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Abstract

selecting the parameters (settings") for biological image analysis systems using the common notions of coverage and complexity. This is essential for broadening the scope of applicability of these algorithms. The problem of automated parameter selection is formulated as a variational optimization based on mathematical descriptions of segmentation conciseness ad coverage, a probabilistic vesselness measure, scale, and the Minimum Description Length (MDL) principle. Of specific interest to us is the analysis of tube-like biological objects such as tumor micro-vasculature and neurons. Next, the efficient Recursive Random Search (RRS) global optimization algorithm is utilized to explore the parameter space in an effic ient and parallel manner. As an example, the resulting parameter set after 1000 RRS steps for all nine parameters of our automated tracing algorithm produces segmentation results that are within 5% of the globally optimal result, as compared to more than 300,000 steps that would be needed by an exhaustive search. Several examples representing progressively more complex search spaces are provided to demonstrate the power and broad applicability of this method. The parameter sets that produce optimal segmentation across different biological objects and imaging modalities, using different segmentation algorithms are presented to demonstrate the generality of this approach. Instead of manually hand-tuning empirical parameters that still inevitably contain human subjectivity, users can now objectively obtain optimal segmentation result automatically, across a broad domain of applications.

Technical Approach

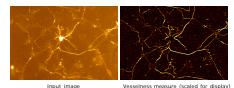
The variational formulation for image segmentation is expressed mathematically using the Minimum Description Length (MDL) formulation [1]. This is analogous to the problem of fitting a polynomial to a set of points, where the goal is to find a trade-off between the degree of the polynomial (conciseness) and the corresponding fitting error (coverage).

Let an image segmentation algorithm be a function $f:\{L?\}\to M$ that maps the object(s) imaged in the image I to the segmented object(s) M using a parameter vector $?\in \Omega$ where Ω is the space containing all possible parameter vectors. Now, the goal becomes to obtain the optimal M , called \hat{M} , in terms of conciseness and coverage

$$L_{2-p}(I|M) = \frac{1}{1} \log_{\frac{p}{2}} \underbrace{P}_{\text{coverage}}^{\{I\}} \underbrace{AM}_{\text{pl}}) + \frac{k}{24243} \log_{2} n$$
conciseness

 $-\log_2 P(I|M)@\mathbf{m}$ $\mathbf{x} = \{\mathbf{x} \mid \mathbf{x} \in M^C, P(\mathbf{x}) \ge \mathbf{a}\}$

Since the likelihood term is not defined by the MDL principle, we use the probabilistic Vesselness measure [2] for tube-like objects.



Then, the image is segmented using a particular set of parameters, and the resulting segmented image is evaluated using the MDL-based m

An efficient global optimization algorithm, the Recursive Random Search [3] is used to explore the parameter space using the metric as defined using the

Conclusions

The presented approach removes the human subjectivity factor when selecting parameters/settings for an image analysis system. It also reveals the practical limitations of a particular image analysis system, especially when the system is

The broad applicability of this method is demonstrated by the experimental results. Essentially, all the parameters of an image analysis system are mapped onto user -friendlier terms.

Automated Parameter Selection for Optimal Biological Image Segmentation



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Input

Optimal:

Evhaustivo

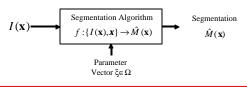
different

application:

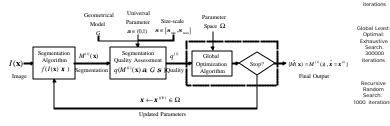
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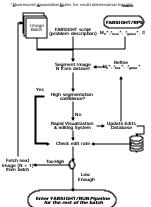


The Traditional Approach to Image Segmentation



The Automated Parameter Selection Approach to Image Segmentation





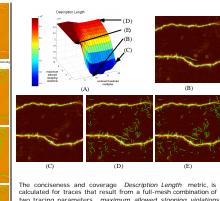
The FARSIGHT* Framework

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neurons from noisy confocal microscope images. Information Technology in Biomedicine, IEEE Transactions on, 2003, 7 (4): p. 302-317. [14] Abdul-Karim, M.A., et al., Automated tracing and change analysis of angiogenic vasculature from in vivo multiphoton confocal image time series, Microvasc Res, 2003. 66 (2): p. 113-25.

Experimental Results: Automated Tracing Algorithm [13,14]



two tracing parameters. maximum allowed stopping violations and contrast threshold multiplier. The traces are shown in (B), (C), (D), and (E), and the corresponding metric value is pointed in (A), where (B) is the optimal trace. (Image courtesy of Andreas Jeromin, Baylor College of Medicine, Texas)

Recursive Random Search

150 200

50

Global Least- Optimal

Experimental Results: Simple Intensity Thresholding Description Length vs. Intensity Threshold 150,000

Global Optimal

Input Image

Intensity Threshold Description Length vs. Intensity Threshold 400,000 300,000 200,000 100,000 50 100 150 200 250

Illustrates the segmentation result using a global intensity threshold value for tube-like images: upper row displays the result for a neuron image (source: Nataliae lowell, Wadsworth Center, New York), and the lower row displays the result for a retinal blood wessels image (source: The STAKE Project). The left-most column contains the input image, the middle column contains the optimal segmentation image, and the right-most column shows plots of the description length (the conciseness and coverage metric) versus the intensity threshold value.

parameters is analogous to the variational formulation for image References

State of the Art

Metrics for segmentation evaluation [4-10] can be either goal-oriented, i.e. evaluation based on the performance of post-segmentation steps such as pattern classification [9], based on other application-defined criteria, such as the probability of false detection [6, 7, 10], or based on visual inspections [4]. Here, we restrict ourselves to the latter category, since we believe that optimality of the seamentation result should be compatible with and independent

The Minimum Description Length (MDL) principle [1] is chosen since it offers a systematic way to obtain an objective balance between

segmentation conciseness and coverage. We present a different MDL-based formulation, i.e. compared to [11-12], where the

formulation now contains a term to account for the probability of

This search of optimal

of any post-segmentation image analysis steps.

structural presence at each pixel.

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38-54.
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