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## TET proteins help maintain genome integrity

LJI researchers show that TET-loss of function causes rapid development of malignant cancer

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LA JOLLA, CA—Members of the TET (short for ten-eleven translocation) family have been known to function as tumor suppressors for many years, but how they keep a lid on the uncontrolled cell proliferation of cancer cells had remained uncertain. Now, researchers at the La Jolla Institute for Allergy and Immunology demonstrate that TET proteins collectively constitute a major class of tumor suppressors and are required to maintain genome instability.

The study, published in the current edition of *Nature Communications*, shows that mice lacking both Tet2 and Tet3 rapidly develop an aggressive form of myeloid leukemia. "We removed Tet2 and Tet3 by injecting the mice with chemicals that acutely deleted their genes, after which we could see signs of cancer in one week and a full-blown transmissible cancer in four weeks," says senior author Anjana Rao, Ph.D., a professor in the Institute's Division of Signaling and Gene Expression who led the study. "This proves that TET-loss of function directly caused the malignancy."

The gene coding for Tet2 is frequently mutated in patients with myeloid malignancies, a cancer of a type of white blood cells, and even without Tet2 mutations, many human myeloid cancers display low TET function as Rao and her team had shown in an earlier study (Ko et al., Nature 2010.) "The same is true for many other types of cancers," says the study's coauthor Myunggon Ko, Ph.D., formerly a postdoctoral researcher in the Rao lab and now an assistant professor at the Ulsan National Institute of Science and Technology in Ulsan, Korea. "Thus our data imply that TET function is likely to be important for preventing not only myeloid cancers, but also potentially other types of cancers."

Members of the TET family of enzymes help rewrite the epigenome, the regulatory layer of chemical modifications that sits atop the genome and helps determine gene activity without changing the letters of DNA. Specifically, TET enzymes convert a modified form of cytosine, one of the four bases of DNA, called 5-methylcytosine (5mC) to 5-hydroxymethylcytosine (5hmC) and other forms of oxidized methylcytosine. Together these oxidized forms of 5mC facilitate DNA demethylation but are also important in their own right.

"Our findings indicate that loss of TET function contributes to skewing blood stem cells in favor of forming myeloid cells over other blood cell types by regulating the expression of lineage-specific genes," says Ko. "Surprisingly, although cytosine methylation has long been thought to inhibit the activity of genes, there was no clear correlation between DNA methylation and gene expression. They were independently controlled and we may have to change our view about the relation between DNA methylation and gene expression." Rao adds, "One possibility is that

the precipitating factor is loss of oxidized methylcytosines rather than changes in DNA methylation per se."

Deleting just one of the three members of the TET family (*Tet1*, *Tet2* and *Tet3*)—although insufficient to induce full-blown leukemia in a model system—leads to the expansion of the stem cells that give rise to all blood cells. This process is exacerbated in stem cells with more severely impaired TET function, such as that seen in this study in mice lacking both Tet2 and Tet3. The capacity for self-renewal is enhanced and the cells preferentially differentiate into cells belonging to the myeloid lineage, such as macrophages and neutrophils.

The researchers also observed a strong impairment of DNA damage repair, which suggests an important role for TET proteins in maintaining genome stability. "Genome instability is a defining characteristic of most cancers and a direct result of defective DNA damage repair," says Ko. "What we don't know yet is how TET loss of function leads to impaired DNA damage repair."

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## **Full citation:**

Acute loss of TET function results in aggressive myeloid cancer in mice. Jungeun An, Edahi Gonzales-Avalos, Ashu Chawla, Mira Jeong, Isaac F. Lopez-Moyado, Wei Ll, Margaret A. Goodell, Lukas Chavez, Myunggon Ko, Anjana Rao. *Nature Communications* (2015). DOI: 10.1038/ncomms10071

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