



## Cell Segmentation and Tracking in Phase Contrast Images using Graph Cut with Asymmetric Boundary Costs

Robert Bensch and Olaf Ronneberger

Computer Science Department and BIOSS Centre for Biological Signalling Studies, University of Freiburg, Germany



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- UNI FREIBURG Introduction •
  - Method •
    - Segmentation
    - Tracking
  - Experiments
  - Conclusion •



## Phase contrast microscopy





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(B) Phase-contrast 50 μm

Phase-contrast

Figure: B. Alberts et al., Molecular Biology of the Cell, 4th Edition, 2002.

#### Visualize transparent objects with high contrast at cell borders

### Phase contrast microscopy





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Shade-off

Halo pattern

Strong edges inside and outside the cell

#### Drawback: Various artifacts

# Standard segmentation algorithms



Cyan: Graph cut segmentation result Yellow: Our manual ground truth

- Standard edge-based segmentation algorithms fail
- Traditional graph cut with symmetric boundary costs.

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UNI FREIBURG Our approach



#### UNI FREIBURG True cell borders appear as **dark-to-bright** transition (positive phase contrast microscopy)





Yellow: Cell outwards direction Green: True cell border Red: Wrong cell border

Our approach



True cell borders appear as dark-to-bright transition (positive phase contrast microscopy)





Yellow: Cell outwards direction Green: True cell border Red: Wrong cell border

- Search for segmentation mask that favors dark-tobright transitions at its boundary
- Graph cut with asymmetric boundary costs

#### **Related work**



- UNI FREIBURG Kanade et al.: Two-step reconstruction approach
  - Reconstruct abs. phase image & apply basic threshold techniques
  - Fails if sample contains light absorbing structures
  - <u>Ambühl et al.</u>: Morphological image processing and level sets
    - Handle halo artifacts by changing image during level set evolution
  - Magnusson et al.: Winner ISBI Cell Tracking Challenge 2014
    - Strong tracking approach & Segmentation based on bandpass filtering, thresholding and watershed transform

- (1) K. Li and T. Kanade, "Nonnegative mixed-norm preconditioning for microscopy image segmentation," Proceedings of IPMI, pp. 362–373, 2009.
- (2) M.E. Ambül, C. Brepsant, J.-J. Meister, A.B. Verkhovsky, and I.F. Sbalzarini, "High-resolution cell outline segmentation and tracking from phase-contrast microscopy images," JOM, vol. 245, no.2, pp. 161–170, 2012.
- (3) K. Magnusson, J. Jaldén, and H. M. Blau, Cell tracking using bandpass filtering and the viterbi algorithm, Description of the algorithm available at: http://www.codesolorzano.com/celltrackingchallenge/

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    - Strong tracking approach & Segmentation based on bandpass filtering, thresholding and watershed transform
  - <u>Boykov et al.</u>: Asymmetric boundary costs in min-cut
    - Propose asymmetric boundary costs for segmentation

#### $\rightarrow$ Never been applied to phase contrast microscopy

- (1) K. Li and T. Kanade, "Nonnegative mixed-norm preconditioning for microscopy image segmentation," Proceedings of IPMI, pp. 362–373, 2009.
- (2) M.E. Ambül, C. Brepsant, J.-J. Meister, A.B. Verkhovsky, and I.F. Sbalzarini, "High-resolution cell outline segmentation and tracking from phase-contrast microscopy images," JOM, vol. 245, no.2, pp. 161–170, 2012.
- (3) K. Magnusson, J. Jaldén, and H. M. Blau, Cell tracking using bandpass filtering and the viterbi algorithm, Description of the algorithm available at: http://www.codesolorzano.com/celltrackingchallenge/
- (4) Y. Boykov and G. Funka-Lea, "Graph cuts and efficient n-d image segmentation," IJCV, vol. 70, no. 2, pp. 109-131, 2006.



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# Segmentation energy functional

UNI FREIBURG Cost function (Region & boundary term)

$$E(M) = \lambda \cdot R(M) + B(M)$$

 $\operatorname{Mask} M : \Omega \to \{0, 1\},\$  $\Omega \subset \mathbb{R}^2$ 

Boundary term



# Segmentation energy functional

UNI FREIBURG Cost function (Region & boundary term)

$$E(M) = \lambda \cdot R(M) + B(M)$$

 $\operatorname{Mask} M : \Omega \to \{0, 1\},\$  $\Omega \subset \mathbb{R}^2$ 

Boundary term

$$B(M) = \int_{\Omega} C_{\text{edge}} \left( \underbrace{\langle \nabla M(\mathbf{x}), -\nabla I(\mathbf{x}) \rangle}_{\text{intensity derivative } d} \right) d\mathbf{x} \qquad \text{Image } I$$
  
(perpendicular to mask boundary) 
$$O(I) = \int_{\Omega} C_{\text{edge}} \left( \underbrace{\langle \nabla M(\mathbf{x}), -\nabla I(\mathbf{x}) \rangle}_{\text{intensity derivative } d} \right) d\mathbf{x}$$
  
(perpendicular to mask boundary) 
$$O(I) = \int_{\Omega} C_{\text{edge}} \left( \underbrace{\langle \nabla M(\mathbf{x}), -\nabla I(\mathbf{x}) \rangle}_{\text{intensity derivative } d} \right) d\mathbf{x}$$

Asymmetric boundary penalties (dark-to-bright)

$$C_{\text{edge}}(d) = \begin{cases} \exp\left(-\frac{d^2}{2\sigma^2}\right) & \text{if } d > 0\\ 1 & \text{else.} \end{cases}$$

 $\rightarrow$  directed graph with asymmetric edge weights

## Symmetric vs. asymmetric penalties



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> 3x3 pixel neighborhood, Edges and weights (only outwards edges shown)

# Symmetric boundary penalties





BURG

3x3 pixel neighborhood, Edges and weights (only outwards edges shown)



 Low costs at wrong cell borders (bright-to-dark transitions)

# Asymmetric boundary penalties





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> 3x3 pixel neighborhood, Edges and weights (only outwards edges shown)



• Low costs at **correct cell borders** (dark-to-bright transitions)



## Asymmetric boundary penalties





Cyan mask: Segmentation result of graph cut with **symmetric costs** Yellow: Our manual ground truth



Red mask: Segmentation result of **proposed method** Yellow: Our manual ground truth

# **Regional penalties**

- 55
- UNI FREIBURG Standard graph cut (negative log-likelihood)

$$R(A) = \sum_{p \in \mathcal{P}} R_p(A_p) \quad \text{(regional term)}$$

 $R_p(\text{"obj"}) = -\ln \Pr(I_p | \text{"obj"}) \text{ (object penalty)}$  $R_p("bkg") = -\ln \Pr(I_p|"bkg")$  (background penalty)  $\rightarrow$  hard constraint

# **Regional penalties**

UNI FREIBURG Standard graph cut (negative log-likelihood)

$$R(A) = \sum_{p \in \mathcal{P}} R_p(A_p) \quad \text{(regional term)}$$

 $R_p(\text{"obj"}) = -\ln \Pr(I_p | \text{"obj"}) \text{ (object penalty)}$  $R_p("bkg") = -\ln \Pr(I_p|"bkg")$  (background penalty)  $\rightarrow$  hard constraint

In our approach

$$R(M) = \int_{\Omega} M(\mathbf{x}) \cdot C_{\text{obj}}(I(\mathbf{x})) d\mathbf{x} \quad \text{(regional term)}$$
$$C_{\text{obj}}(v) = \frac{P(v|\mathcal{B}) - P(v|\mathcal{O})}{P(v|\mathcal{O}) + P(v|\mathcal{B})} \quad \text{(data costs)} \quad \begin{array}{l} \text{Intensity } v \\ P(v|\mathcal{O}) \text{ and } P(v|\mathcal{B}) \\ \text{from fore-/background} \end{array}$$

#### $\rightarrow$ soft constraint

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intensity histograms





$$\begin{split} E(M) &= \lambda \int_{\Omega} M(\mathbf{x}) \cdot C_{\text{obj}}(I(\mathbf{x})) d\mathbf{x} \\ &+ \int_{\Omega} C_{\text{edge}} \left( \langle \nabla M(\mathbf{x}), -\nabla I(\mathbf{x}) \rangle \right) d\mathbf{x} \end{split}$$

• Enery minimization problem

Optimization

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Discretize edge term into 8 directions
 → combinatorial optimization problem



• Solve efficiently by a **min-cut approach** 



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# Tracking: Segmentation propagation





#### 1) Propagate Segmentation Information

#### a) Foreground information

using eroded mask

 $\rightarrow$  hard foreground constraint

#### b) Partitioning information

using borders of "support regions"  $\rightarrow$  hard background constraint

# Tracking: Label propagation





2) Propagate Labels to overlapping Segments using max. **IoU** 

a) Resolve one-to-many correspondences

start new tracks (with new label)

b) Resolve many-to-one correspondences

stop other tracks



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# Datasets: ISBI cell tracking challenge<sup>1,2</sup>



- Strong shape variations
- Weak outer borders, strong irrelevant inner borders
- Cytoplasm has same structure as background

Glioblastoma-astrocytoma U373 cells on a polyacrylimide substrate\* Phase contrast microscopy

(1) ISBI Cell Tracking Challenge, Available at: http://www.codesolorzano.com/celltrackingchallenge.
(2) M. Maška, V. Ulman, D. Svoboda, P. Matula, and P. Matula, et al., "A benchmark for comparison of cell tracking algorithms," Bioinformatics, vol. 30, no. 11, pp. 1609–1617, 2014.
\*Data provided by Dr. Sanjay Kumar, Department of Bioengineering University of California at Berkeley, Berkeley CA (USA).

Boundary	Seq. 1			Seq. 2			
costs	F-meas.	Recall	Prec.	F-meas.	Recall	Prec.	
Symm.	0.863	0.838	0.889	0.768	0.732	0.808	
Asymm. (Equ. 2)	0.896	0.894	0.897	0.835	0.822	0.847	
Boundary detection F-measure, recall and precision (4 pixels tolerance)							

- Boundary detection recall and precision\*
- Symmetric vs. asymmetric boundary costs

\*Computed using code from "The Berkeley Segmentation Dataset and Benchmark", Available at: http://www.eecs.berkeley.edu/Research/Projects/CS/vision/bsds/.

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### Stability of results





F-measure isolines



Boundary detection results across varying min-cut parameters lambda and sigma.

#### Experiments: Symmetric vs. asymmetric costs



Cyan masks: Graph cut with symmetric costs, Red masks: Our approach with asymmetric costs, Yellow borders: Our manual ground truth

- Improved detection of very weak boundaries
- Halo boundaries are handled well

# Preliminary results: ISBI cell tracking challenge

Group	Av. SEG	Av. TRA
KTH-SE	0.7953	0.9818
HOUS-US	0.5323	0.9206
IMCB-SG	0.2669	0.9595

Reported results on the "challenge dataset"

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Sequence Av. SEG Av. TRA

Seq. 1	0.8648	0.9830
Seq. 2	0.7563	0.9150
Seq. 1+2	0.8105	0.9490

Our preliminary results on the "training dataset"

- Comparison against top ranked methods from last years ISBI cell tracking challenge
- Phase contrast dataset: PhC-C2DH-U373

#### Submitted results: PhC-C2DH-U373











## Conclusion



- UNI FREIBURG Direction dependent boundary costs improve segmentation in phase contrast microscopy
  - Our approach outperforms standard min-cut segmentation with symmetric costs
  - Preliminary results suggest competitive performance with top-ranked methods in the ISBI CTC



 $\rightarrow$  Profit for cell segmentation in other modalities

 $\rightarrow$  Open-source MATLAB code (and ImageJ plugin): http://lmb.informatik.uni-freiburg.de/resources/opensource/CellTracking/







### Thank you!



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