

MCFIP - The following article supports our assertion that the future of computational biology will be guided by various algorithms; i.e. EpiMogrify.

As explained in the following, when principles for biomedical science were not available to address interactions and imbalances that included particle physics, they required a new algorithm.

<https://www.mcfip.net/upload/7%20-%20Transitioning%20of%20Computational%20Biology.pdf>

Three people working 15 years each were required to design an algorithm for quantum biology (QB) for use as a foundation for addressing the causes of chronic diseases for use by computational biologists. Refer to the following for discussion.

<https://www.mcfip.net/upload/Integrating%20Science.pdf>

<https://phys.org/news/2020-10-approach-optimising-culture-conditions-required.html>

OCTOBER 12, 2020

## **Computational approach shows promise for optimising culture conditions required for cell therapy**

by [Duke-NUS Medical School](#)

Cellular therapy is a powerful strategy to produce patient-specific, personalized cells to treat many diseases, including heart disease and neurological disorders. But a major challenge for cell therapy applications is keeping cells alive and well in the lab.

That may soon change as researchers at Duke-NUS Medical School, Singapore, and Monash University, Australia have devised an algorithm that can predict what molecules are needed to keep cells healthy in laboratory cultures. They developed a computational approach called EpiMogrify, that can predict the molecules needed to signal stem cells to change into specific tissue cells, which can help accelerate treatments that require growing patient cells in the lab.

"Computational biology is rapidly becoming a key enabler in cell therapy, providing a way to short-circuit otherwise expensive and time-consuming discovery approaches with cleverly designed algorithms," said Assistant Professor Owen Rackham, a computational biologist at Duke-NUS, and a senior and corresponding author of the study, published today in the journal *Cell Systems*.

In the laboratory, cells are often grown and maintained in cell cultures, formed of a substance, called a medium, which contains nutrients and other molecules. It has been an ongoing challenge to identify the necessary molecules to maintain high-quality cells in culture, as well as finding molecules that can induce stem cells to convert to other cell types.

The research team developed a computer model called EpiMogrify that successfully identified molecules to add to cell culture media to maintain healthy nerve cells, called astrocytes, and heart cells, called cardiomyocytes. They also used their model to successfully predict molecules that trigger stem cells to turn into astrocytes and cardiomyocytes.

"Research at Duke-NUS is paving the road for cell therapies and regenerative medicine to enter the clinic in Singapore and worldwide; this study leverages our expertise in computational and systems biology to facilitate the good manufacturing practice (GMP) production of high-quality cells for these much needed therapeutic applications," said Associate Professor Enrico Petretto, who leads the Systems Genetics group at Duke-NUS, and is a senior and corresponding author of the study.

The researchers added existing information into their model about genes tagged with epigenetic markers whose presence indicates that a gene is important for cell identity. The model then determines which of these genes actually code for proteins necessary for a cell's identity. Additionally, the model incorporates data about proteins that bind to cell receptors to influence their activities. Together, this information is used by the computer model to predict specific proteins that will influence different cells' identities.

"This approach facilitates the identification of the optimum cell culture conditions for converting cells and also for growing the high-quality cells required for cell therapy applications," said ARC Future Fellow Professor Jose Polo, from Monash University's Biomedicine Discovery Institute and the Australian Research Medicine Institute, who is also a senior and corresponding author of the study.

The team compared cultures using protein molecules predicted by EpiMogrify to a type of commonly used cell culture that uses a large amount of unknown or undefined complex molecules and chemicals. They found the EpiMogrify-predicted cultures worked as well or even surpassed their effectiveness.

The researchers have filed for a patent on their computational approach and the cell culture factors it predicted for maintaining and controlling cell fate. EpiMogrify's predicted molecules are available for other researchers to explore on a [public database](#).

"We aim to continue to develop tools and technologies that can enable cell therapies and bring them to the clinic as efficiently and safely as possible," said Asst Prof Rackham.

"The developed technology can identify cell culture conditions required to manipulate cell fate and this facilitates growing important cells in chemically-defined cultures for cell therapy applications," added Dr. Uma S. Kamaraj, lead author of the study and a graduate of Duke-NUS' Integrated Biology and Medicine Ph.D. Program.