

// Determinte the size of the grid (of blocks)
dim3 dimGrid( (int) ceil((float)width/(float)BLOCKDIM_X),
             (int) ceil((float)height/(float)BLOCKDIM_Y),
             1);

// Allocate some device memory for the output
uchar4 *d_out;
cudaMalloc((void **)&d_out, sizeof(uchar4)*width*height);

// Setup texture and 2D array
texRef4.addressMode[0] = cudaAddressModeClamp;
texRef4.addressMode[1] = cudaAddressModeClamp;
texRef4.filterMode = cudaFilterModePoint;
texRef4.normalized = false;
cudaArray *d_tex;
cudaMallocArray( (cudaArray**)&d_tex, &uchar4Desc, width, height );
cudaMemcpyToArray(d_tex, 0, 0, _in, sizeof(uchar4)*width*height,
cudaMemcpyHostToDevice);
cudaBindTextureToArray(texRef4, d_tex, uchar4Desc);

// Launch the CUDA kernel
kernel_boxcar_texture<<< dimGrid, dimBlock >>>(d_out, width, height,
halfkernelsize);
cutilCheckMsg("kernel_boxcar_texture execution failed.\n");
```

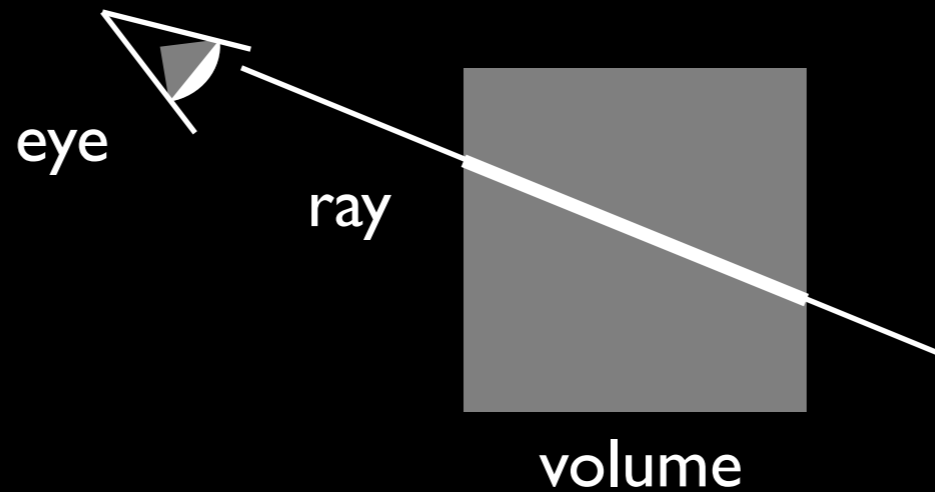
# The current reality of volume visualization

file format  
parsing

compilation  
and linking

user  
interface

resource  
allocation



memory  
corruption  
debugging

CPU-GPU  
communication

data  
interpolation

progressive  
refinement

# Enter: DSLs

- Domain Specific Languages (DSLs) promise the solution
- Two approaches:
  - Provide environment to develop DSLs (Stanford)
  - Develop small DSLs for specific target domains (Harvard)

# DSLs for GPUs

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- Restrict ourselves to something modest but feasible
- A high-level language for GPU computing?
  - We don't know the solution yet.
- A high-level language for GPU visualization of volumetric data for radiation therapy purposes?
  - Yes!



A 3D medical visualization of a human head and neck. The skull is rendered in a semi-transparent blue color, revealing the internal brain and vascular structures. The brain tissue is shown in a reddish-brown color, and the surrounding soft tissue and vessels are in shades of blue and white. The image is set against a black background.

# Shadie: A DSL for Radiation Oncology

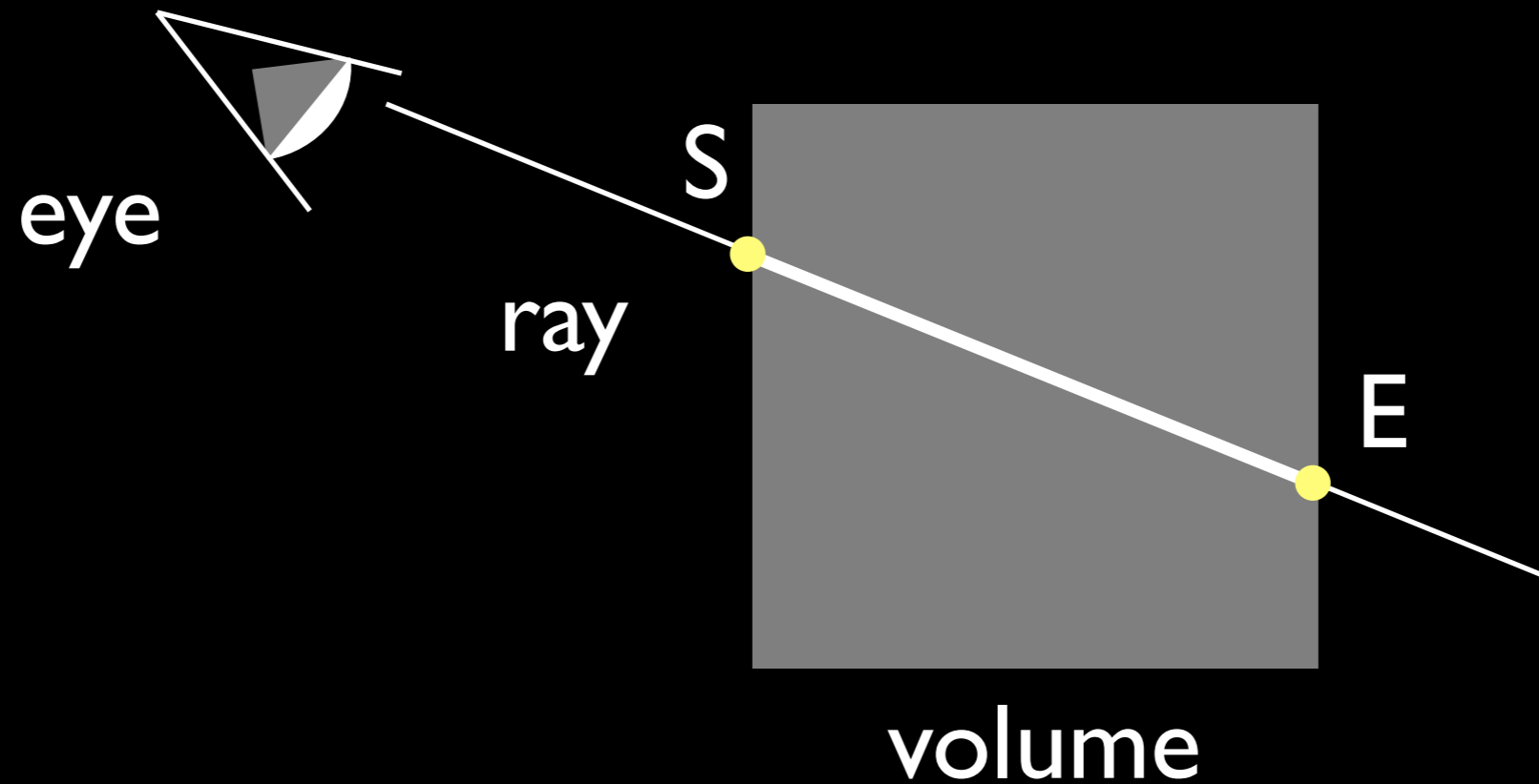


PLUNC  
3D Slicer



Shadie

# The Ray Model

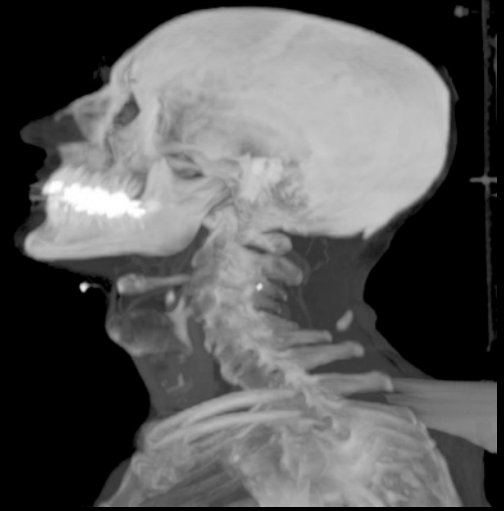


User supplies a function  $f : (\text{RayStart}, \text{RayEnd}) \rightarrow \text{Color}$

# Maximum intensity projection (MIP)



# MIP: The Shader



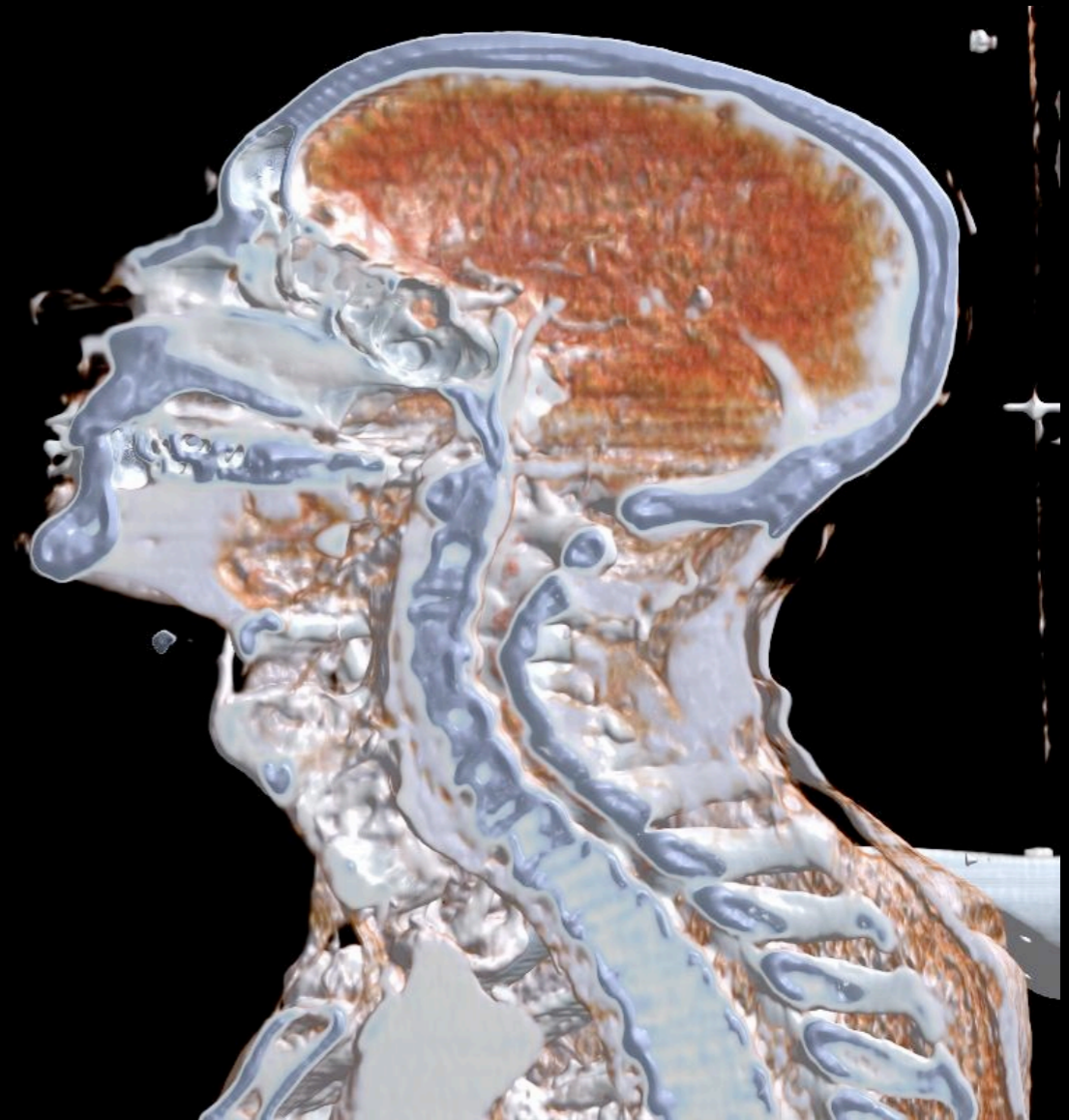
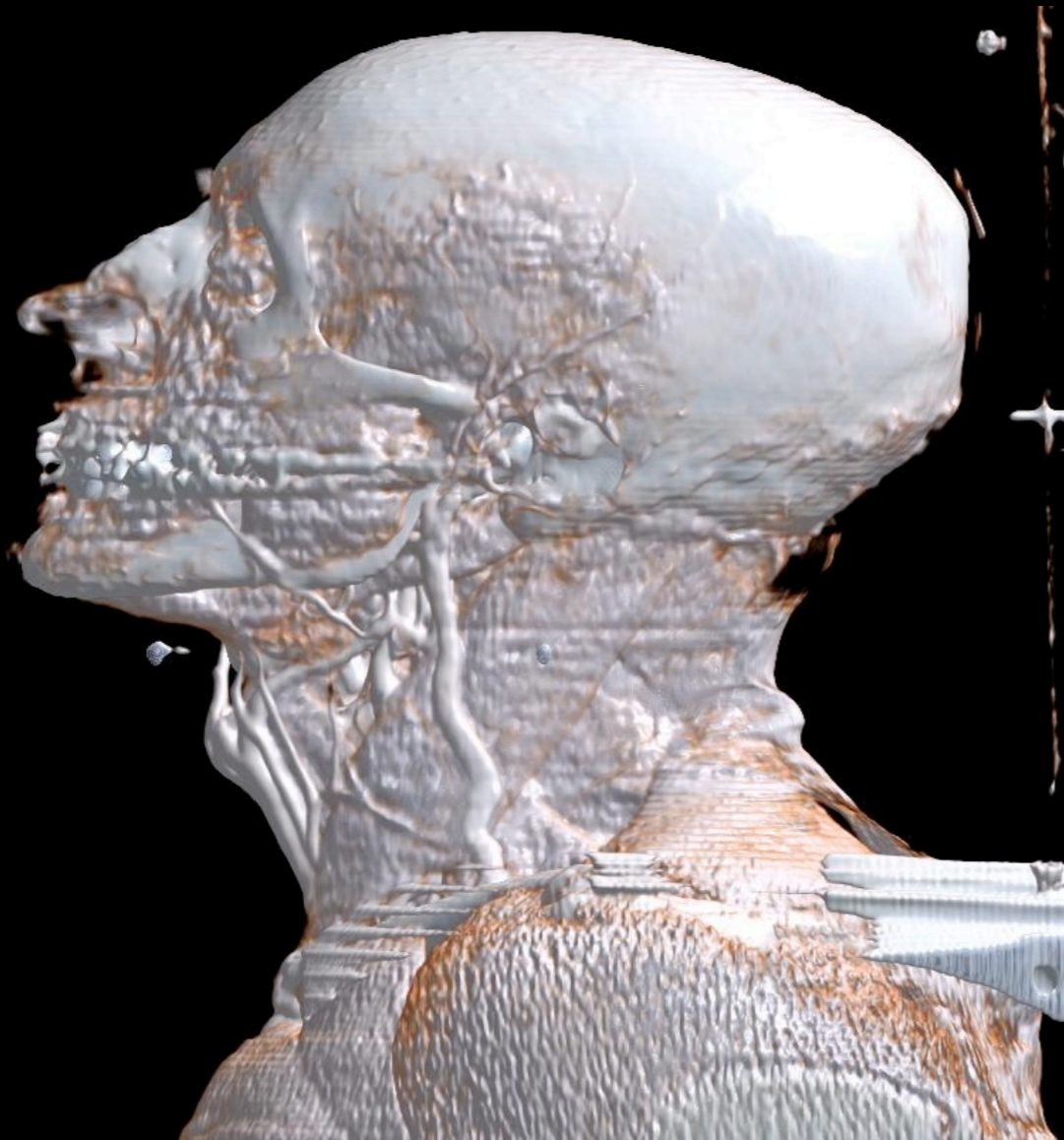
```
data = data3d('data/ct', 'short')
m = 0.0

for t in linspace(0.0, 1.0, 1000):
    # find position along ray
    P = (1-t) * S + t * E

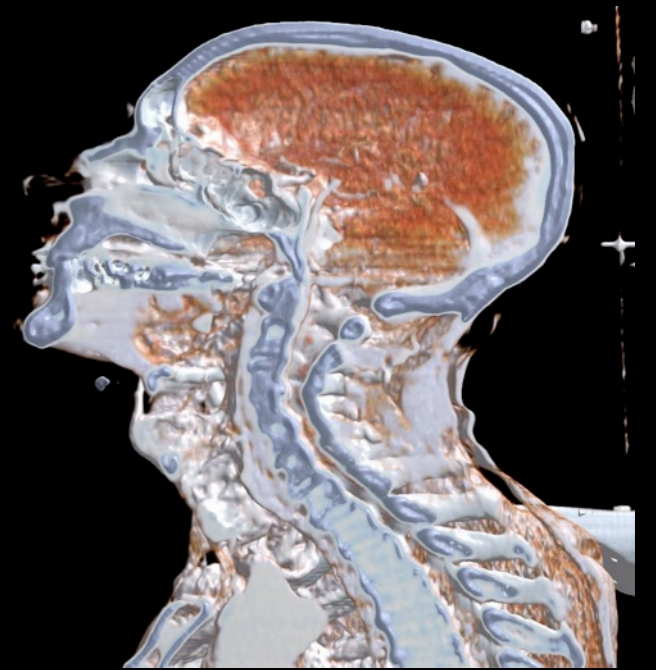
    # update maximum
    m = max(m, linear_query_3d(data, P))

return m
```

# Phong lighting + cut plane



# Phong lighting + cut plane



```
# query CT
density = cubic_query_3d_cut(data, P, D, cut)

# apply transfer function
tf_query = (density - tf_pos) / tf_width
if tf_query < 0: continue
rgba = linear_query_1d_rgba(tf, tf_query*2 - 1)

# apply phong shading
N = -normalize(cubic_gradient_3d_cut(data, P, D, cut))
L = normalize(lightpos - P)
color = phong(L, N, C, rgba.xyz, 1, 50, 0.5)
```

# CT + Dose Iso-Surface





# Multi-pass computation

```
ct = data3d(...)
```

```
def compute_rpl_slice(X, prev_slice):  
    # RPL computation at point X...
```

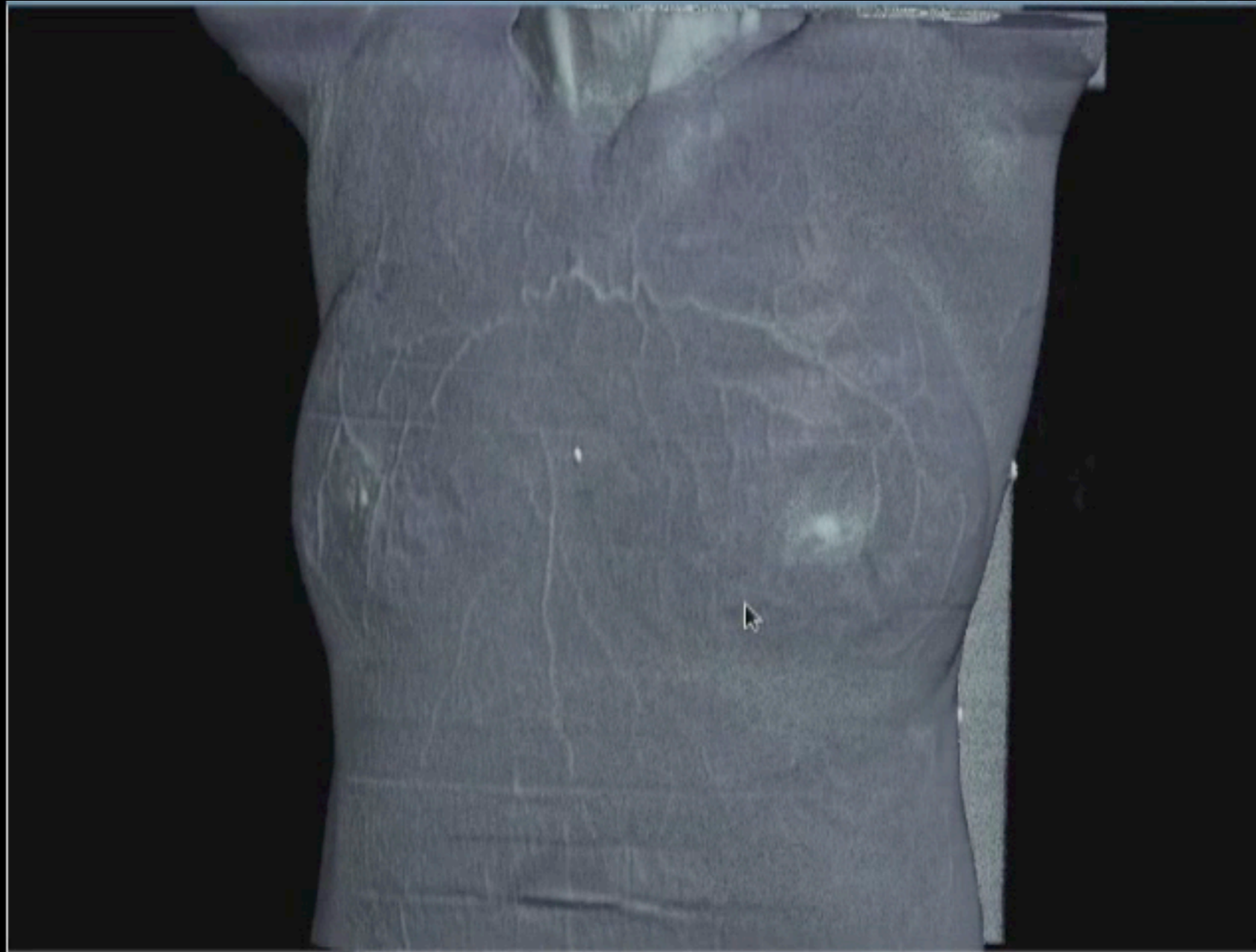
```
rpl = compute3d(compute_rpl_slice, rpl_size)
```

```
def compute_dose(X):  
    # dose computation at point X...
```

```
dose = compute3d(compute_dose, dose_size)
```

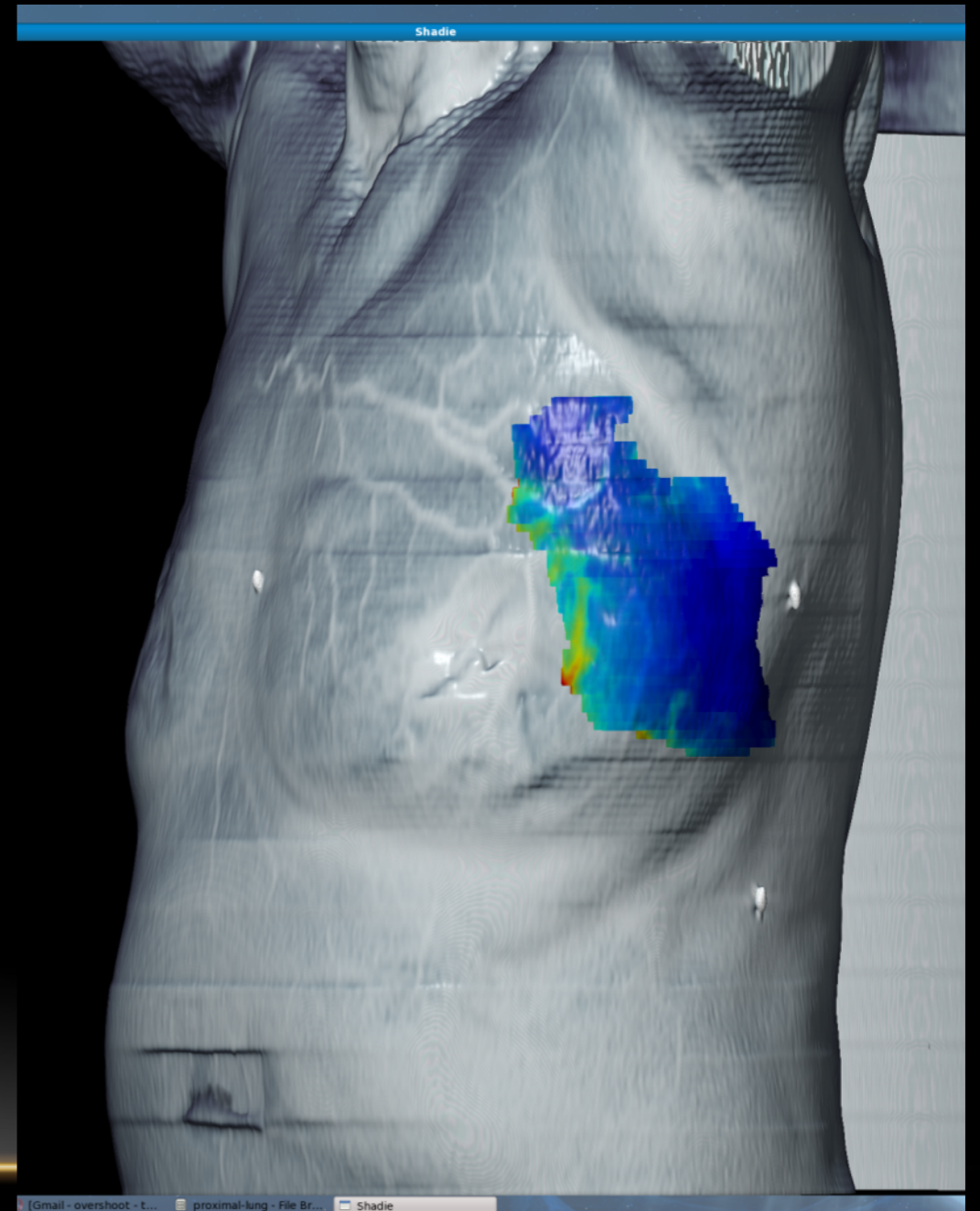
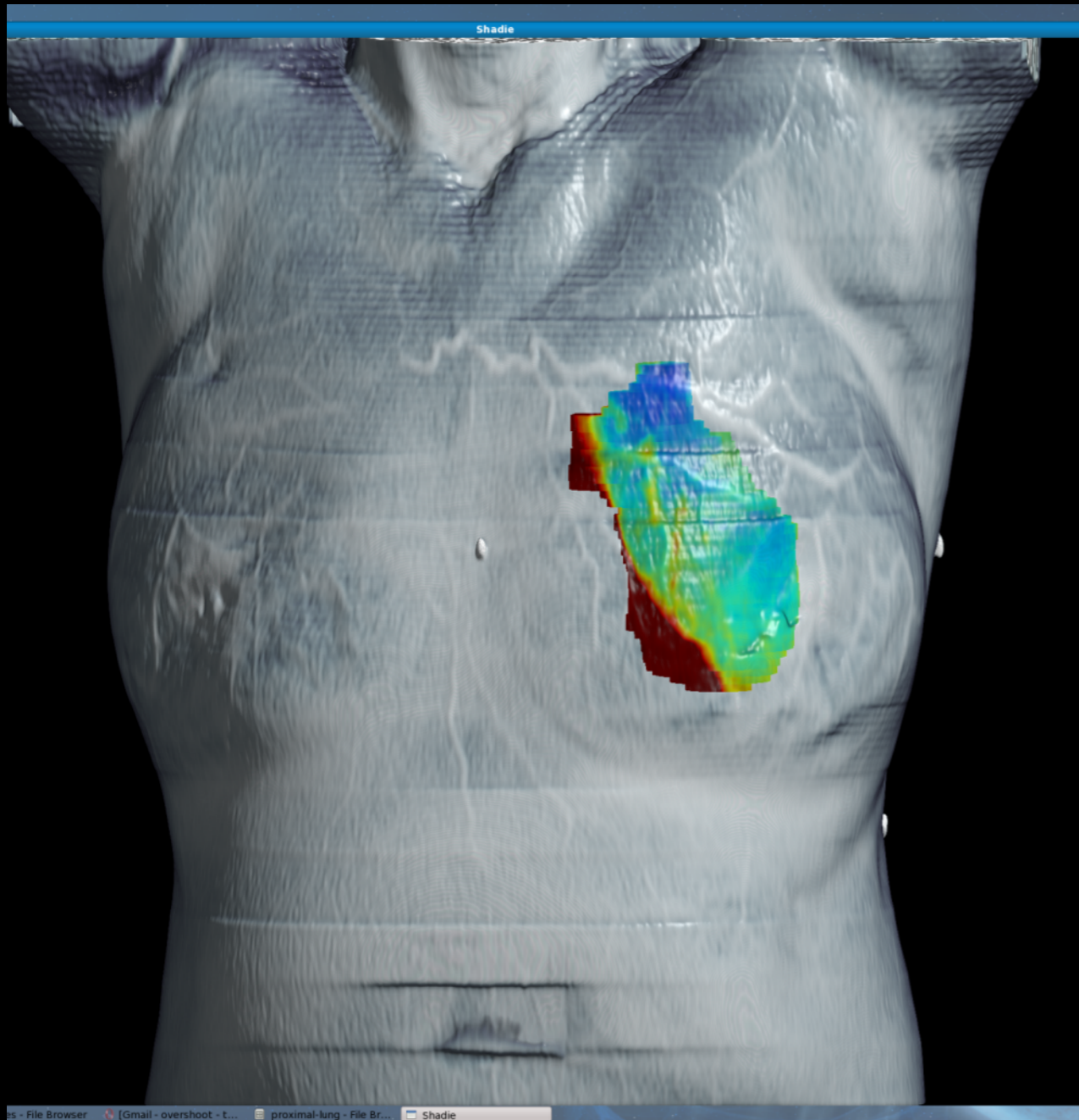
```
def main(X, D):  
    # the main visualization function  
    # compute RGBA value based on CT and dose
```

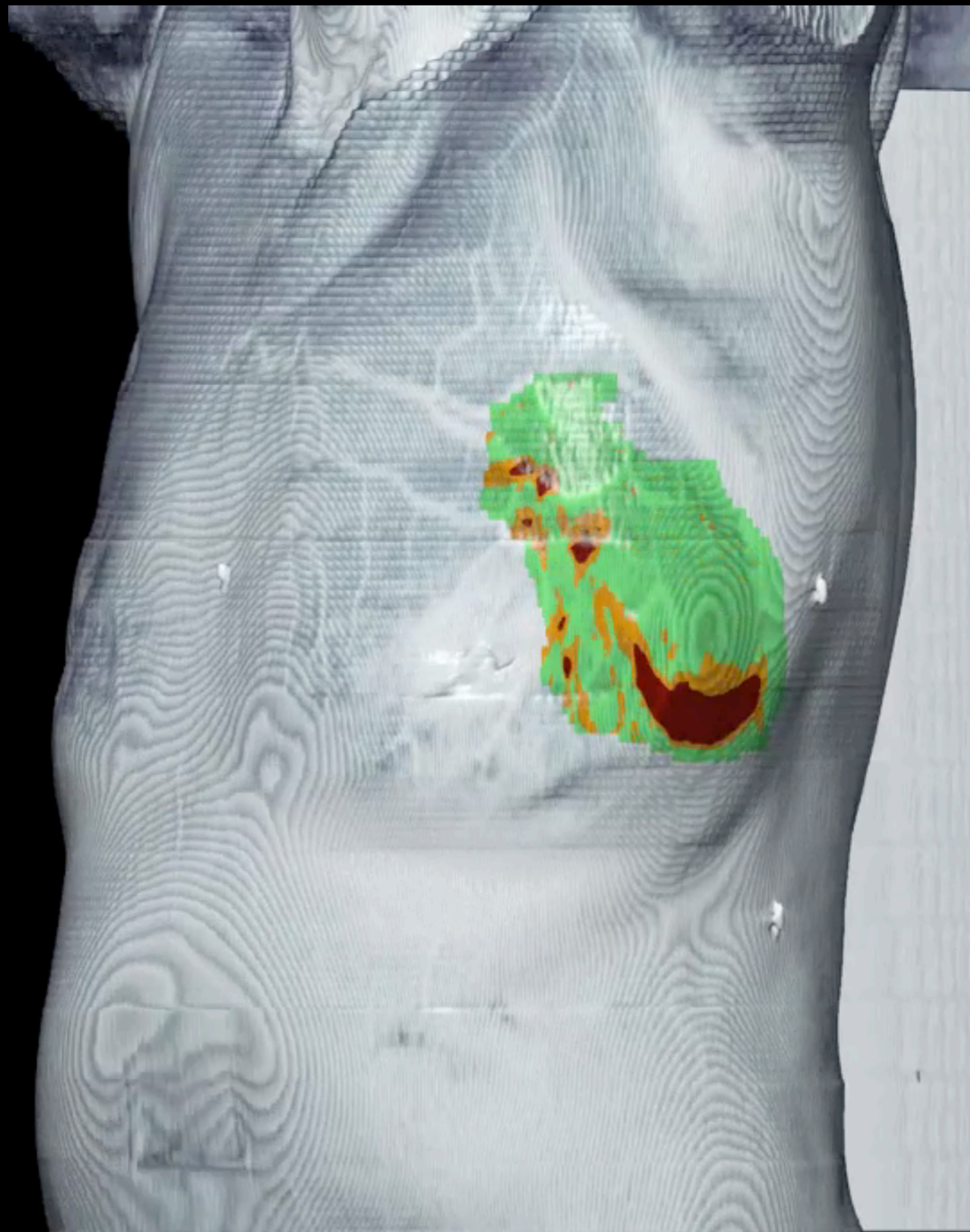
# Radiological Pathlength



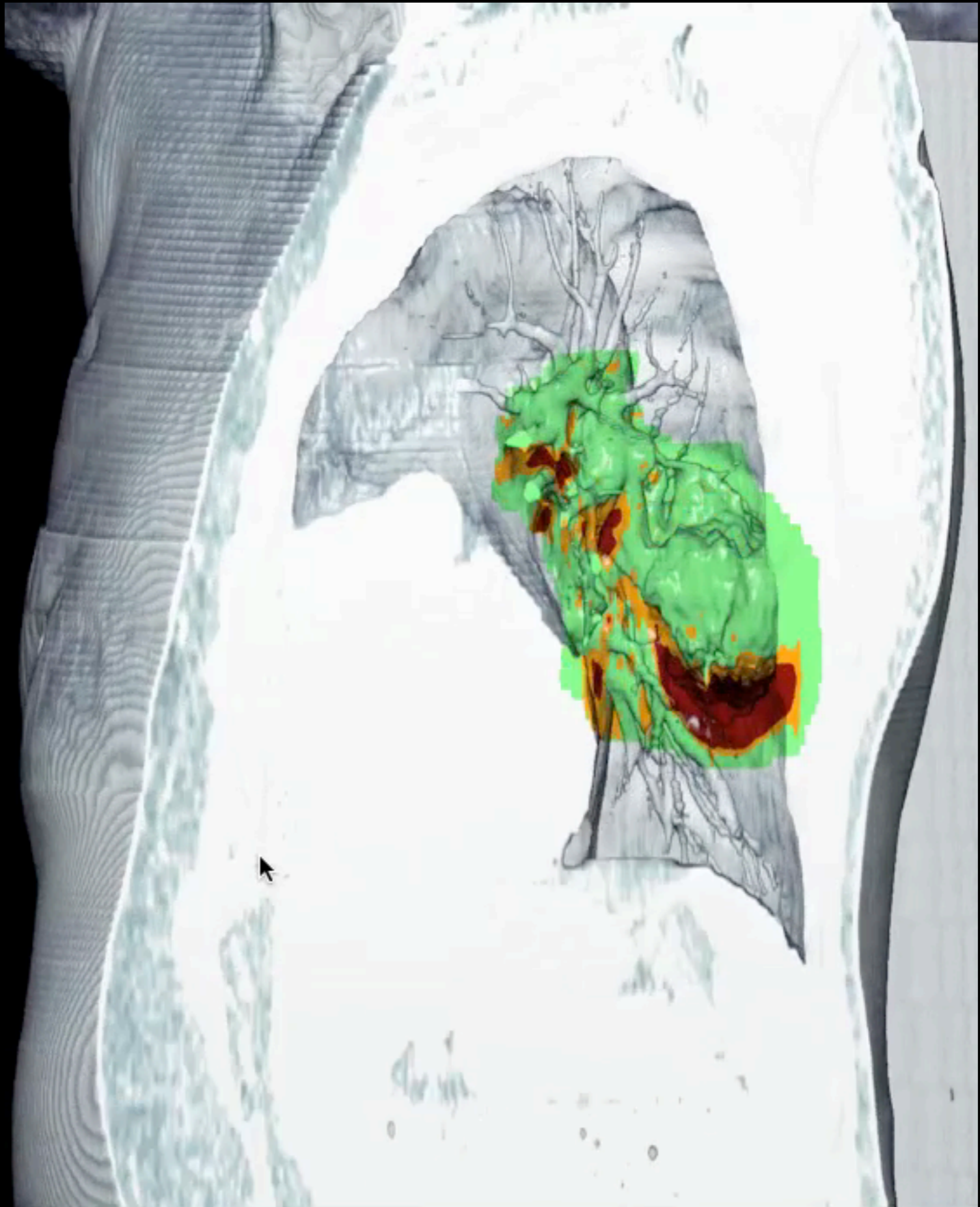
# CW to proximal ITV Shader

(includes intersected heart; uses ITV contours)

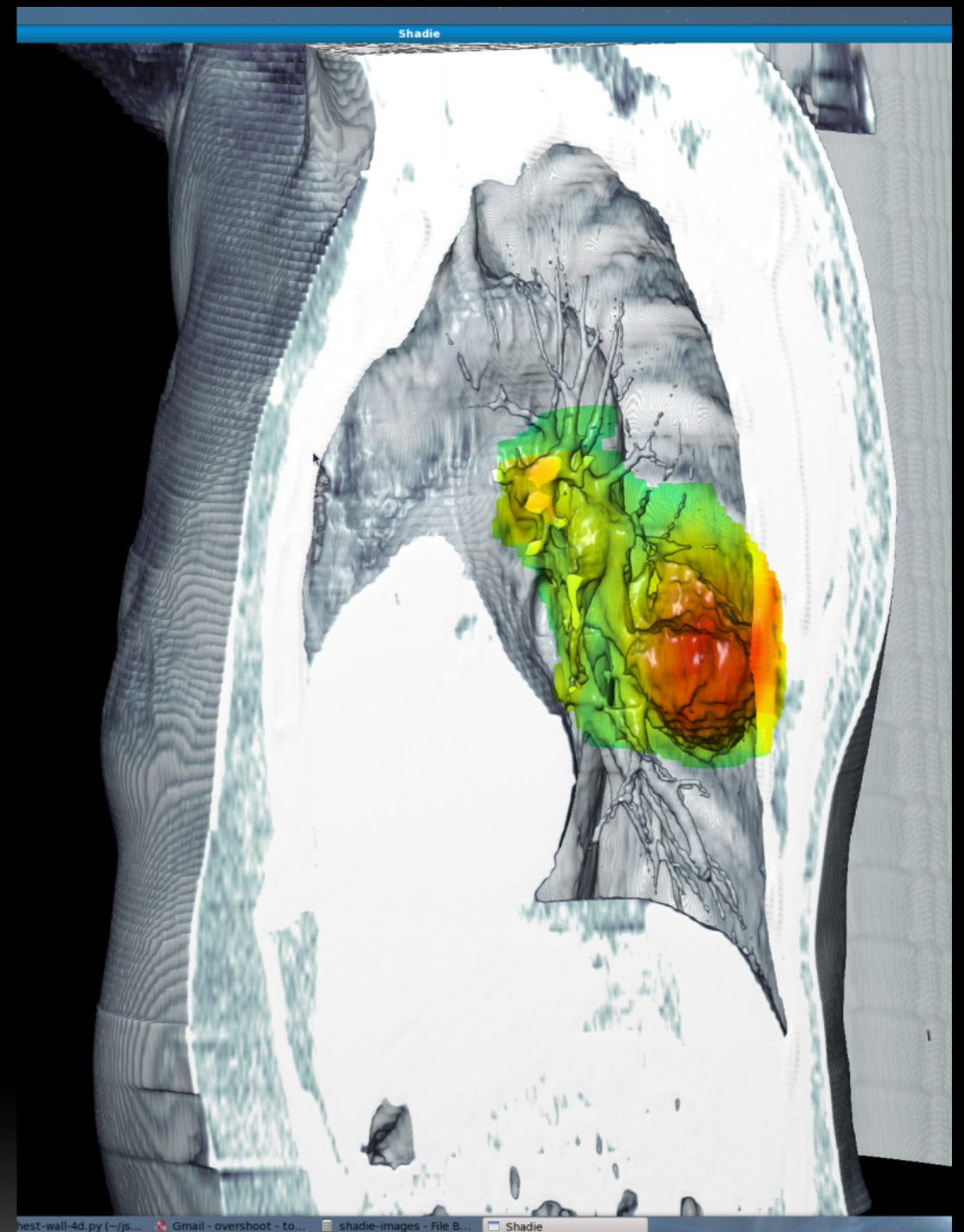
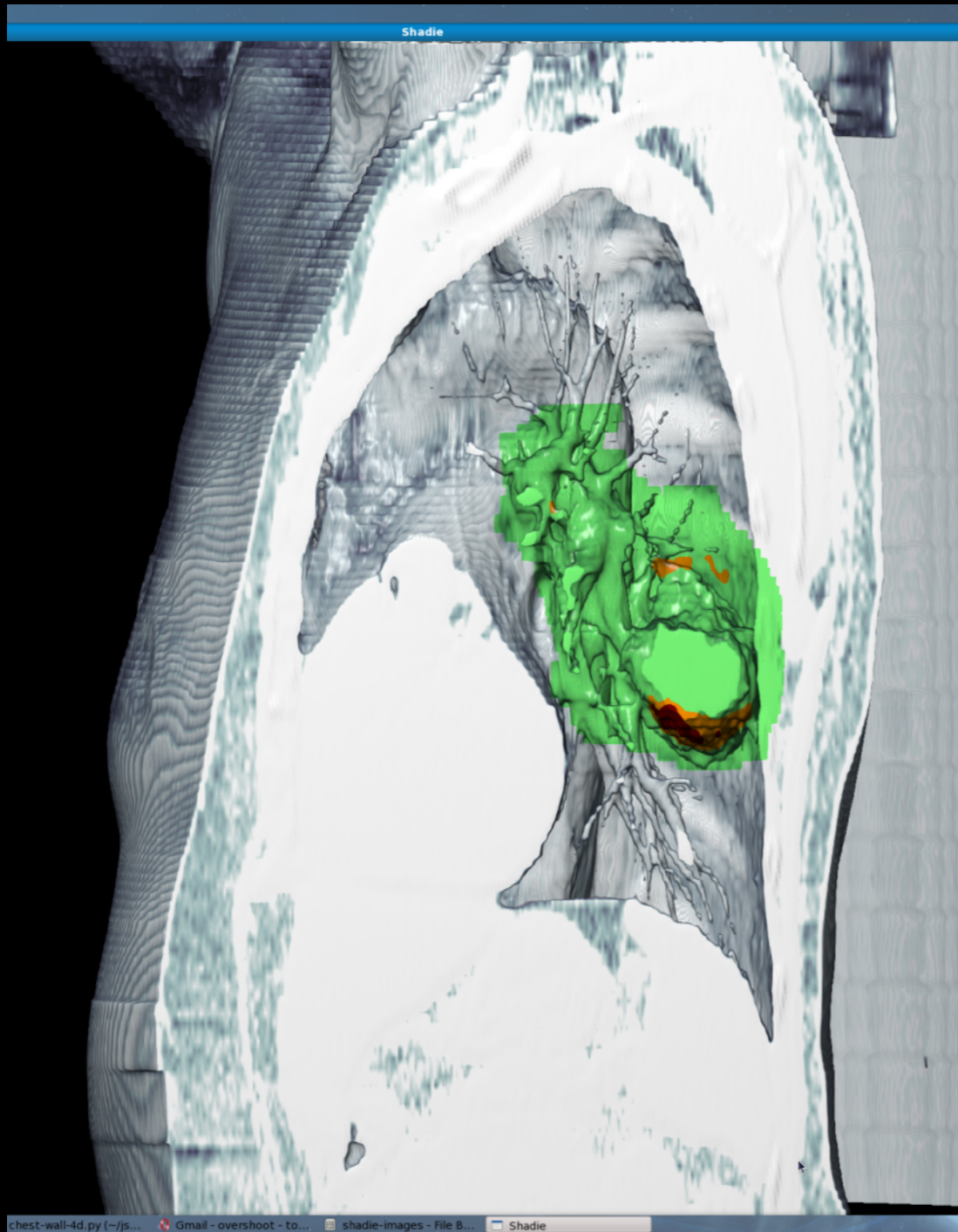




Green:  $<2\text{mm}$ ; Yellow  $<5\text{mm}$ ; Red  $>5\text{mm}$



# Overshoot



Time averaged

$\sigma$

# Status & Future

- First implementation finished and online
  - <http://code.google.com/p/shadie/>
- Several shaders implemented at MGH
- Much more to on CS and medical side
  - Back ends, runtime system, computational kernels, uncertainty, more applications, etc. etc. etc.

# Thank You

- The project was supported by the Federal Share of program income earned by Massachusetts General Hospital on C06 CA059267, Proton Therapy Research and Treatment Center

