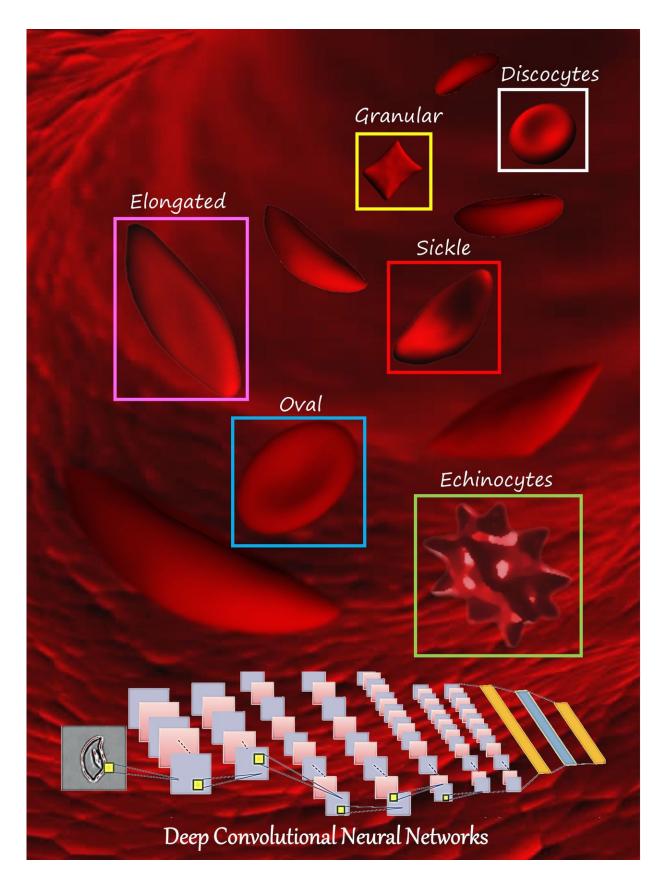


New machine learning system can automatically identify shapes of red blood cells

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Classifying Sickle Cell Anemia RBC in an automated manner with high accuracy based on Deep Convolutional Neural Network method for 8 SCD patients (over 7,000 single RBC images) for both oxygenated and deoxygenated RBCs. Credit: Xu et al.

Using a computational approach known as deep learning, scientists have developed a new system to classify the shapes of red blood cells in a patient's blood. The findings, published in *PLOS Computational Biology*, could potentially help doctors monitor people with sickle cell disease.

A person with sickle cell disease produces abnormally shaped, stiff red blood cells that can build up and block blood vessels, causing pain and sometimes death. The disease is named after sickle-shaped (crescentlike) red blood cells, but it also results in many other shapes, such as oval or elongated red blood cells. The particular shapes found in a given patient can hold clues to the severity of their disease, but it is difficult to manually classify these shapes.

To automate the process of identifying <u>red blood cell</u> shape, Mengjia Xu of Northeastern University, China, and colleagues developed a computational framework that employs a machine-learning tool known as a deep <u>convolutional neural network</u> (CNN).

The new framework uses three steps to classify the shapes of red blood cells in microscopic images of blood. First, it distinguishes red blood cells from the background of each image and from each other. Then, for each cell detected, it zooms in or out until all cell images are a uniform size. Finally, it uses deep CNNs to categorize the cells by shape.

The researchers validated their new tool using 7,000 microscopy images from eight <u>sickle cell disease</u> patients. They found that the automated



method successfully classified red blood cell shape for both oxygenated and deoxygenated cells (red blood cells transport oxygen to tissues throughout the body).

"We have developed the first <u>deep learning</u> tool that can automatically identify and classify red blood cell alteration, hence providing direct quantitative evidence of the severity of the disease," says study co-author George Karniadakis.

The research team plans to further improve their deep CNN tool and test it in other blood diseases that alter the shape and size of red blood cells, such as diabetes and HIV. They also plan to explore its usefulness in characterizing <u>cancer cells</u>.

More information: Xu M, Papageorgiou DP, Abidi SZ, Dao M, Zhao H, Karniadakis GE (2017) A deep convolutional neural network for classification of red blood cells in sickle cell anemia. *PLoS Comput Biol* 13(10): e1005746. <u>doi.org/10.1371/journal.pcbi.1005746</u>

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