

Scientists discover 'fickle' DNA changes in brain

September 30 2011

Johns Hopkins scientists investigating chemical modifications across the genomes of adult mice have discovered that DNA modifications in nondividing brain cells, thought to be inherently stable, instead underwent large-scale dynamic changes as a result of stimulated brain activity. Their report, in the October issue of *Nature Neuroscience*, has major implications for treating psychiatric diseases, neurodegenerative disorders, and for better understanding learning, memory and mood regulation.

Specifically, the researchers, who include a husband-and-wife team, found evidence of an epigenetic change called demethylation — the loss of a methyl group from specific locations — in the non-dividing brain cells' DNA, challenging the scientific dogma that even if the DNA in non-dividing adult neurons changes on occasion from methylated to demethylated state, it does so very infrequently.

"We provide definitive evidence suggesting that DNA demethylation happens in non-dividing neurons, and it happens on a large scale," says Hongjun Song, Ph.D., professor of neurology and neuroscience and director of the Stem Cell Program in the Institute for Cell Engineering of the Johns Hopkins University School of Medicine. "Scientists have previously underestimated how important this epigenetic mechanism can be in the adult brain, and the scope of change is dramatic."

DNA comprises the fixed chemical building blocks of each person or animal's <u>genome</u>, but the addition or removal of a methyl group at the



specific location chemically alters DNA and regulates gene expression, enabling cells with the same genetic code to acquire and activate separate functions.

In previously published work, the same Hopkins researchers reported that electrical brain stimulation, such as that used in electroconvulsive therapy (ECT) for patients with drug resistant depression, resulted in increased brain cell growth in mice, due likely to changes in DNA methylation status.

This time, they again used electric shock to stimulate the brains of live mice. A few hours after administering the brain stimulation, the scientists analyzed two million of the same type of neurons from the brains of stimulated mice, focusing on what happens to one building block of DNA — cytosine — at 219,991 sites. These sites represented about one percent of all cytosines in the whole mouse genomes.

In collaboration with genomic biologist Yuan Gao, now at the Lieber Institute for Brain Development, the scientists used the latest DNA sequencing technology and compared neurons in mice with or without brain stimulation. About 1.4 percent of the cytosines measured showed rapid active demethylation or became newly methylated.

"It was mind-boggling to see that so many methylation sites — thousands of sites — had changed in status as a result of <u>brain activity</u>," Song says. "We used to think that the brain's epigenetic DNA methylation landscape was as stable as mountains and more recently realized that maybe it was a bit more subject to change, perhaps like trees occasionally bent in a storm. But now we show it is most of all like a river that reacts to storms of activity by moving and changing fast."

The majority of the sites where the methylation status of the cytosine changed as a result of the <u>brain</u> activity were not in the expected areas of



the genome that are traditionally believed to control gene expression, Song notes. Rather, they were in regions where cytosines are low in density, in genomic regions where the function of DNA methylation is not well understood.

Because DNA demethylation can occur passively during cell division, the scientists targeted radiation to the sections of mouse brains they were studying, permanently preventing passive cell division, and still found evidence of DNA demethylation. This confirms, they say, that the DNA methylation changes they measured occurred independently of cell division.

"Our finding opens up new opportunities to figure out if these epigenetic modifications are potential drug targets for treating depression and promote regeneration, for instance," says Guo-li Ming, M.D., Ph.D., professor of neurology and neuroscience.

Provided by Johns Hopkins Medical Institutions

Citation: Scientists discover 'fickle' DNA changes in brain (2011, September 30) retrieved 5 May 2024 from <u>https://medicalxpress.com/news/2011-09-scientists-fickle-dna-brain.html</u>

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