

# Semi-Automatic Central-Chest Lymph-Node Definition from 3D MDCT Images

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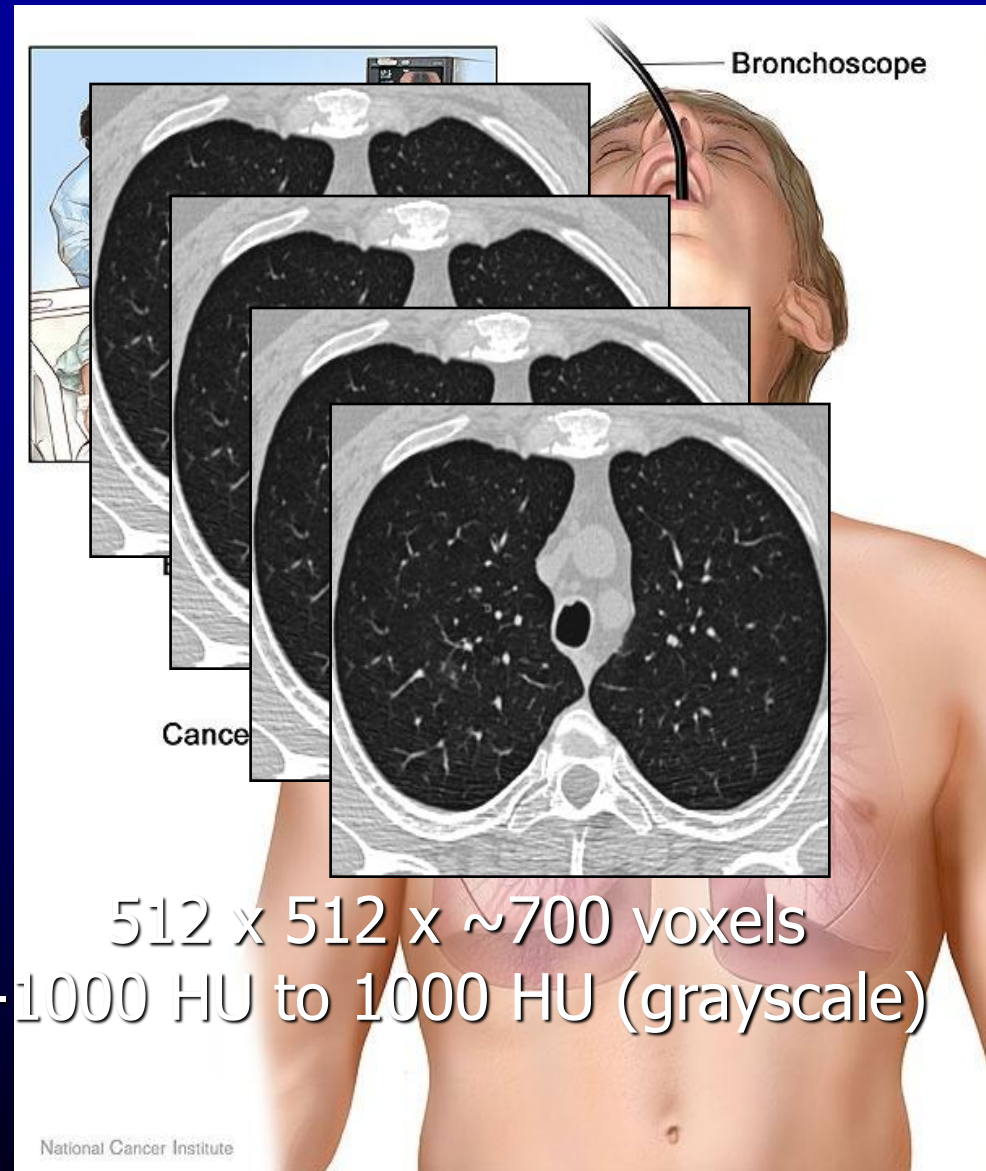
*Penn State University*

*University Park and Hershey, PA, USA*

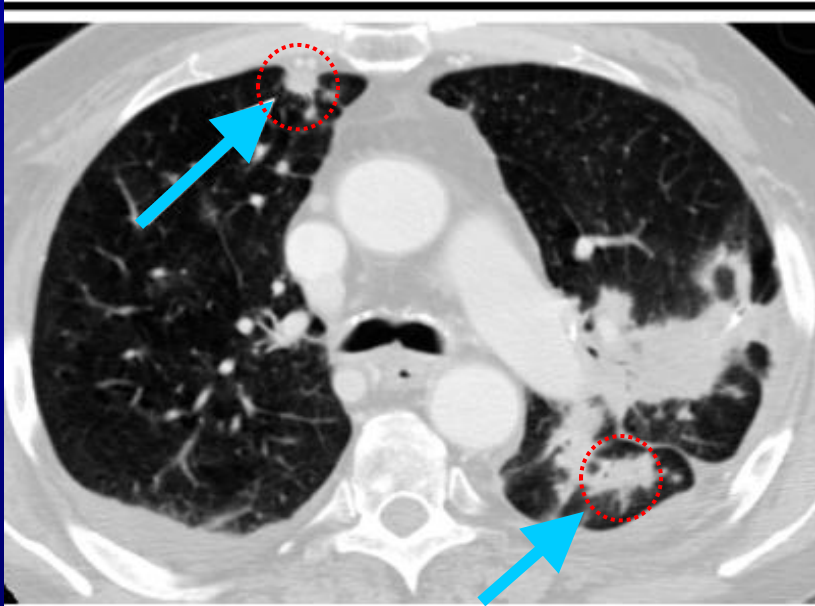


# Motivation

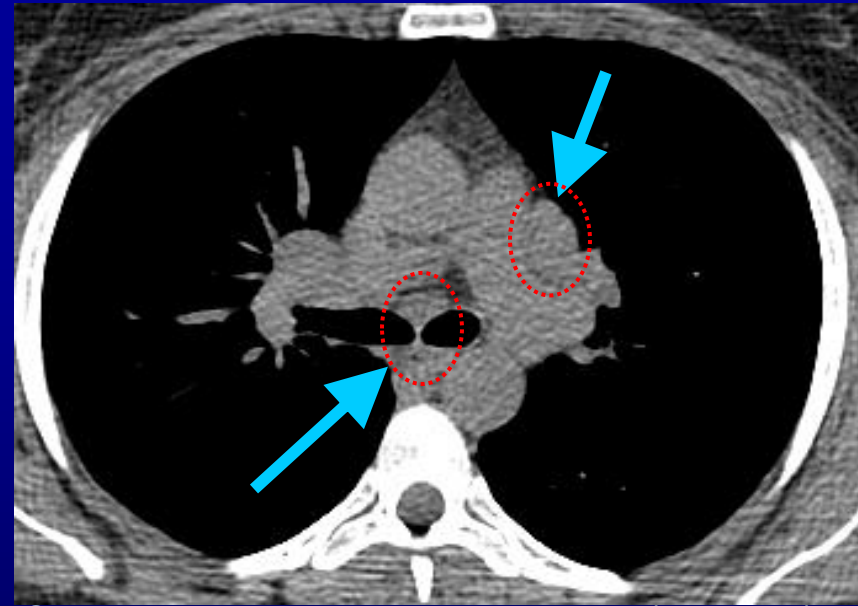
- Lung cancer is the leading cause of cancer death
- Early lung cancer detection could increase survival rate
- Diagnosis procedures:
  - 3D MDCT chest image assessment
  - Follow-on diagnostic bronchoscopy



# Motivation



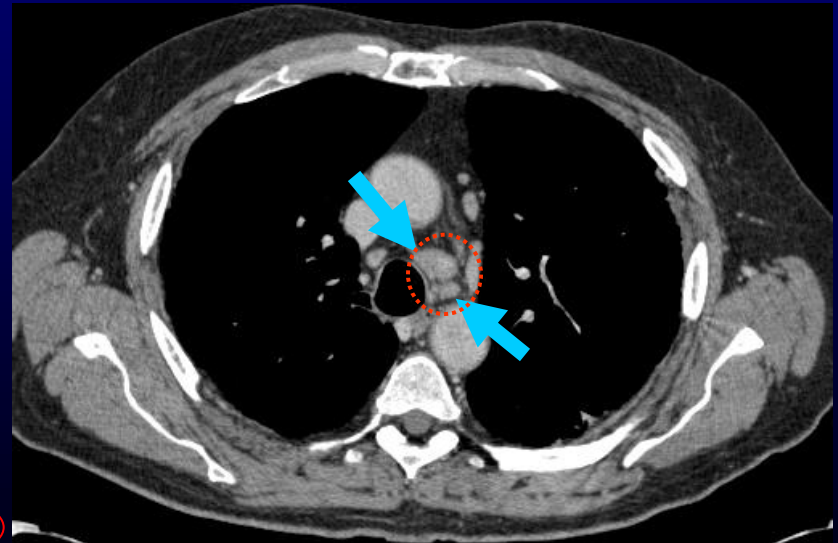
Armato *et al.*, *Radiology* 2004



Subaortic and subcarinal lymph node (p2h012b)

- ROIs involve complex phenomena
- Manual slice tracing – not feasible
- Automatic methods
  - application dependent
- Semi-automatic methods
  - more practical

**Live Wire**



Lower para-tracheal lymph node (IRB20349.3.3)

# Prior Work

## 2D Live Wire:

- Mortensen and Barrett (*Graphical Models and Medical Imaging 1998*)
- Falcão *et al.* (*Graphical Models and Medical Imaging 1998*)
- Lu and Higgins (*Int. J. CARS 2007*)

## 3D Live Wire:

- Falcão and Udupa (*Medical Image Analysis 2000*)
- Hamarneh *et al.* (*SPIE Medical Imaging 2005*)
- König and Hesser (*SPIE Medical Imaging 2005*)
- Souza *et al.* (*SPIE Medical Imaging 2006*)
- Lu and Higgins (*Int. J. CARS 2007*)
- Poon *et al.* (*SPIE Medical Imaging 2007 and CMIG 2008*)

# Prior Work

## Cost Function:

$$l(p, q) = w_z f_z(q) + w_G f_G(q) + w_D f_D(p, q) + w_{Df} f_{Df}(p, q)$$

## Dynamic Graphic Search:

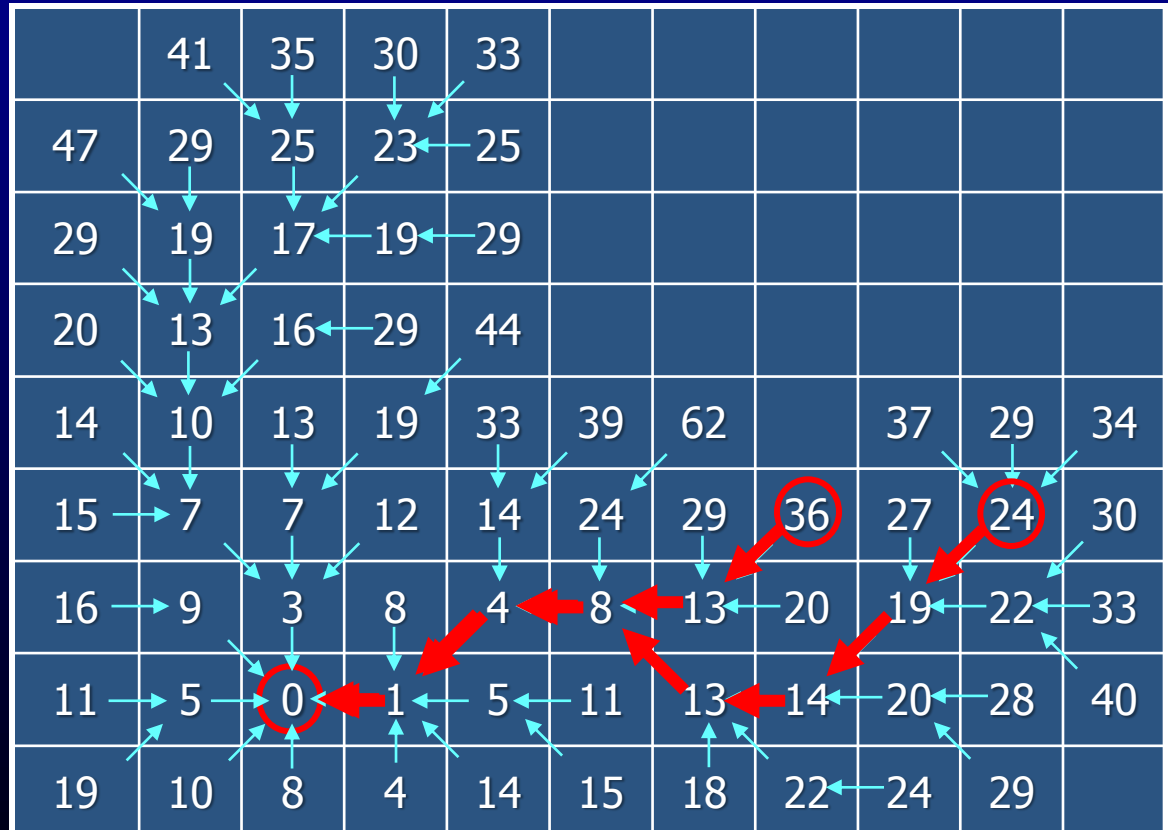
**Input:**  
 $s$   
 $l(p, r)$

**Data Structures:**  
 $L$   
 $N(p)$   
 $e(p)$   
 $g(p)$

**Output**  
 $pt$

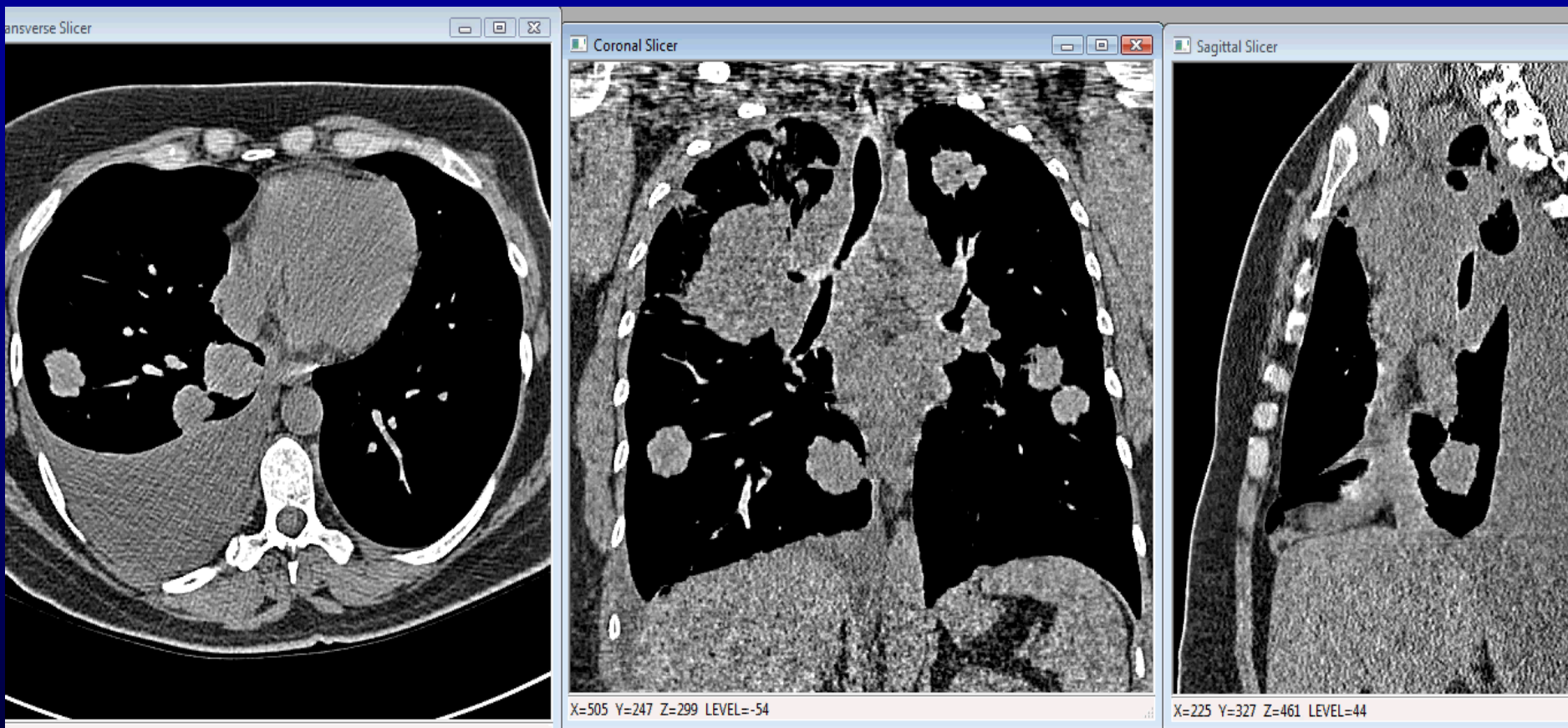
**Algorithm:**

1.  $g(s) \leftarrow 0; L \leftarrow s;$
2. While  $L \neq \Phi$  do begin
3.      $pt \leftarrow \min(L);$
4.      $e(p) \leftarrow TRUE;$
5.     for each  $r \in N(p)$  such that  $\neg e(r)$  do begin
6.          $gtemp \leftarrow g(p) + l(p, r);$
7.         if  $r \in L$  and  $gtemp < g(r)$  then
8.              $g(r) \leftarrow gtemp; pt(r) \leftarrow p;$
9.         if  $r \notin L$  then begin
10.              $g(r) \leftarrow gtemp; pt(r) \leftarrow p; L \leftarrow r;$
11.         end
12.     end
13. end
14. end
15. end



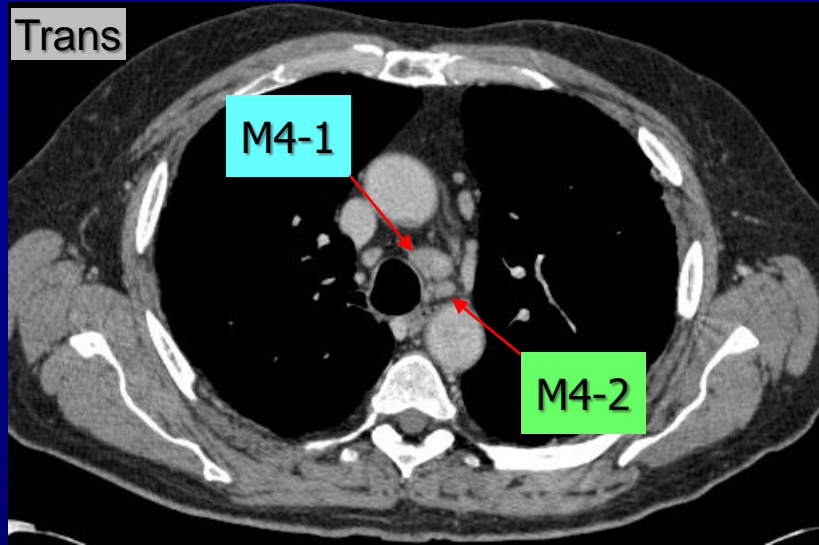


# Prior Work

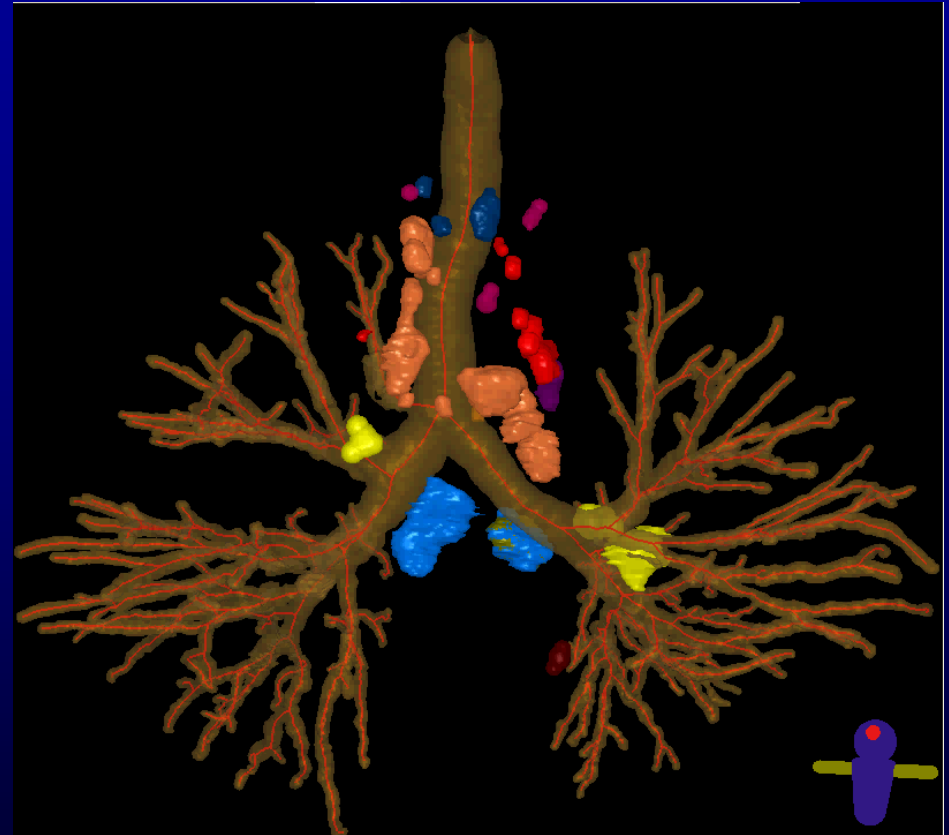
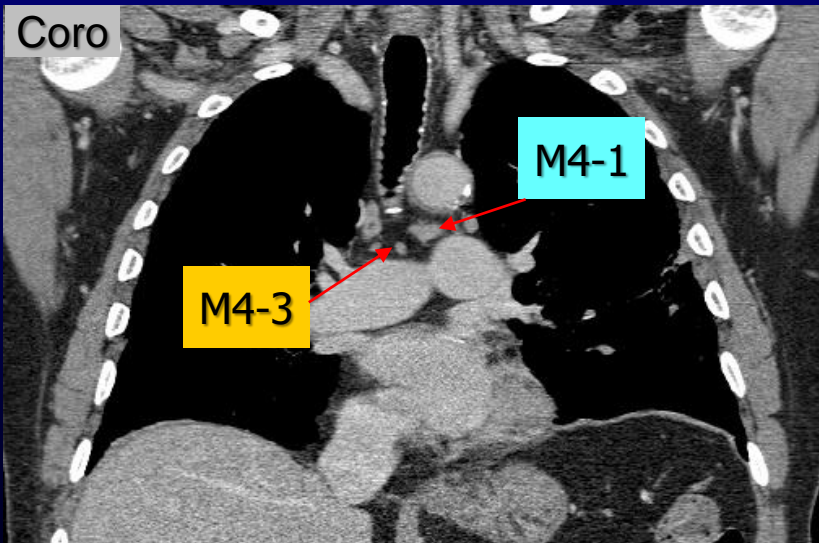


Segment ROI using 3D Live Wire

# Central-Chest Lymph Nodes in 3D MDCT Image



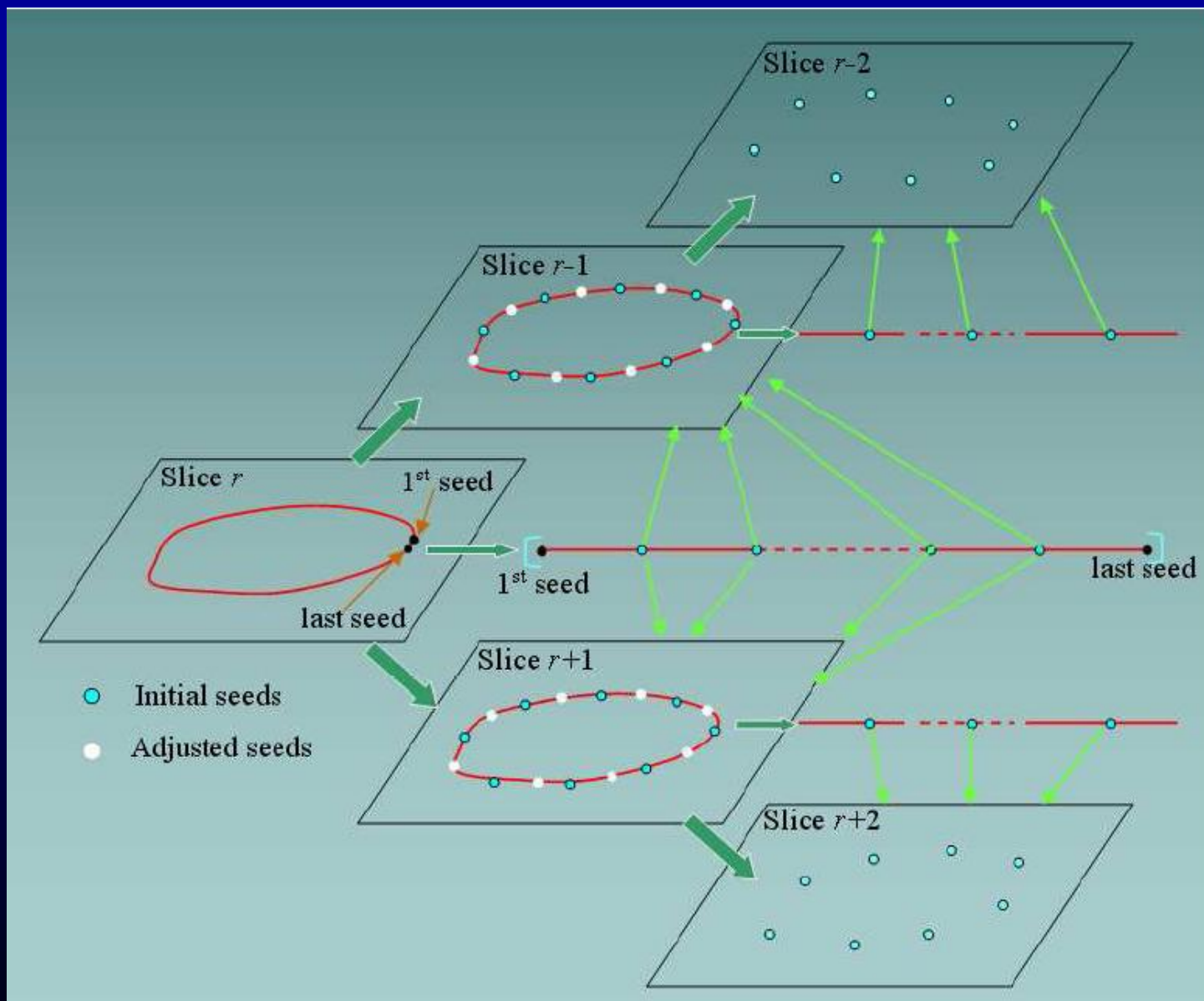
Case IRB20349.3.3



Complete segmentation of  
central-chest lymph nodes

Typical lower paratracheal (M4)  
lymph nodes

# Single-Section Live Wire





# Single-Section Live Wire

```

Inputs:
i, j /* Iterators */
n /* Counter */
r, Br /* Reference section index and associated reference boundary */
imin, imax /* Min and max section indices for running automatic 3D
live wire in direction Dtw */
Si /* Seed set for section i, defined based on Bref */

Data Structures:
Bref(i), Cobj /* Reference object boundary for section i and its accumulated cost */
Lobj, N /* List of pixels along Bobj(i) and its size. */
Tn, Tcost /* Pixel-number and cost thresholds. By default, Tn = 16. */

Output
Bobj(i) /* Final ROI boundary for section i, imin ≤ i ≤ imax */

Algorithm:
01. i = r + 1, k = 1 /* Compute 2D boundaries for upper portion of ROI */
02. Bobj = Bref(i) = Br /* Use Br as the reference section for section i */
03. while i ≤ imax do
04. do
05. if N ≤ T then n = 4 /* Reduce the size of Si if N is small enough */
06. else n = 8
07. end
08. while j ≤ n do
09. S(i) ← Lobj(j * N/n) /* Select pixels evenly along Bref */
10. end
11. 2DLW(Si) → Bobj(i), cobj /* Connect seeds in Si using 2D live wire and output section
i's boundary and cobj */
12. Update(Lobj) → Lobj /* Add boundary pixels to Lobj starting from the midpoint
from the segment between Si(1) and Si(n) on Bobj */
13. if k == 1 then /* Record the boundary cost as a reference for next iteration */
14. c̃obj = cobj
15. end
16. k = k + 1
17. while  $|c_{obj} - \tilde{c}_{obj}| > \frac{\tilde{c}_{obj}}{100}$ 
18. if Stopcheck() → FALSE then /* Process next section unless the boundary cost is
too small or stopping constraints are met. */
19. i = i + 1 and Bref(i) = Bobj
20. end
21. end
22. i = r - 1, k = 1 /* Compute 2D boundaries for lower portion of ROI */
23. Bobj = Bref(i) = Br /* Use Br as the reference section for section i */
24. while i ≥ imin do
25. do
26. if N ≤ T then n = 4 /* Reduce the size of Si if N is small enough */
27. else n = 8
28. end
29. while j ≤ n do
30. S(i) ← Lobj(j * N/n) /* Select pixels evenly along Bref */
31. end
32. 2DLW(Si) → Bobj(i), cobj /* Connect seeds in Si using 2D live wire and output section
i's boundary and cobj */
33. Update(Lobj) → Lobj /* Add boundary pixels to Lobj starting from the mid point
from the segment between Si(1) and Si(n) on Bobj */
34. if k == 1 then /* Record the boundary cost as a reference for next iteration */
35. c̃obj = cobj
36. end
37. k = k + 1
38. while  $|c_{obj} - \tilde{c}_{obj}| > \frac{\tilde{c}_{obj}}{100}$ 
39. if Stopcheck() → FALSE then /* Process next section unless the boundary cost is
too small or stopping constraints are met. */
40. i = i - 1 and Bref(i) = Bobj
41. end
42. end

```

➤ Adjust working area adaptively

➤ Refine seed set iteratively

➤ Terminate iterations when

$$|c_{obj} - \tilde{c}_{obj}| < \frac{\tilde{c}_{obj}}{100};$$

➤ Terminate 3D process when

➤ Section limits reached

➤ Boundary costs vary greatly

$$|c_{obj} - c_r| > \frac{c_r}{10}.$$

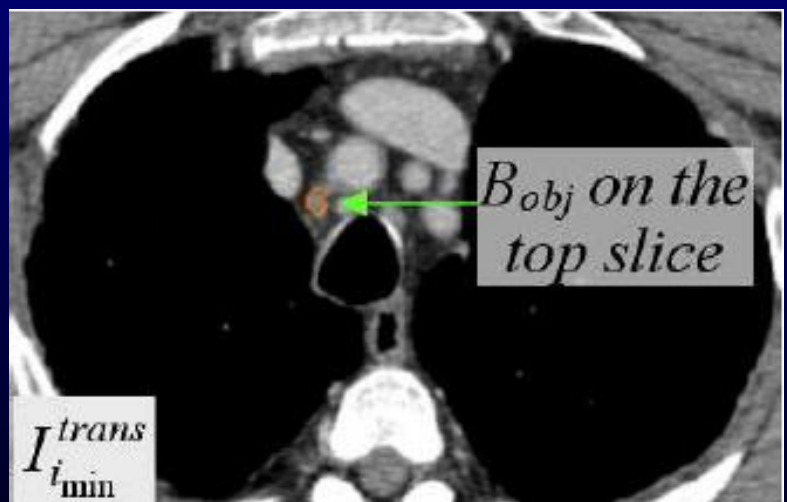
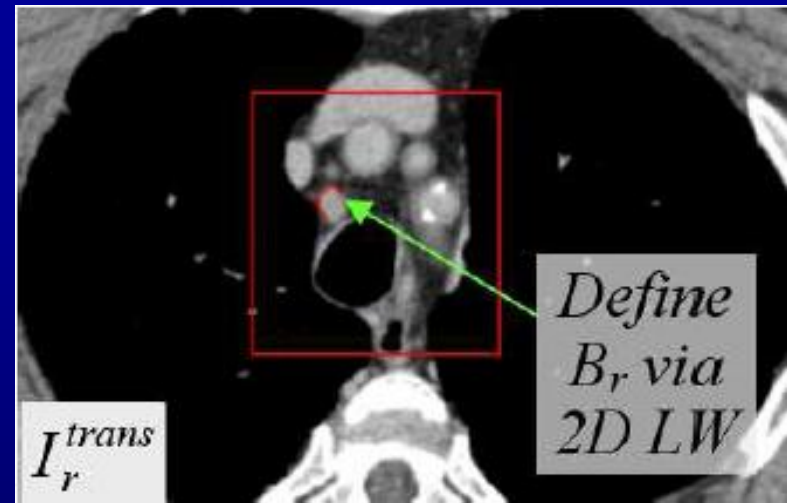
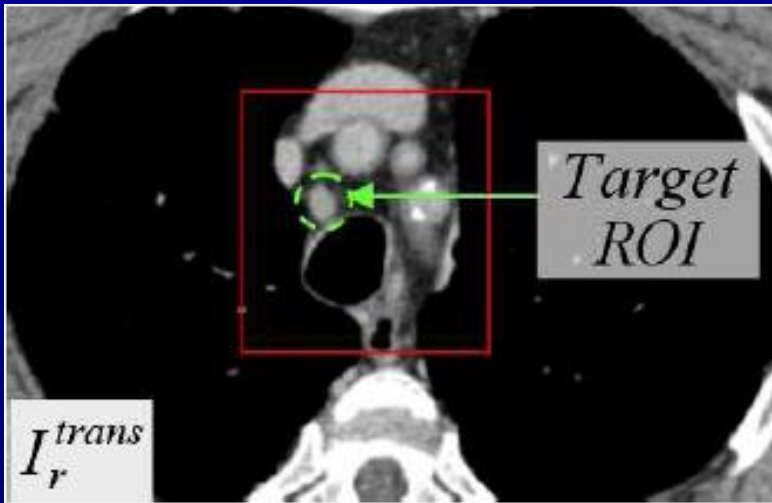
➤ Boundary is too small

➤ Segmented region is too small

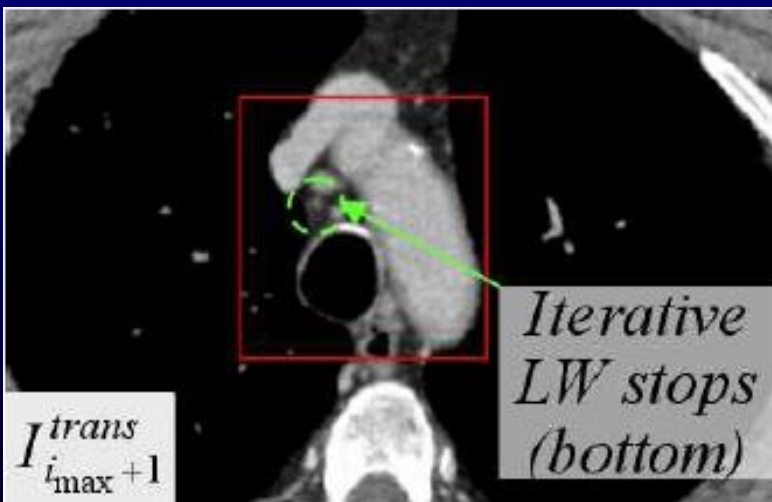
➤ Intensity distribution varies greatly

$$N_i^{out} > 0.5 \cdot N_i^{obj}$$

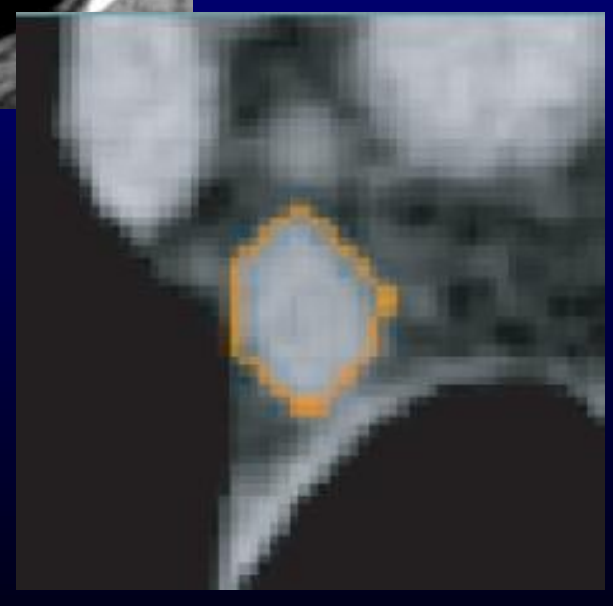
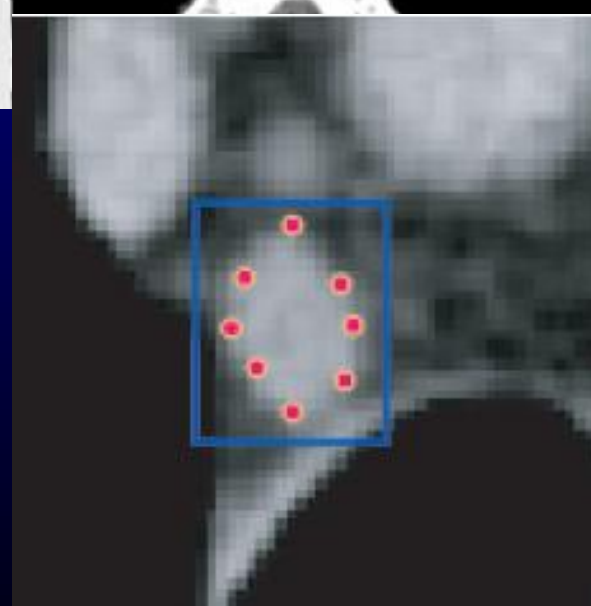
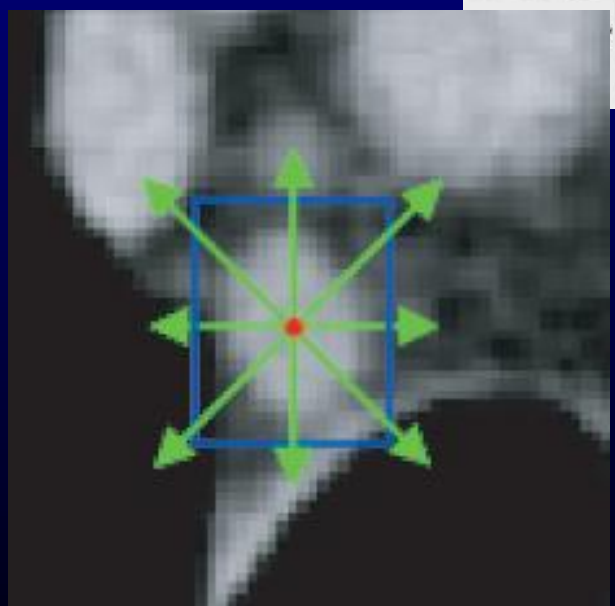
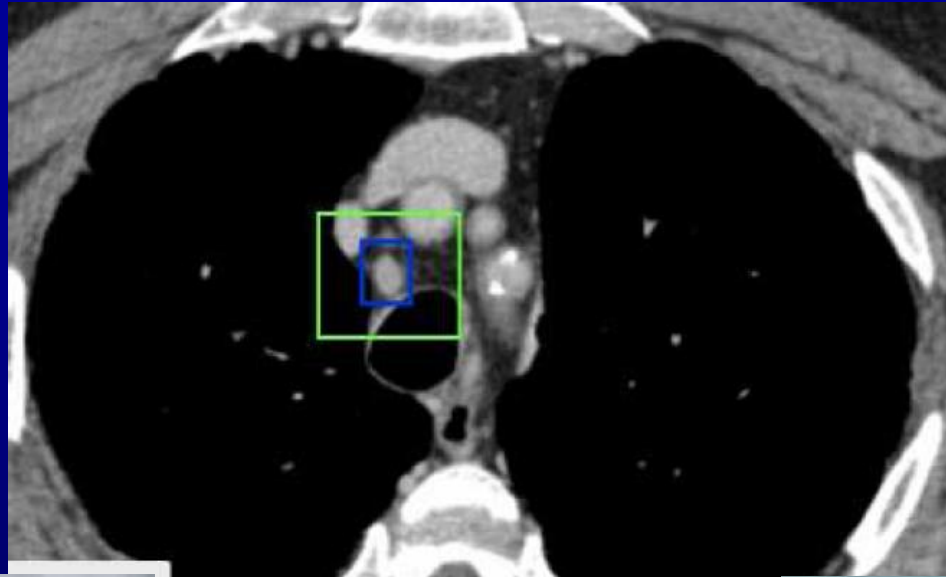
# Single-Section Live Wire



# Single-Section Live Wire

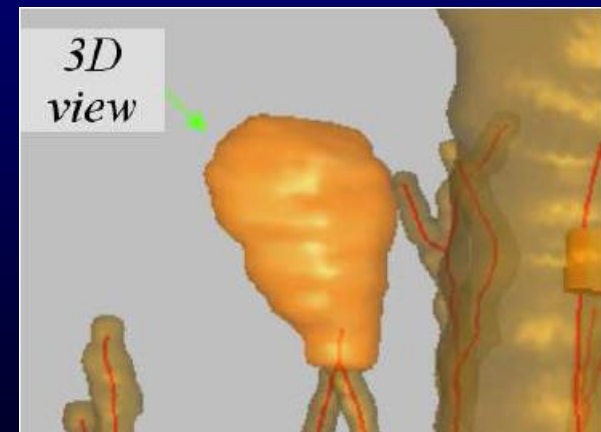
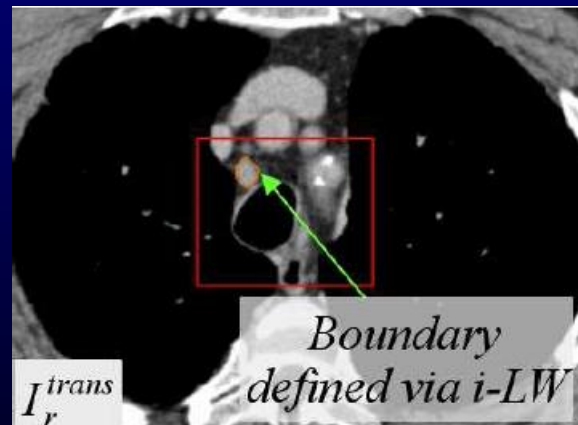
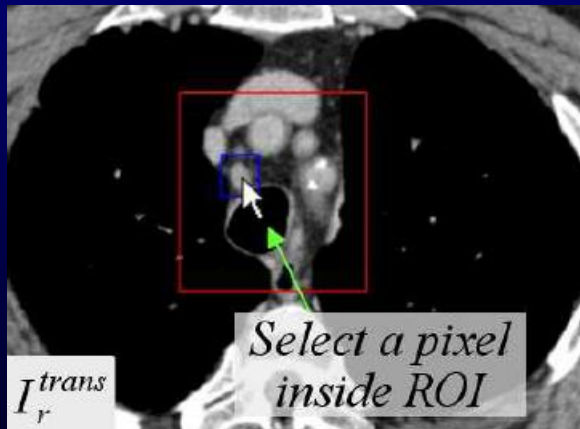
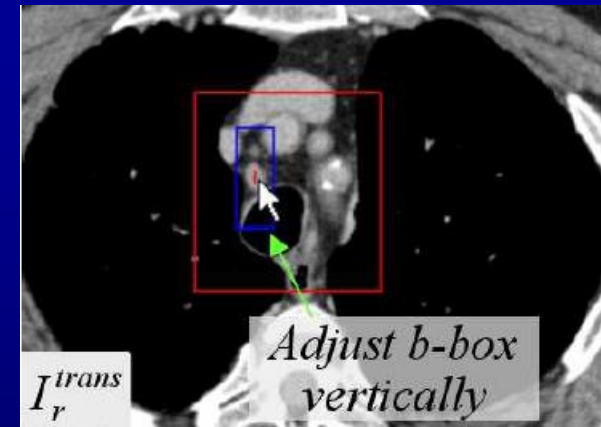
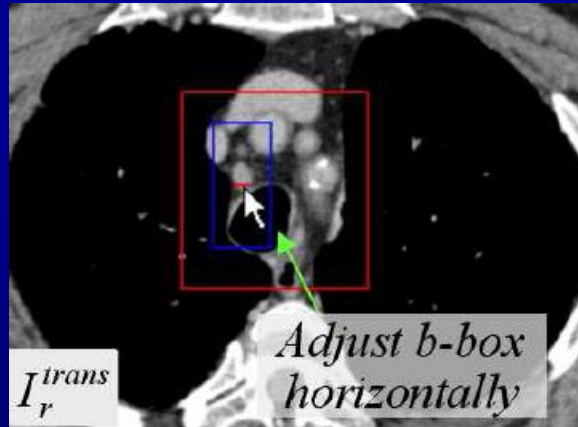
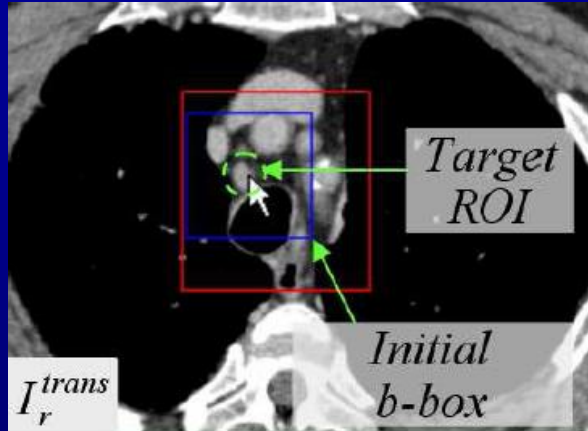


# Single-Click Live Wire

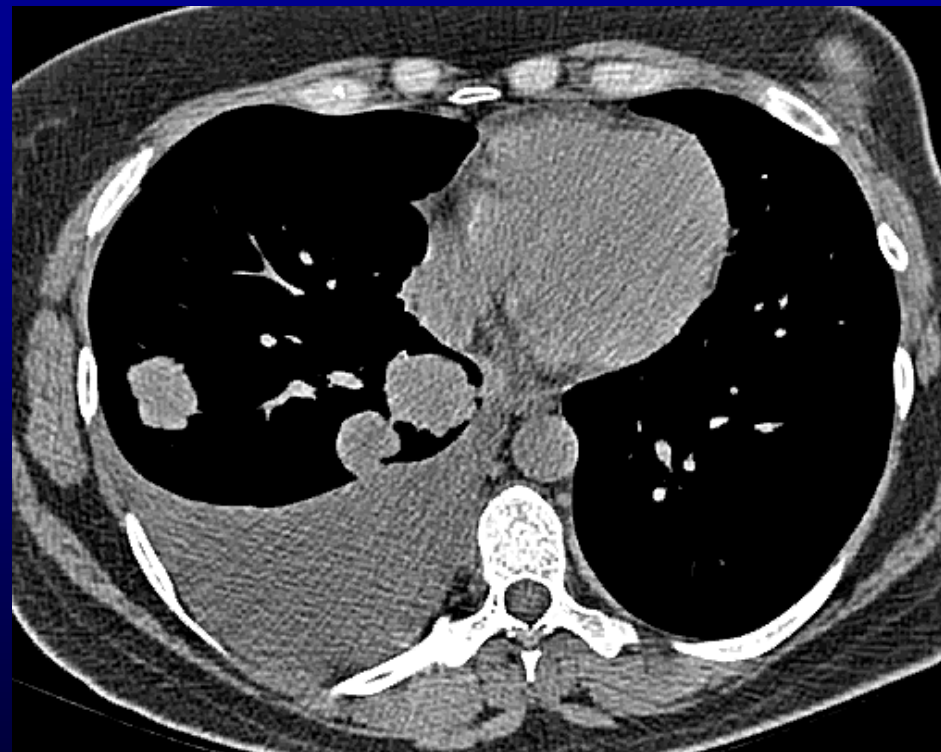




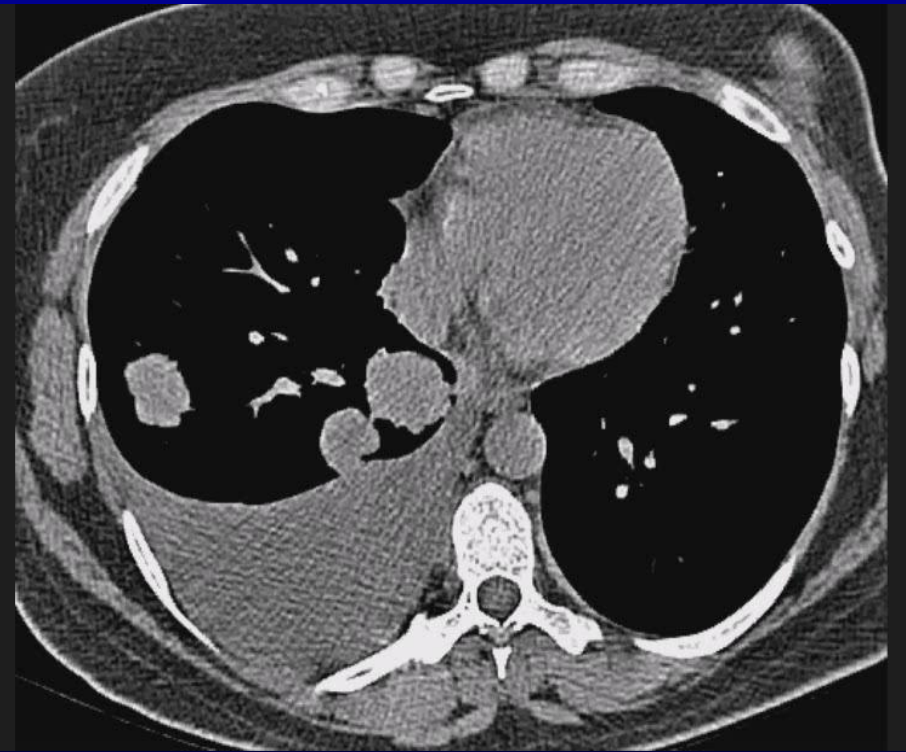
# Single-Click Live Wire



# Single-Section and Single-Click LW



Single-Section



Single-Click

# Performance Evaluation

## MDCT Data:

Case	Voxel Dim.(mm)			# of slice in scan	Contrast- Enhanced	Observer $\mathcal{O}$	Scan ID
	$\Delta X$	$\Delta Y$	$\Delta Z$				
21405_64	0.64	0.64	0.5	702	N	$\mathcal{O}_1$	1
20349_3_3	0.72	0.72	0.5	578	N	$\mathcal{O}_1$	2
20349_3_15	0.68	0.68	0.5	757	N	$\mathcal{O}_1, \mathcal{O}_2$	3
20349_3_27	0.67	0.67	0.5	752	Y	$\mathcal{O}_1$	4
21405_67	0.69	0.69	0.5	716	N	$\mathcal{O}_1, \mathcal{O}_2$	5

Computer - Dell Precision 650 workstation:

- Dual Intel Xeon 3.2GHz, 3GB RAM, Windows XP

# Results

Method	Single-Section	Single-Click
Number of Nodes	50	
Success Rate	90 % (45/50)	80% (40/50)
Accuracy	81±7%	79±8%
Inter-Trial Reproducibility	88±7%	86±9%
Processing Time	16±4s	20±5s

## Segmented Nodes:

- ❑ Short-Axis Length: 5.8±1.5 mm
- ❑ Long-Axis Length: 11±4.0 mm
- ❑ Volume: 256±210 mm<sup>3</sup>
- ❑ Number of Voxels: 1089±861

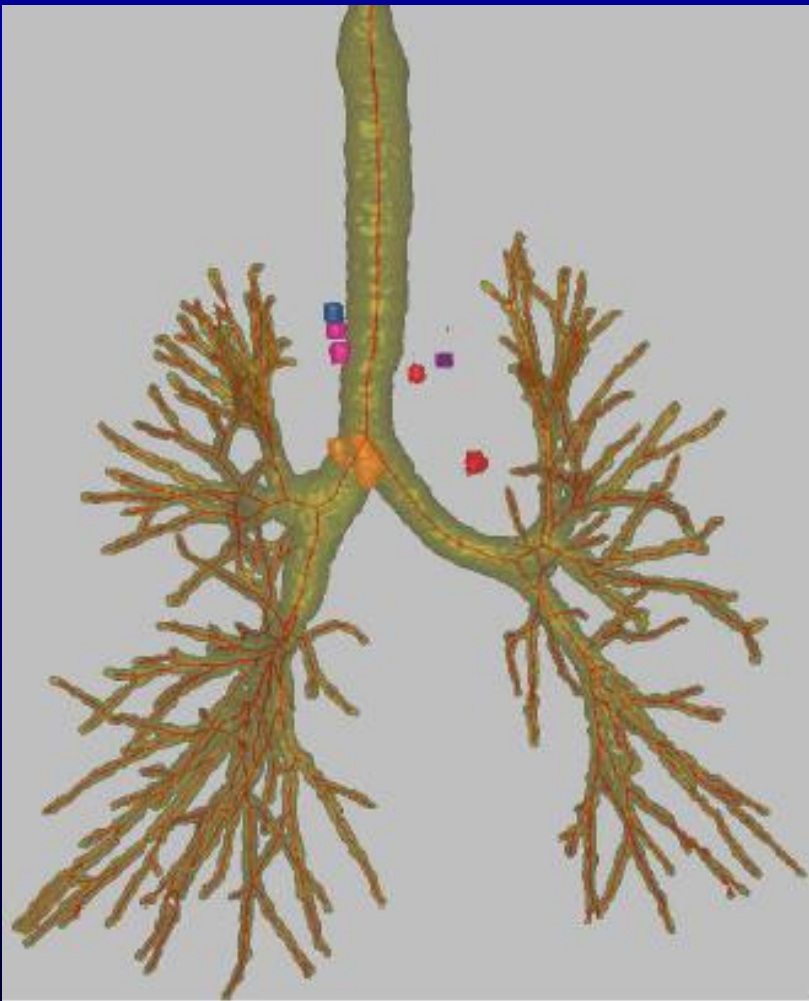


# Comparison of Two Observers

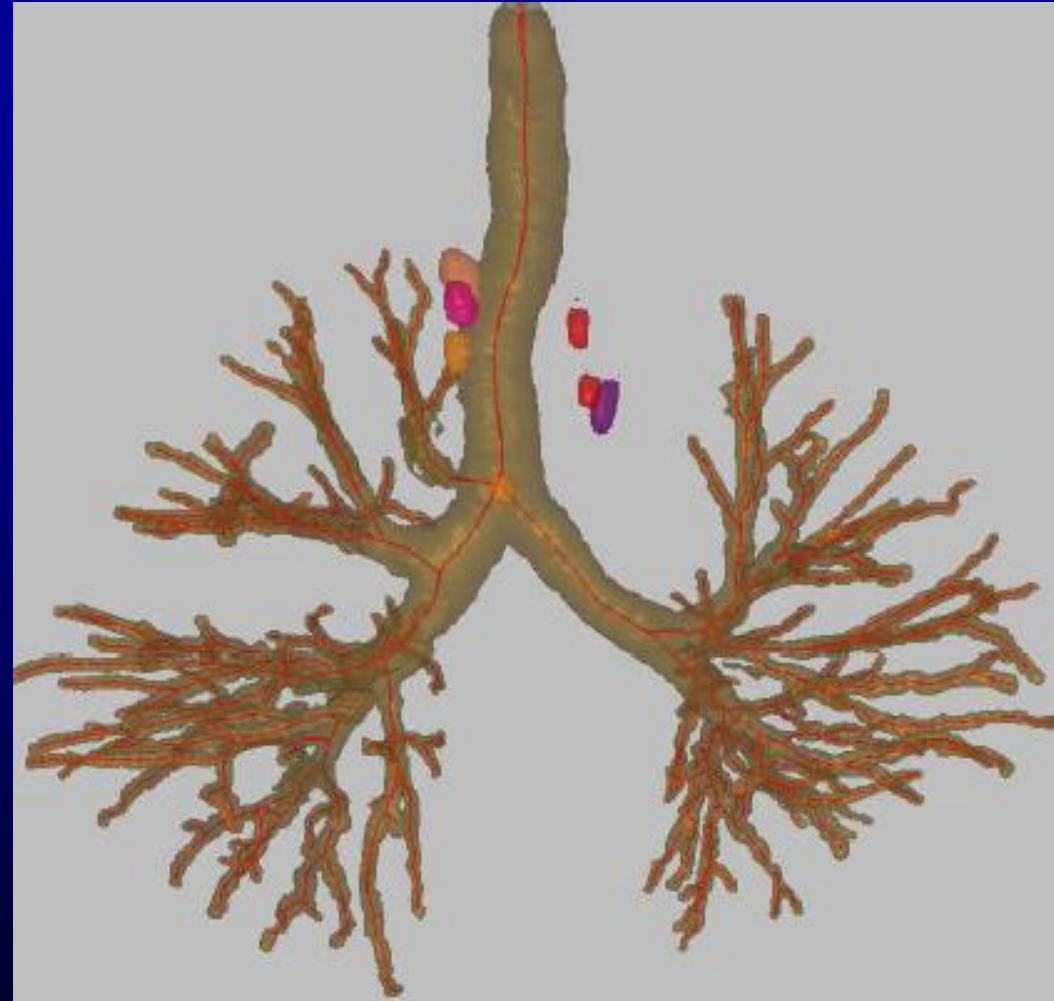
	20349.3.15		21405.67	
	$O_1$	$O_2$	$O_1$	$O_2$
Accuracy	83%	80%	78%	79%
Reproducibility	87%	88%	89%	85%
Processing Time	16s	20s	19s	22s

→ Operator Independent

# Example Lymph Node Segmentations



21405.64



20349.3.3

\* Segmentable nodes: both observers and both methods

# Summary

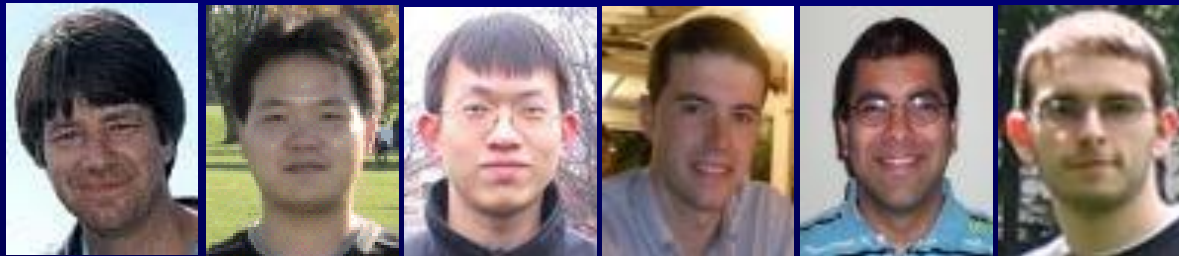
## Single-Section and Single-Click Live Wire

- Reduce human interaction
- Efficient and reliable
- Operator independent
- Can handle typical lymph nodes and other ROIs
  - 90%          Single-Section
  - 80%          Single-Click

# Acknowledgments

- NIH NCI grant #R01-CA074325
- Pinyo Taeprasartsit - experiment participant

The Multidimensional Imaging Processing Lab at Penn State





**Thanks!**

# Single-Section and Single-Click LW Evaluation

Lymph node #: 50

Succ. Seg. :  $M_1$ -45 (90%)  $M_2$ -40 (80%)

Accuracy: >80%

Inter-trial Reprod.: >85%

Lymph Node	Processing Time (Second)				Axis Length (mm)		Volume ( $mm^3$ )	# of Voxel in Lymph Node
	$M_1, \mathcal{E}_1$	$M_1, \mathcal{E}_2$	$M_2, \mathcal{E}_1$	$M_2, \mathcal{E}_2$	Short	Long		
$\mu$	15	17	19	20	5.8	10.5	255.6	1089
$\sigma$	3.9	4.5	5.1	5.1	1.5	4.0	209.6	861.1
Min	7	9	7	9	3.7	5.5	63	273
Max	28	30	29	35	10.7	20.1	970	3796

	$a(M_1, \mathcal{E}_1)$	$a(M_1, \mathcal{E}_2)$	$r(M_1, \mathcal{E}_1, \mathcal{E}_2)$	$a(M_2, \mathcal{E}_1)$	$a(M_2, \mathcal{E}_2)$	$r(M_2, \mathcal{E}_1, \mathcal{E}_2)$
$\mu$	81	81	88	79	79	86
$\sigma$	6.8	6.5	6.7	7.6	8.2	8.5
Min	62	69	66	63	54	60
Max	95	97	98	93	92	99

$M_1, M_2$ : Single-section and single-click LW;  $\mathcal{E}_1, \mathcal{E}_2$ : Trials;  $a(M_i, \mathcal{E}_j)$ : accuracy;  $r(M_i, \mathcal{E}_j, \mathcal{E}_k)$ : reproducibility

# Single-Section and Single-Click LW Evaluation

	Processing Time (Second)			
	$\mathcal{M}_1, \mathcal{E}_1$	$\mathcal{M}_1, \mathcal{E}_2$	$\mathcal{M}_2, \mathcal{E}_1$	$\mathcal{M}_2, \mathcal{E}_2$
$\mu_{\text{Scan3}, \mathcal{O}_1}$	15	14	17	19
$\mu_{\text{Scan3}, \mathcal{O}_2}$	16	19	20	21
$\mu_{\text{Scan5}, \mathcal{O}_1}$	19	20	19	20
$\mu_{\text{Scan5}, \mathcal{O}_2}$	15	18	26	27
Overall	15	17	19	20

	Accuracy and Reproducibility (%)					
	$a(\mathcal{M}_1, \mathcal{E}_1)$	$a(\mathcal{M}_1, \mathcal{E}_2)$	$r(\mathcal{M}_1, \mathcal{E}_1, \mathcal{E}_2)$	$a(\mathcal{M}_2, \mathcal{E}_1)$	$a(\mathcal{M}_2, \mathcal{E}_2)$	$r(\mathcal{M}_2, \mathcal{E}_1, \mathcal{E}_2)$
$\mu_{\text{Scan3}, \mathcal{O}_1}$	84.3	85.5	88.5	81.2	80.5	85.6
$\mu_{\text{Scan3}, \mathcal{O}_2}$	78.3	78.5	82.9	79.5	78.2	86.2
$\mu_{\text{Scan5}, \mathcal{O}_1}$	78.8	79.1	91.8	76.3	77.1	85.6
$\mu_{\text{Scan5}, \mathcal{O}_2}$	80.9	81.1	89.8	77.1	80.1	85.2
Overall	80.7	81.2	88.3	79.2	79.3	85.6

$M_1, M_2$ : Single-section and single-click LW;  $\varepsilon_1, \varepsilon_2$ : Trials;  $a(M_i, \varepsilon_j)$ : accuracy;  $r(M_i, \varepsilon_j, \varepsilon_k)$ : reproducibility