

Physarum Computations

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Abstract

Physarum is a slime mold. It was observed over the past 10 years that the mold is able to solve shortest path problems and to construct good Steiner networks [9, 11, 8]. In a nutshell, the shortest path experiment is as follows: A maze is covered with mold and food is then provided at two positions s and t and the evolution of the slime is observed. Over time, the slime retracts to the shortest s - t -path. A video showing the wet-lab experiment can be found at <http://www.youtube.com/watch?v=tL02n3YMcXw&t=4m43s>. We strongly recommend to watch this video.

A mathematical model of the slime's dynamic behavior was proposed in 2007 [10]. Extensive computer simulations of the mathematical model confirm the wet-lab findings. For the edges on the shortest path, the diameter converges to one, and for the edges off the shortest path, the diameter converges to zero.

We review the wet-lab and the computer experiments and provide a proof for these experimental findings. The proof was developed over a sequence of papers [6, 7, 4, 2, 1, 3]. We recommend the last two papers for first reading.

An interesting connection between Physarum and ant computations is made in [5].

1998 ACM Subject Classification G.2.2 Graph Theory (Path and circuit problems)

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References

- 1 Vincenzo Bonifaci. Physarum can compute shortest paths: A short proof. *Inf. Process. Lett.*, 113(1-2):4–7, 2013.
- 2 Vincenzo Bonifaci, Kurt Mehlhorn, and Girish Varma. Physarum can compute shortest paths. In *SODA*, pages 233–240, 2012. full version to appear in *Journal of Theoretical Biology*.
- 3 Michael Dirnberger and Kurt Mehlhorn. A convergence proof for nonuniform directed Physarum. January 2013.
- 4 Kentaro Ito, Anders Johansson, Toshiyuki Nakagaki, and Atsushi Tero. Convergence properties for the physarum solver. arXiv:1101.5249v1, January 2011.
- 5 Qi Ma, Anders Johansson, Atsushi Tero, Toshiyuki Nakagaki, and David J. T. Sumpter. Current reinforced random walks for biological problem solving. 2012.
- 6 T. Miyaji and Isamu Ohnishi. Mathematical analysis to an adaptive network of the plasmodium system. *Hokkaido Mathematical Journal*, 36:445–465, 2007.
- 7 T. Miyaji and Isamu Ohnishi. Physarum can solve the shortest path problem on riemannian surface mathematically rigourously. *International Journal of Pure and Applied Mathematics*, 47:353–369, 2008.
- 8 T. Nakagaki, M. Iima, T. Ueda, Y. Nishiura, T. Saigusa, A. Tero, R. Kobayashi, and K. Showalter. Minimum-risk path finding by an adaptive amoebal network. *Physical Review Letters (PRL)*, 99(068104):4, 2007.



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- 9 T. Nakagaki, H. Yamada, and Á. Tóth. Maze-solving by an amoeboid organism. *Nature*, 407:470, 2000.
- 10 A. Tero, R. Kobayashi, and T. Nakagaki. A mathematical model for adaptive transport network in path finding by true slime mold. *Journal of Theoretical Biology*, pages 553–564, 2007.
- 11 A. Tero, S. Takagi, T. Saigusa, K. Ito, D. Bebbler, M. Fricker, K. Yumiki, R. Kobayashi, and T. Nakagaki. Rules for biologically inspired adaptive network design. *Science*, 327:439–442, 2010.