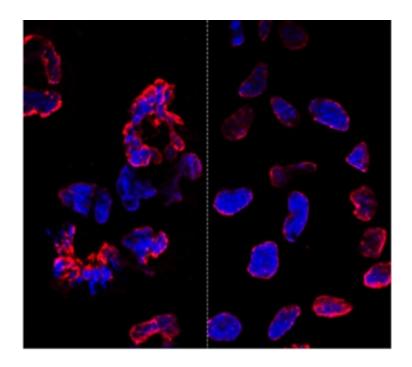


## No extra mutations in modified stem cells, study finds

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The new study shows that gene-editing technologies are specific to their targets and do not introduce harmful mutations, clearing the way for the development of safe therapies in the clinic. The left panel shows misshapen nuclear envelopes (red) from induced pluripotent stem cells derived from cells with Parkinson's disease (DNA in blue). The right panel shows similarly induced cells that have been gene-edited to restore the cells. Credit: The Salk Institute for Biological Studies

The ability to switch out one gene for another in a line of living stem cells has only crossed from science fiction to reality within this decade.



As with any new technology, it brings with it both promise—the hope of fixing disease-causing genes in humans, for example—as well as questions and safety concerns. Now, Salk scientists have put one of those concerns to rest: using gene-editing techniques on stem cells doesn't increase the overall occurrence of mutations in the cells. The new results were published July 3 in the journal *Cell Stem Cell*.

"The ability to precisely modify the DNA of <u>stem cells</u> has greatly accelerated research on human diseases and cell therapy," says senior author Juan Carlos Izpisua Belmonte, professor in Salk's Gene Expression Laboratory. "To successfully translate this technology into the clinic, we first need to scrutinize the safety of these modified stem cells, such as their genome stability and mutational load."

When scientists want to change the sequence of a stretch of DNA inside cells—either for research purposes or to fix a genetic mutation for therapeutic purposes—they have their choice of two methods. They can use an engineered virus to deliver the new gene to a cell; the cell then integrates the new DNA sequence in place of the old one. Or scientists can use what's known as custom targeted nucleases, such as TALEN proteins, which cut DNA at any desired location. Researchers can use the proteins to cut a gene they want to replace, then add a new gene to the mix. The cell's natural repair mechanisms will paste the new gene in place.

Previously, Belmonte's lab had pioneered the use of modified viruses, called helper-dependent adenoviral vectors (HDAdVs) to correct the gene mutation that causes <u>sickle cell disease</u>, one of the most severe blood diseases in the world. He and his collaborators used HDAdVs to replace the <u>mutated gene</u> in a line of stem cells with a mutant-free version, creating stem cells that could theoretically be infused into patients' bone marrow so that their bodies create healthy blood cells.



Before such technologies are applied to humans, though, researchers like Belmonte wanted to know whether there were risks of editing the genes in stem cells. Even though both common gene-editing techniques have been shown to be accurate at altering the right stretch of DNA, scientists worried that the process could make the cells more unstable and prone to mutations in unrelated genes—such as those that could cause cancer.



This image depicts, starting from the left, Juan Carlos Izpisua Belmonte, Keiichiro Suzuki and Mo Li of the Gene Expression Laboratory. Credit: The Salk Institute for Biological Studies

"As cells are being reprogrammed into stem cells, they tend to accumulate many mutations," says Mo Li, a <u>postdoctoral fellow</u> in Belmonte's lab and an author of the new paper. "So people naturally worry that any process you perform with these cells in vitro—including gene editing—might generate even more mutations."



To find out whether this was the case, Belmonte's group, in collaboration with BGI and the Institute of Biophysics, Chinese Academy of Sciences in China, turned to a line of stem cells containing the mutated gene that causes sickle cell disease. They edited the genes of some cells using one of two HDAdV designs, edited others using one of two TALEN proteins, and kept the rest of the cells in culture without editing them. Then, they fully sequenced the entire genome of each cell from the four edits and control experiment.

While all of the cells gained a low level of random gene mutations during the experiments, the cells that had undergone gene-editing—whether through HDAdV- or TALEN-based approaches—had no more mutations than the cells kept in culture.

"We were pleasantly surprised by the results," Keiichiro Suzuki, a postdoctoral fellow in Belmonte's lab and an author of the study, says. "People have found thousands of mutations introduced during iPSC reprogramming. We found less than a hundred single nucleotide variants in all cases."

The finding, Li adds, doesn't necessarily mean that there are no inherent risks to using stem cells with edited genes, but that the editing process doesn't make the stem cells any less safe.

"We concluded that the risk of mutation isn't inherently connected to gene editing," he says. "These cells present the same risks as using any other cells manipulated for cell or gene therapy." He adds that two other papers published in the same issue support their results (one by Johns Hopkins University and one from Harvard University and collaborators).

The Belmonte group is planning more studies to address whether generepair in other cell types, using other approaches, or targeting other genes could be more or less likely to cause unwanted mutations. For



now, they hope their findings encourage those in the field to keep pursuing gene-editing techniques as a potential way to treat genetic diseases in the future.

Other researchers on the study were Jing Qu, April Goebl, Emi Aizawa, Rupa Devi Soligalla, Jessica Kim, Na Young Kim, Hsin-Kai Liao, Chris Benner, and Concepcion Rodriguez Esteban of the Salk Institute for Biological Studies; Chang Yu, Xiaotian Yao, Senwei Tang, Fan Zhang, Feng Chen, Yabin Jin, and Yingrui Li of BGI; and Jing Qu, Tingting Yuan, Ruotong Ren, Xiuling Xu, and Guang-Hui Liu of the Institute of Biophysics, Chinese Academy of Sciences.

**More information:** www.sciencedirect.com/science/ ... ii/S1934590914002641

## Provided by Salk Institute

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