

Introduction to Image Analysis

Image Segmentation

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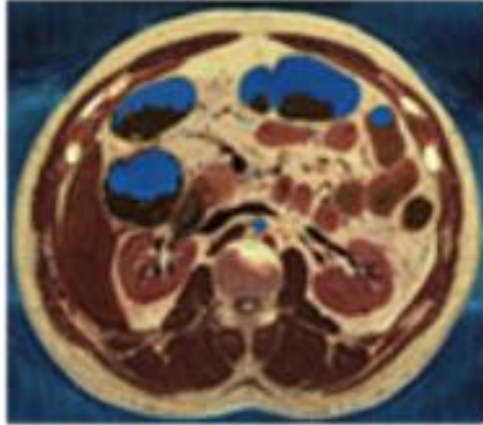
University of Bern

ARTORG Center for Biomedical Engineering Research

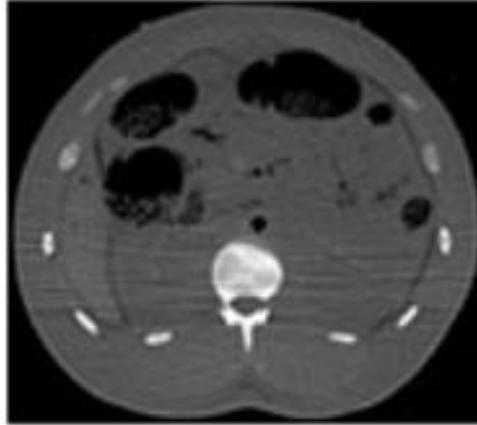
Medical Image Analysis Group

BASICS OF IMAGE SEGMENTATION

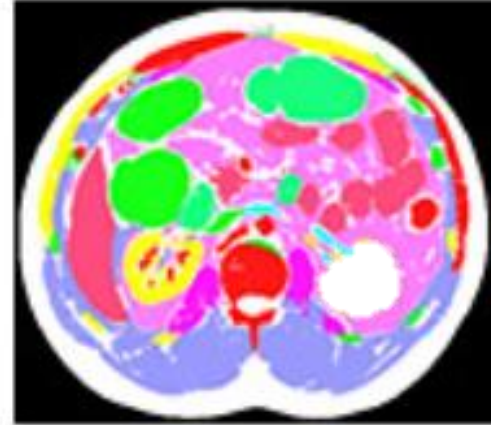
Segmentation – The Goal



Picture



CT Scan

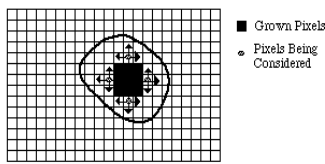
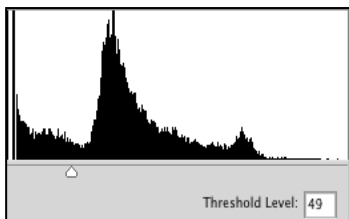


Labels

Visible Human Project

www.nlm.nih.gov/research/visible/visible_human.html

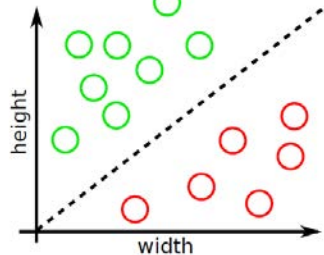
Segmentation Overview – Big Picture



(b) Growing Process After a Few Iterations

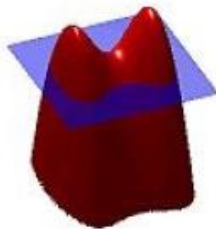
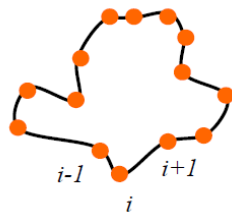
Simple Methods

- > Thresholding
- > Region-Growing
- > ...



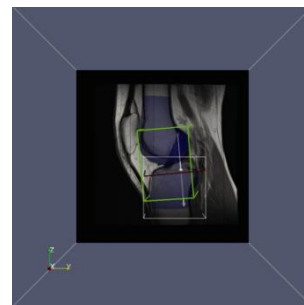
Classification + Clustering

- > kNN
- > SVM
- > ...



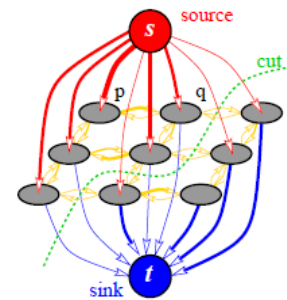
Deformable Models

- > Snakes
- > Level-Sets
- > ...



Active Models

- > ASM
- > AAM

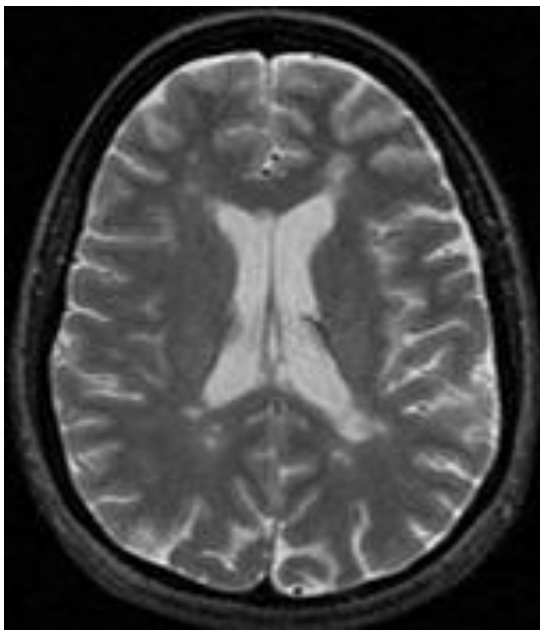


Random Fields & Graph-Cuts

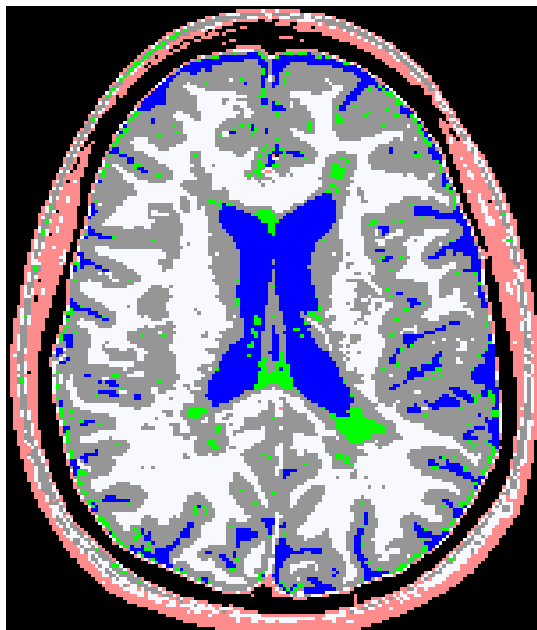
- > MRF
- > CRF

Classification of Voxels

- > Intuitively, we associate different voxels to different structures, depending on their intensity values



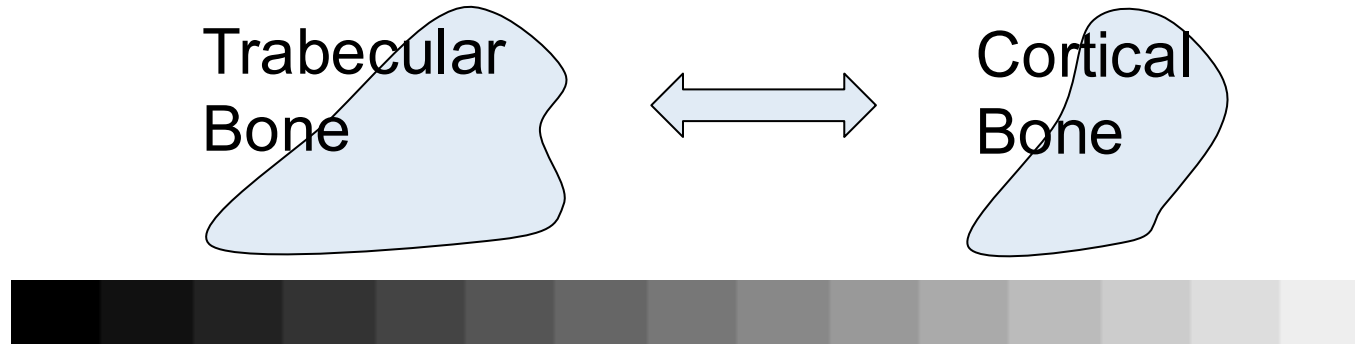
T2w



Segmentation

Classification of voxels

- > Less intuitively, we group voxels trying to maximize the “distance” between the groups



Computationally this can modeled through the **Image Histogram**

Image Histogram

- > Image histogram: representation of the frequency of $P(i)$ intensity values i in the image.

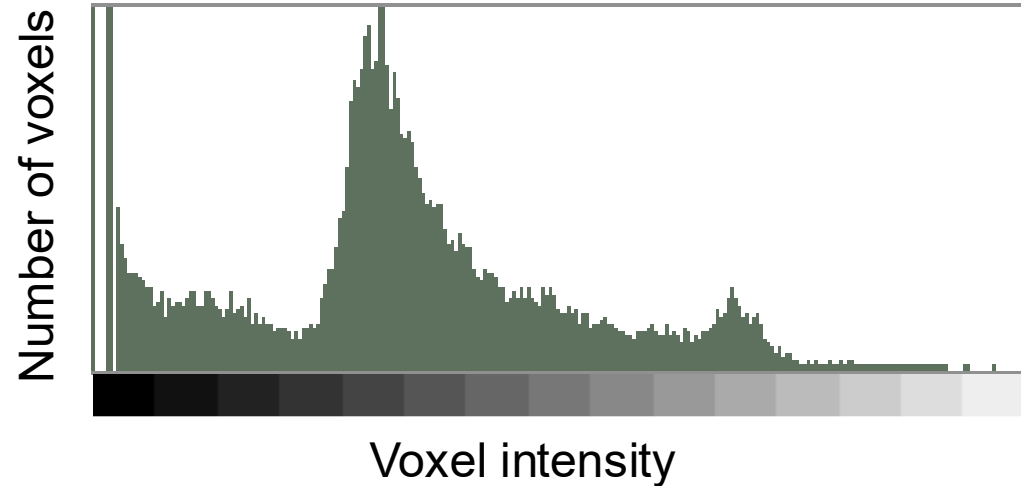
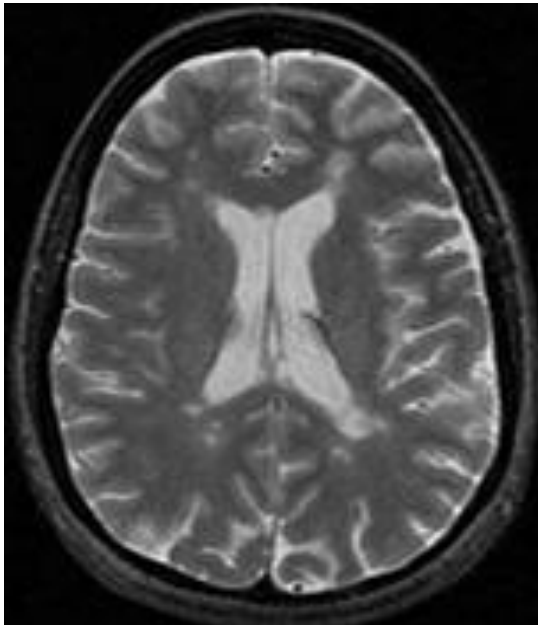
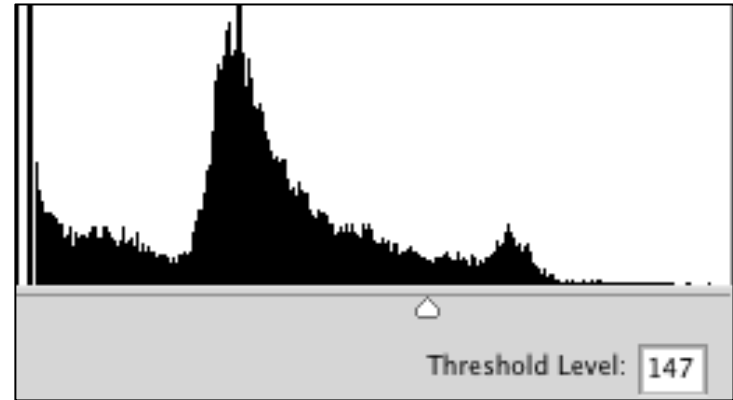
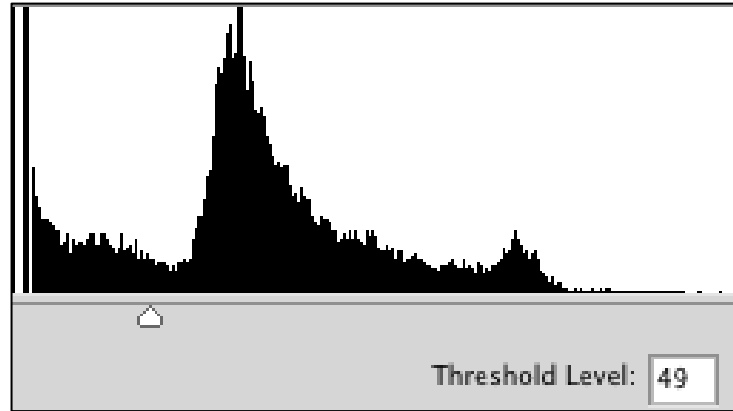
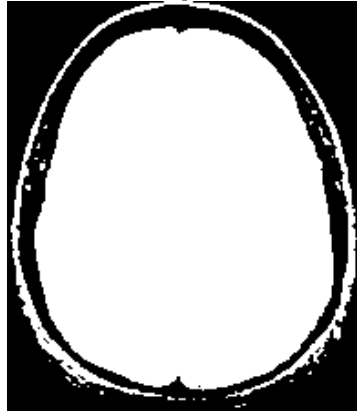
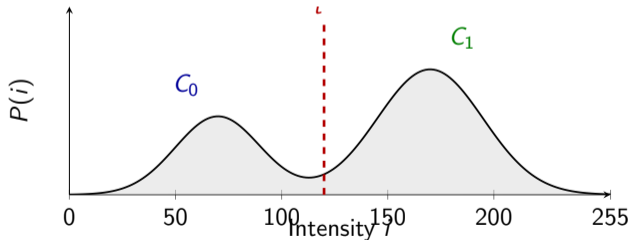


Image Thresholding

- > The very basics of image segmentation
 - **Global**, adaptive/local, Optimal



- > Given an image histogram, how to **automatically** find the best threshold for binary segmentation?
- > Manual threshold selection is subjective and not reproducible
- > Otsu (1979): choose t^* that maximizes the **between-class variance** $\sigma_B^2(t)$



- > Let the image have L intensity levels $\{0, 1, \dots, L - 1\}$
- > Normalized histogram: $p_i = n_i/N$, where $N =$ total pixels
- > Choose threshold t to partition pixels into two classes:

$$C_0 = \{0, 1, \dots, t\} \quad \text{and} \quad C_1 = \{t + 1, \dots, L - 1\}$$

Class probabilities:

$$\omega_0(t) = \sum_{i=0}^t p_i, \quad \omega_1(t) = \sum_{i=t+1}^{L-1} p_i = 1 - \omega_0(t)$$

Class means:

$$\mu_0(t) = \frac{1}{\omega_0(t)} \sum_{i=0}^t i \cdot p_i, \quad \mu_1(t) = \frac{1}{\omega_1(t)} \sum_{i=t+1}^{L-1} i \cdot p_i$$

Key insight: Total variance = within-class + between-class variance

$$\sigma_{\text{total}}^2 = \sigma_W^2(t) + \sigma_B^2(t)$$

Within-class variance:

$$\sigma_W^2(t) = \omega_0(t) \sigma_0^2(t) + \omega_1(t) \sigma_1^2(t)$$

Between-class variance:

$$\sigma_B^2(t) = \omega_0(t) \omega_1(t) [\mu_0(t) - \mu_1(t)]^2$$

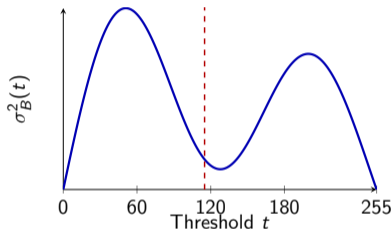
- > Since σ_{total}^2 is constant, minimizing $\sigma_W^2 \iff$ maximizing σ_B^2
- > $\sigma_B^2(t)$ is **much cheaper** to compute (no per-pixel variance needed)

$$t^* = \arg \max_{0 < t < L-1} \sigma_B^2(t)$$

1. Compute the normalized histogram p_i
2. For each candidate threshold t :
 - Compute $\omega_0(t), \omega_1(t)$
 - Compute $\mu_0(t), \mu_1(t)$
 - Compute $\sigma_B^2(t)$
3. Select $t^* = \arg \max_t \sigma_B^2(t)$
4. Segment:

$$g(x, y) = \begin{cases} 1 & \text{if } f(x, y) > t^* \\ 0 & \text{otherwise} \end{cases}$$

Complexity: $O(L)$ – single pass using cumulative sums

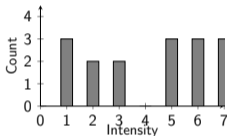
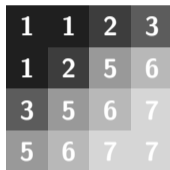


Between-class variance curve

Peak = optimal threshold

Otsu Segmentation – Worked Example

Consider a 4×4 image with $L = 8$ intensity levels:



Evaluating $\sigma_B^2(t)$ for each t :

t	1	2	3	4	5	6
σ_B^2	1.69	3.57	5.14	6.53	4.08	0.95

$\implies t^* = 4$ separates low-intensity pixels $\{1, 2, 3\}$ from high $\{5, 6, 7\}$

Strengths

- > Fully automatic – no parameters
- > Computationally efficient: $O(L)$
- > Optimal for bimodal histograms
- > Equivalent to 1D Fisher's discriminant analysis
- > Widely implemented (OpenCV, scikit-image, MATLAB)

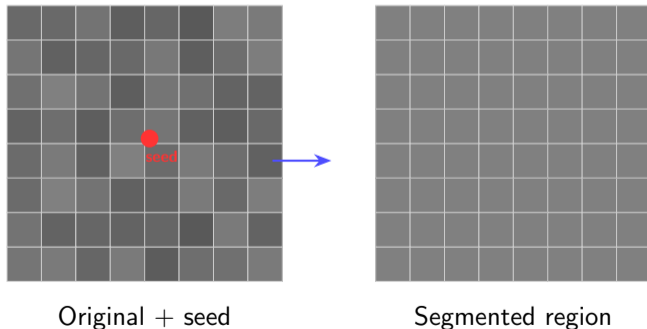
Limitations

- > Assumes bimodal histogram
- > Poor performance with uneven illumination
- > Sensitive to noise
- > No spatial information used
- > Biased toward classes with larger variance

Variations: multi-level Otsu, 2D Otsu, local/adaptive Otsu, Kittler–Illingworth method

Region Growing – Motivation

- > Thresholding uses only **intensity**, ignoring **spatial** relationships
- > We want to group spatially connected pixels that share similar properties
- > Idea: start from **seed points**, grow regions by appending similar neighbors



Let R be the entire image region. A segmentation partitions R into n regions R_1, R_2, \dots, R_n such that:

- (a) $\bigcup_{i=1}^n R_i = R$ (every pixel belongs to a region)
- (b) R_i is a connected region for all i (spatial connectivity)
- (c) $R_i \cap R_j = \emptyset$ for $i \neq j$ (no overlaps)
- (d) $P(R_i) = \text{TRUE}$ for all i (homogeneity predicate)
- (e) $P(R_i \cup R_j) = \text{FALSE}$ for adjacent R_i, R_j (distinct regions)

> $P(\cdot)$ is a **homogeneity predicate**, e.g.:

$$P(R_i) = \text{TRUE} \iff |I(x, y) - \mu_{R_i}| < \tau \quad \forall (x, y) \in R_i$$

1. **Select seed point(s)** s (manual or automatic)
2. Initialize region: $R = \{s\}$
3. For each unvisited neighbor q of R :
 - If similarity criterion is met:
 $|I(q) - \mu_R| < \tau$
 - Add q to R , update μ_R
4. Repeat until no more pixels can be added
5. Choose new seed (if unassigned pixels remain)

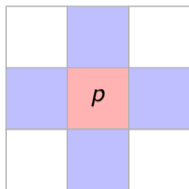
42	48	88
44	47	91
43	45	87
41	44	92
40	45	95

Seed Grown Candidate

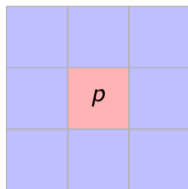
Similarity criteria: intensity difference, gradient magnitude, texture features, color distance, ...

Growing from seed (45) with $\tau = 5$.
Right side (high intensity) is **not** included.

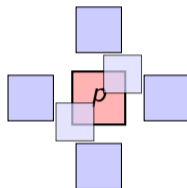
Region Growing – Neighborhood & Connectivity



4-connected



8-connected



6-connected (3D)

- > **4-connected** (2D): shares edge → 4 neighbors
- > **8-connected** (2D): shares edge or corner → 8 neighbors
- > **6/18/26-connected** (3D): for volumetric data
- > Choice of connectivity affects the shape of grown regions

The choice of predicate P is **critical** to the quality of segmentation.

Common criteria:

- > **Intensity difference from seed:**

$$|I(q) - I(s)| < \tau$$

- > **Intensity difference from region mean** (adaptive):

$$|I(q) - \mu_R| < \tau, \quad \mu_R = \frac{1}{|R|} \sum_{p \in R} I(p)$$

- > **Statistical criterion:**

$$|I(q) - \mu_R| < k \cdot \sigma_R$$

where σ_R is the standard deviation of intensities in R

- > **Multi-feature:** combine intensity, gradient, texture

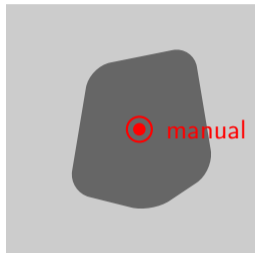
$$d(\mathbf{f}(q), \mathbf{f}_R) < \tau, \quad \mathbf{f} = (I, |\nabla I|, \text{texture}, \dots)^\top$$

Region Growing – Seed Selection Strategies

Seed selection significantly influences results.

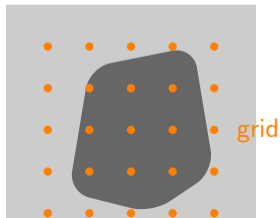
Manual:

- > User clicks on region of interest
- > Common in medical imaging (e.g., 3D Slicer)



Automatic:

- > Grid-based: seeds at regular spacing
- > Histogram-based: select pixels near histogram peaks
- > Edge-based: seeds placed away from edges
- > Clustering: use k-means centroids as seeds



Region Growing – Step-by-Step Illustration

Iteration 1

52	48	45	43	80	85
50	46	44	42	82	88
51	47	43	44	79	90
53	49	45	46	83	87
55	51	48	47	81	92
54	52	50	49	84	91

Seed at pixel with
intensity 43, $\tau = 10$

Iteration 3

52	48	45	43	80	85
50	46	44	42	82	88
51	47	43	44	79	90
53	49	45	46	83	87
55	51	48	47	81	92
54	52	50	49	84	91

Region grows to
similar neighbors

Converged

52	48	45	43	80	85
50	46	44	42	82	88
51	47	43	44	79	90
53	49	45	46	83	87
55	51	48	47	81	92
54	52	50	49	84	91

Growth stops at
intensity boundary

Advantages

- > Simple concept, easy to implement
- > Uses **spatial** information (connected regions)
- > Can segment objects with irregular shapes
- > Can handle multiple criteria simultaneously
- > Results align well with perceived edges

Disadvantages

- > Sensitive to **seed selection**
- > Sensitive to **noise**
- > Order-dependent: first region “dominates”
- > Threshold τ must be chosen carefully
- > **No global view**: local method
- > Can leak through weak boundaries

Seeded Region Growing (SRG)

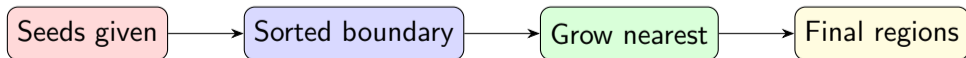
Adams & Bischof, 1994

- > Seeds provided as input
- > Pixels assigned to nearest region based on intensity difference from region mean
- > Ordering: pixel with **smallest** $\delta(q) = |I(q) - \mu_{R_i}|$ is assigned first
- > Deterministic given seeds

Unseeded Region Growing (URG)

Lin et al., 2000

- > No explicit seed selection needed
- > Start with single pixel, grow if $S(p, q) > T$
- > If a pixel cannot be added, it becomes a new seed for another region
- > Avoids seed selection bias
- > Less user interaction required



SRG

Variation: Region Splitting and Merging

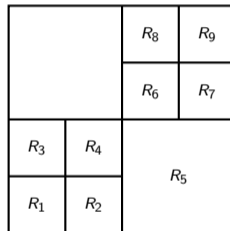
Opposite approach to region growing:

Start with entire image as one region and **split** non-homogeneous parts.

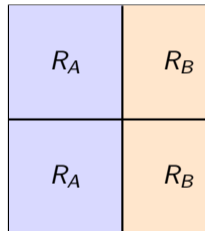
1. If $P(R_i) = \text{FALSE}$, split R_i into 4 quadrants
2. Repeat recursively until all regions satisfy P
3. **Merge** adjacent regions R_i, R_j if $P(R_i \cup R_j) = \text{TRUE}$
4. Stop when no further splits or merges possible

Uses a **quadtree** data structure.

Split



merge
→

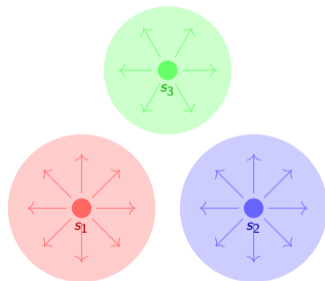


Variation: Simultaneous Region Growing

- > **Problem:** Sequential growing biases toward the first region
- > **Solution:** Grow all regions **simultaneously**

Approach:

- > Multiple seeds placed at once
- > Maintain a **priority queue** of boundary pixels sorted by similarity
- > At each step, assign the globally most similar pixel to its closest region
- > No single region dominates growth
- > Efficient on parallel architectures



Competitive growth

Resolves ambiguities at region boundaries more fairly than sequential methods.

Variation: Adaptive Threshold Region Growing

- > Fixed threshold τ may be too restrictive or too permissive in different parts of the image
- > **Adaptive approach:** update τ as the region grows

Example: Statistical adaptive criterion

$$|I(q) - \mu_R^{(n)}| < k \cdot \sigma_R^{(n)}$$

where $\mu_R^{(n)}$ and $\sigma_R^{(n)}$ are updated after n pixels are added.

Gradient-based adaptation:

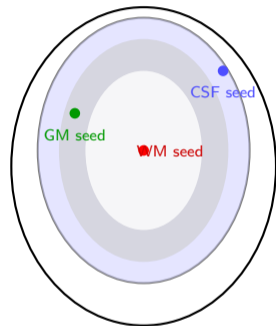
- > Reduce τ near strong gradients (to avoid leaking)
- > Increase τ in smooth homogeneous areas

$$\tau(q) = \tau_0 \cdot \exp\left(-\frac{|\nabla I(q)|^2}{2\beta^2}\right)$$

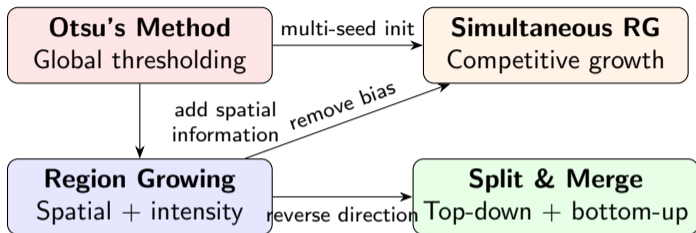
This prevents region leakage through weak boundaries – a common failure mode of basic region growing.

Region growing is widely used in medical image segmentation:

- > **Brain MRI:** segment white matter, gray matter, CSF from T1/T2-weighted images
- > **Vascular segmentation:** trace vessel trees in CT/MR angiography (branch-based strategies)
- > **Tumor segmentation:** delineate lesion boundaries in CT/MRI
- > **Bone segmentation:** separate cortical and trabecular bone in CT
- > **Lung segmentation:** extract airways/parenchyma from chest CT



Brain tissue segmentation

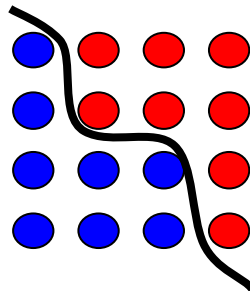


- > **Otsu:** automatic, fast, but no spatial info – good for bimodal images
- > **Region Growing:** incorporates spatial connectivity, flexible criteria, but sensitive to seeds and noise
- > **Variations** (SRG, URG, split&merge, adaptive) address key limitations
- > These methods form the **foundation** for more advanced approaches (deformable models, graph cuts, deep learning)

Live-wire segmentation

Barret et al. Media 1997

- > Motivation: Lack of robustness with automatic methods, time-consuming manual segmentation
- > Works on 2D (i.e. finds contours)
- > Finds the optimal path between starting and ending point (seen as a graph searching problem)
- > Optimal path = **minimal** cost between points = cumulative sum between local segments



Live-wire segmentation

- > Optimal path = **minimal** cost between nodes (pixels)
- > Global cost = cumulative sum between local segments
Local cost $l(p, q)$

$$l(p, q) = \omega_G \cdot f_G(q) + \omega_Z \cdot f_Z(q) + \omega_D \cdot f_D(p, q)$$

⇒ f_G : Gradient information $f_G = 1 - \frac{G}{\max(G)}$

⇒ f_Z : Laplacian (L) information (2nd order derivatives)
 $f_Z = 0$ if $L(I) = 0$ or neighbor(p) shows change of sign
 $f_Z = 1$ otherwise

⇒ f_D : directional derivative

Live-wire segmentation

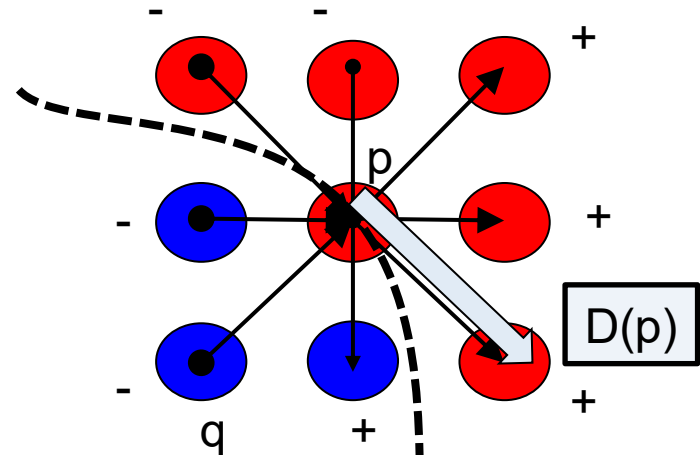
- > Optimal path = **minimal** cost between nodes (pixels)
- > Global cost = cumulative sum between local segments

→ f_d : directional derivative

$$f_D(p, q) = \frac{2}{3\pi} \{ \cos[d_p(p, q)]^{-1} + \cos[d_q(p, q)]^{-1} \}$$

$$d_p(p, q) = D(p) \cdot L(p, q) \quad D(p): \text{normal to gradient direction at point } p$$
$$d_q(p, q) = L(p, q) \cdot D(q)$$

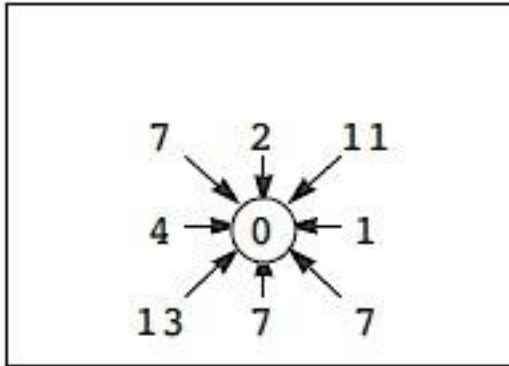
$$L(p, q) = \begin{cases} q - p & \text{if } D(p) \cdot (q - p) \geq 0 \\ p - q & \text{if } D(p) \cdot (q - p) < 0 \end{cases}$$



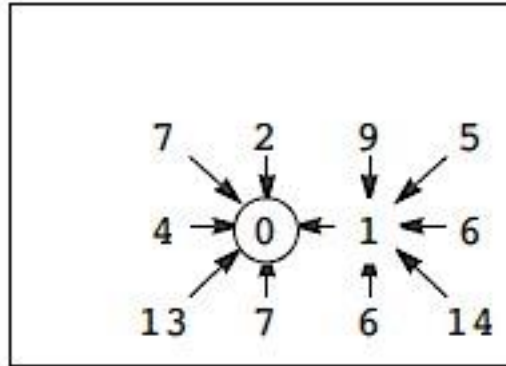
“pq vectors point in the direction of D minimizing the angle between them”

Live-wire segmentation

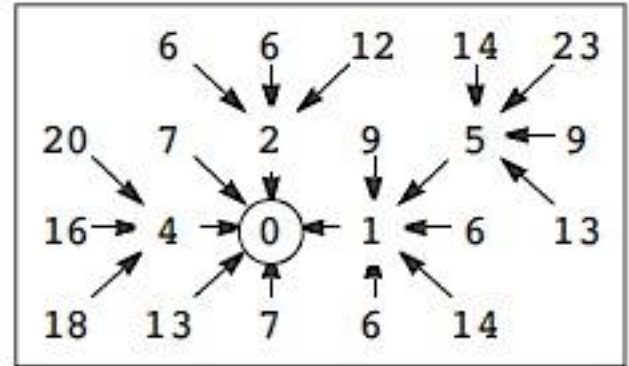
- > Graph searching initialized by user “seed” (i.e. mouse click)
- > Seed is expanded by computing local costs
- > Minimal cost is used as next seed



Initial seed
(one expansion)

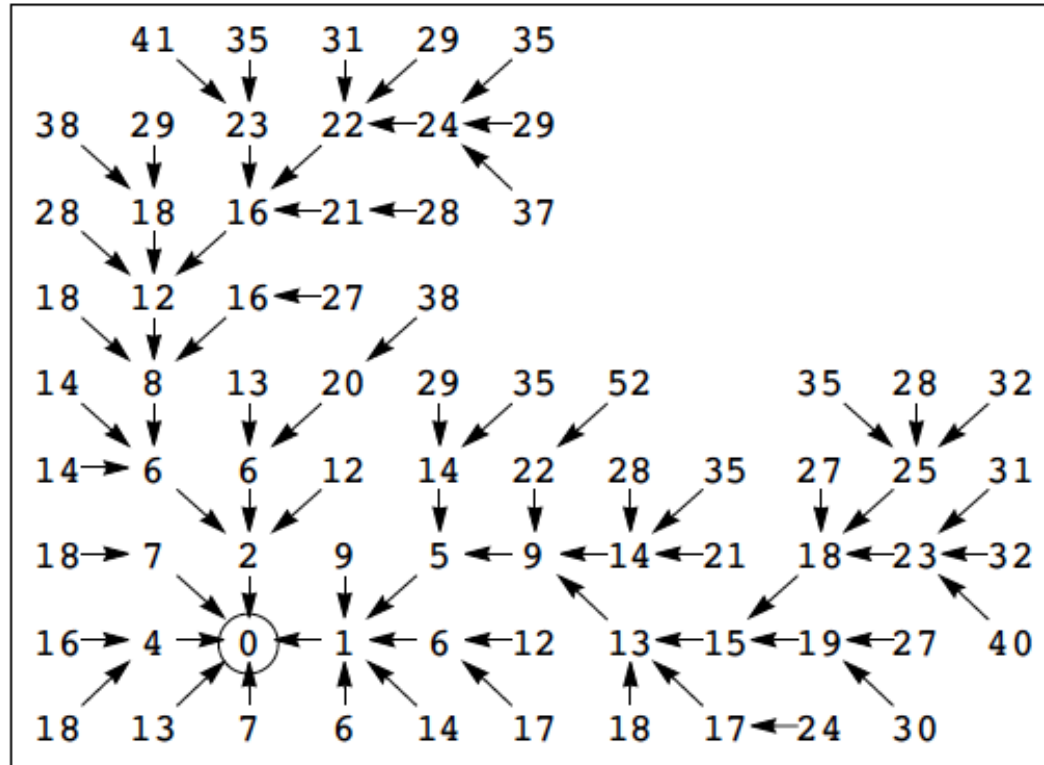


Two expansions



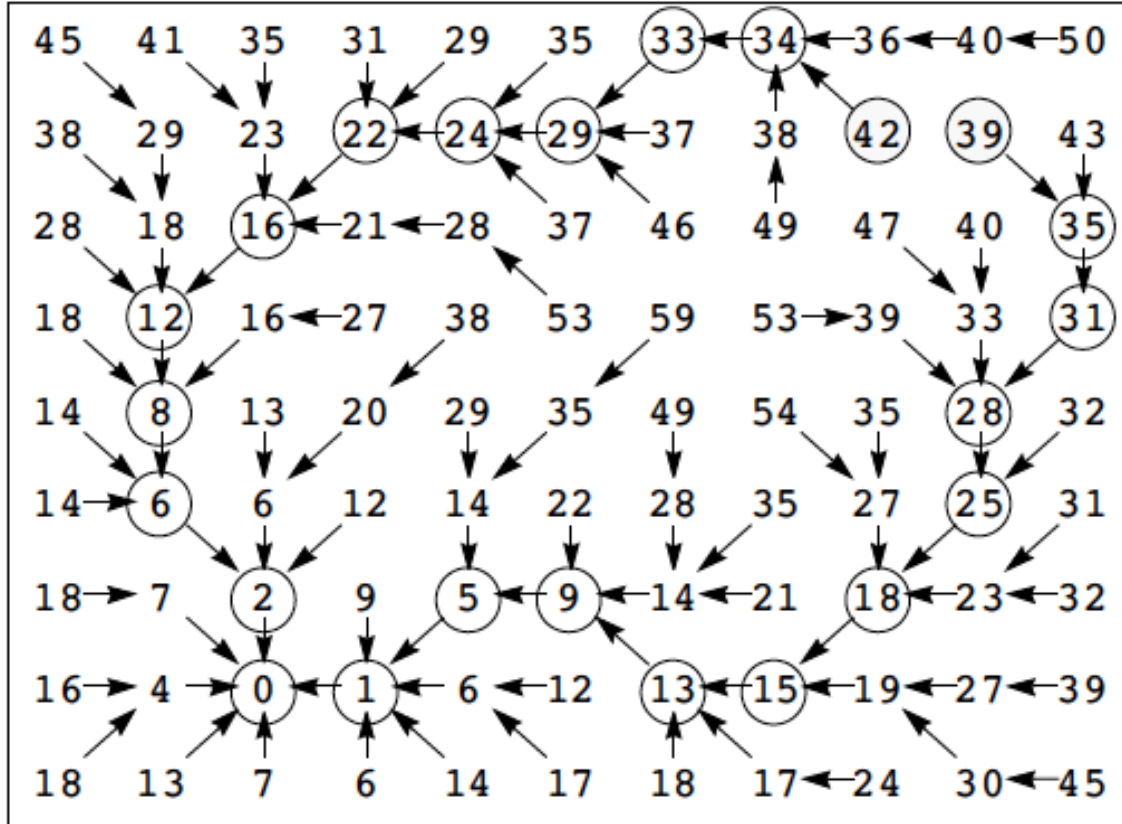
Five expansions

Live-wire segmentation



47 points expanded

Live-wire segmentation



Completed total cost, with total costs of 42 and 39

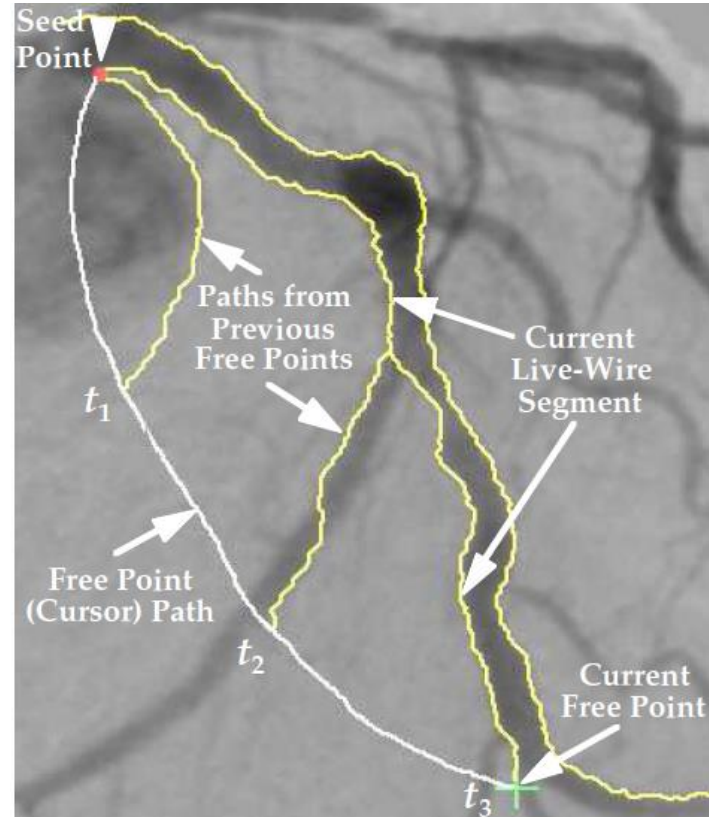
Live-wire segmentation

- > Advantages of live-wire segmentation:
 - Monitor segmentation
 - Correct on the fly
 - No fix-path in advance
 - Speed over other path searching approaches. Enables real time

Live-wire segmentation

> Snap and wrap.

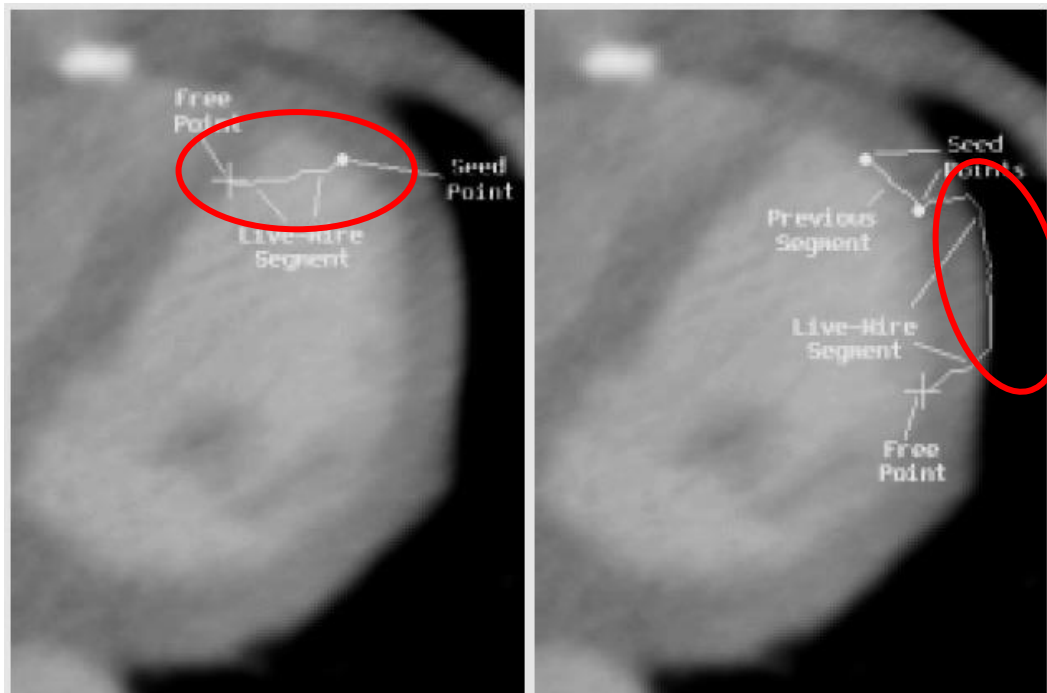
Dynamically changing “free points” elicit a new minimal cost path



Live-wire segmentation

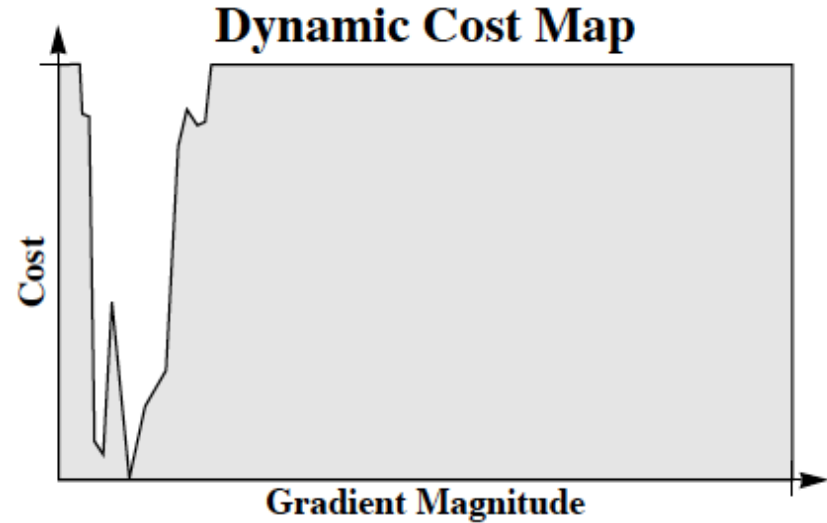
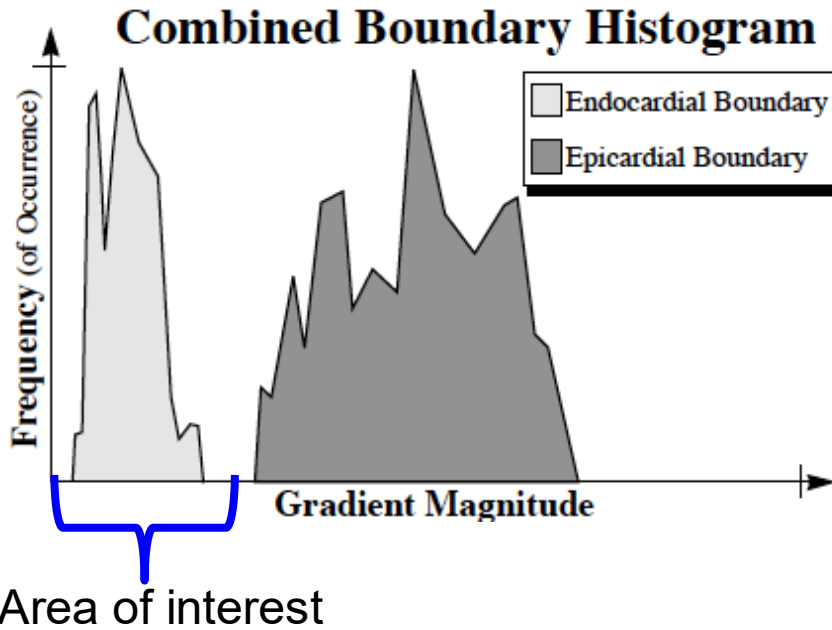
> On the fly training

Goal is to train the cost function to **fit to desired edges** rather than strongest one



Live-wire segmentation

> On the fly training

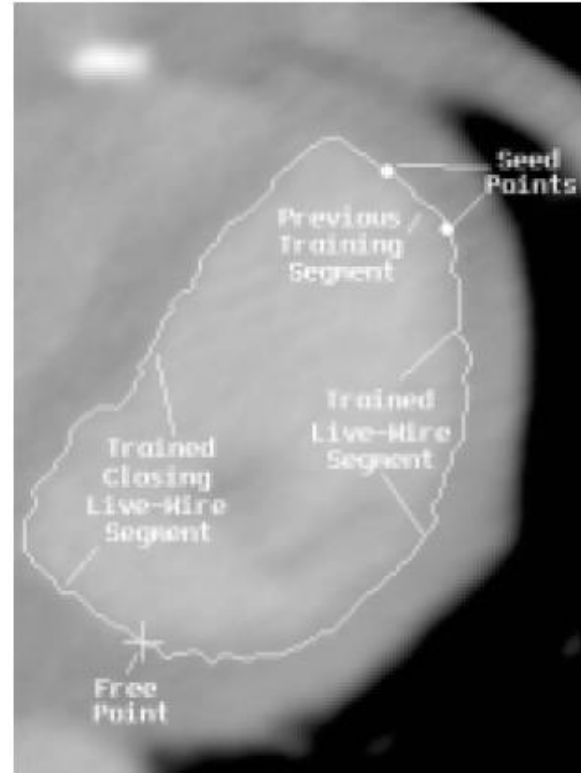


Associated gradient cost
Derived from current segment

Live-wire segmentation

> On the fly training

Resulting segmentation with
on the fly training



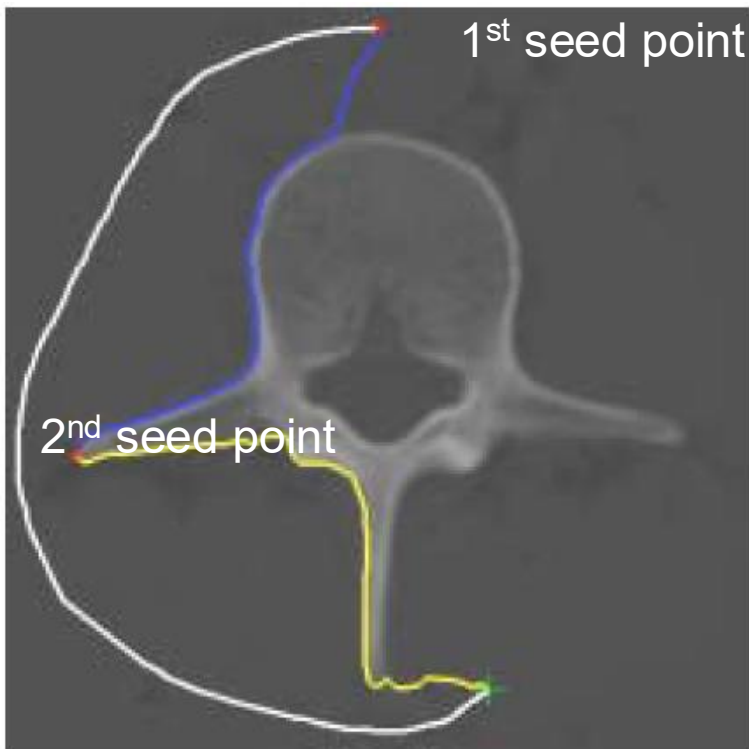
Live-wire segmentation

> Data-driven path cooling

Cooling: path between two fixed points (manual clicks)

Manual = ☹ => **data-driven** cooling!

Data-driven = the **longer** the segment remains “still” the more confidence it has, the cooler it becomes

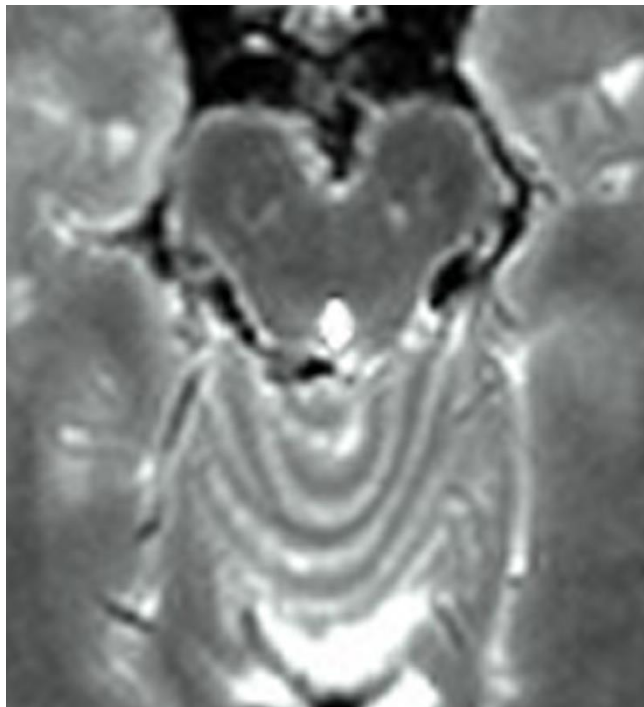


Evaluating segmentation

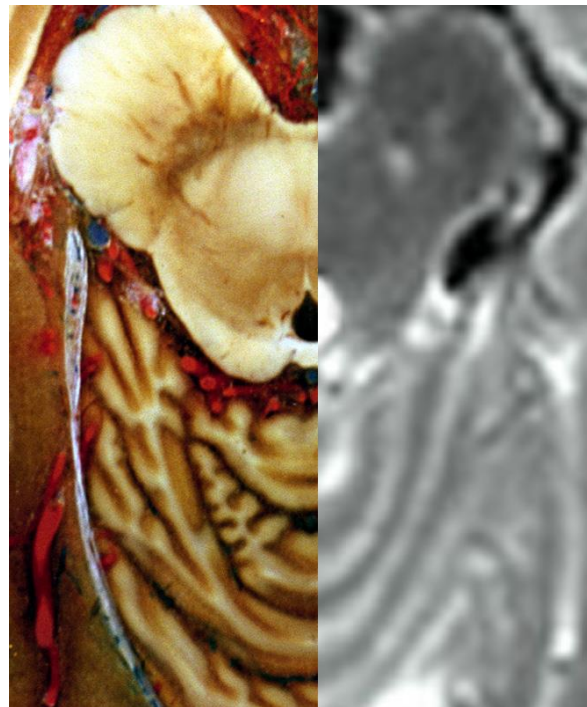
Validation of Image Segmentation

- > Digital phantoms.
 - Ground truth known accurately.
 - Not so realistic.
- > Acquisitions and careful segmentation.
 - Some uncertainty in ground truth.
 - More realistic.
- > Autopsy/histopathology.
 - Addresses pathology directly; resolution
- > Clinical data ?
 - Hard to know ground truth.
 - Most realistic model.

Comparison To Higher Resolution



MRI

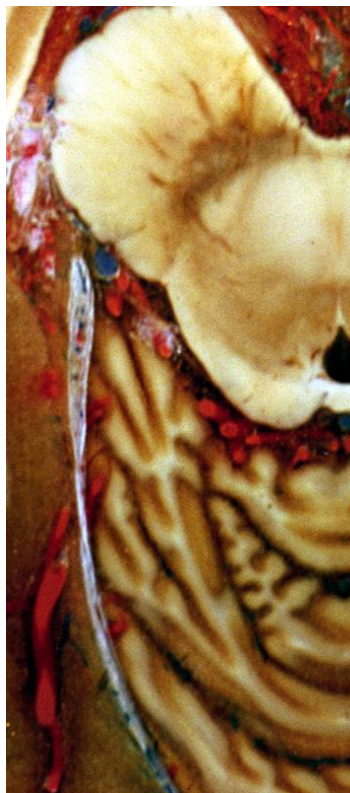


Photograph

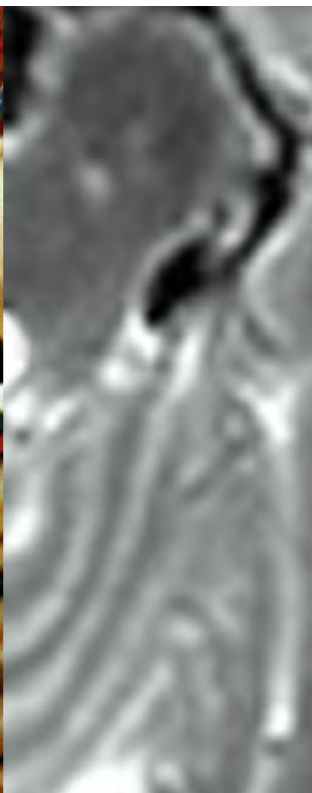
MRI

Source: S. Warfield

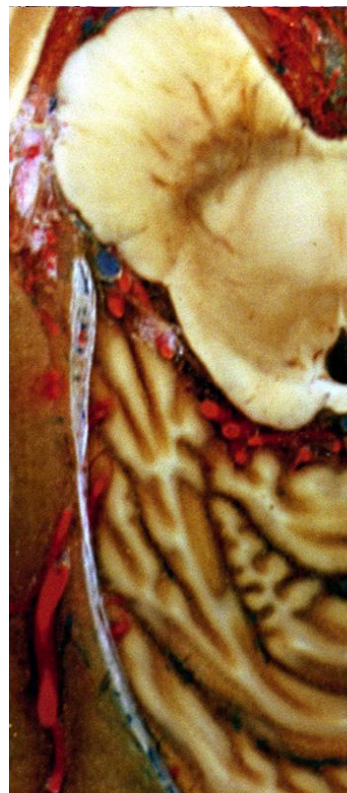
Comparison To Higher Resolution



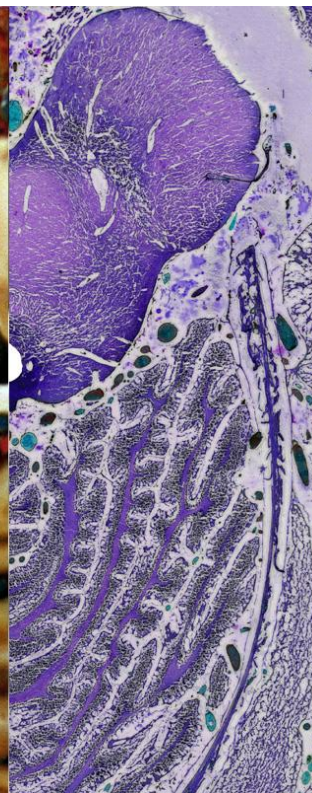
Photograph



MRI



Photograph



Microscopy

Measures of Expert Performance

- > Repeated measures of volume
 - Intra-class correlation coefficient
- > Spatial overlap
 - Jaccard coefficient: Area of intersection over union.
 - Dice coefficient: increased weight of intersection.
 - Vote counting: majority rule, etc.
- > Boundary measures
 - Hausdorff, 95% Hausdorff.
- > Bland-Altman methodology:
 - Requires a reference standard.
- > Measures of correct classification rate:
 - Sensitivity, specificity ($\Pr(D=1|T=1)$, $\Pr(D=0|T=0)$)
 - Positive predictive value and negative predictive value (posterior probabilities $\Pr(T=1|D=1)$, $\Pr(T=0|D=0)$)

Measures of Expert Performance

- > Measures of correct classification rate:
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 - Positive predictive value and negative predictive value (posterior probabilities $\Pr(T=1|D=1)$, $\Pr(T=0|D=0)$)

$$\text{sensitivity} = \frac{\text{number of true positives}}{\text{number of true positives} + \text{number of false negatives}}$$

true positive (TP)

eqv. with hit

true negative (TN)

eqv. with correct rejection

false positive (FP)

eqv. with false alarm, Type I error

false negative (FN)

eqv. with miss, Type II error

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