

Statistical Consulting

Segmentation of Myotubes and Myoblasts

Department of Statistics
Ludwig-Maximilians-Universität München

Giorgi Nozadze, Moustafa Amin, David Gorbunov

Munich, January 30th, 2024



Supervisors: Prof. Dr. David Rügamer, Tobias Weber

Project Partner: Dr. Maximilian Saller

1 Abstract

Lorem ipsum dolor sit amet, consectetur adipiscing elit. Ut purus elit, vestibulum ut, placerat ac, adipiscing vitae, felis. Curabitur dictum gravida mauris. Nam arcu libero, nonummy eget, consectetur id, vulputate a, magna. Donec vehicula augue eu neque. Pellentesque habitant morbi tristique senectus et netus et malesuada fames ac turpis egestas. Mauris ut leo. Cras viverra metus rhoncus sem. Nulla et lectus vestibulum urna fringilla ultrices. Phasellus eu tellus sit amet tortor gravida placerat. Integer sapien est, iaculis in, pretium quis, viverra ac, nunc. Praesent eget sem vel leo ultrices bibendum. Aenean faucibus. Morbi dolor nulla, malesuada eu, pulvinar at, mollis ac, nulla. Curabitur auctor semper nulla. Donec varius orci eget risus. Duis nibh mi, congue eu, accumsan eleifend, sagittis quis, diam. Duis eget orci sit amet orci dignissim rutrum.

Contents

1	Abstract	
2	Project Goal	1
3	Segmentation using Unsupervised Methods	2
3.1	Watershed	2
4	Stardist	3
5	Segmentation of Myotubes with MyoSAM	4
6	Results	5
7	Conclusion	6
A	Appendix	

2 Project Goal

Myogenesis is the process describing the formation, growth, development, and regeneration of muscle tissue in the body. Therein mononucleated, partially differentiated precursor cells, also called myoblasts, proliferate, migrate, differentiate, and eventually fuse and thereby form multinucleated fibers known as myotubes. The myotubes then mature into myofibers.

Given the spatiotemporal nature of myogenesis, it is instrumental to be able to segment both myotubes and myoblasts from microscopy images since it enables more systematic and quantitative research in this area. Other than simply counting the number of cells, creating an overlay of myotubes and myoblasts in order to find how many myoblasts are encapsulated by a myotube can tell you what stage of myogenesis certain cells are in. This also tells you how many cells are quiescent and yet to differentiate. In a similar vein, this holds true for the measurement of the area and diameter of myotubes. This project intends to obtain such quantitative statistics without using heuristics. To this end, two models are needed: one that performs instance segmentation for the myoblasts and one for the myotubes.

These models need to overcome two hurdles. First and foremost, it must be able to differentiate between overlapping instances. Besides coincidence, these overlaps may stem from the myogenesis itself, e.g. during cell fusion. It may also be due to the fact that samples is not truly two dimensional such that cells from underneath can cause such overlaps by shining through. Secondly, it should be robust in its predictions. Many times, microscopy images require preprocessing which can create artifacts, amplifying noise, or blur small scale structures. The instance segmentation should aim to be as independent of the preprocessing as possible.

All in all, the goal is not only to speed up otherwise time consuming analyses, but also to improve reproducibility and eliminate all human bias within the process of segmentation. Human bias can occur at various points. Many times it is difficult to distinguish whether a continuously bright region is one object or several ones in actuality. Furthermore, the quality of preprocessing can influence the number of counted cells because some instances may be too dim too spot with the naked eye.

3 Segmentation using Unsupervised Methods

3.1 Watershed

4 Stardist

5 Segmentation of Myotubes with MyoSAM

6 Results

7 Conclusion

A Appendix