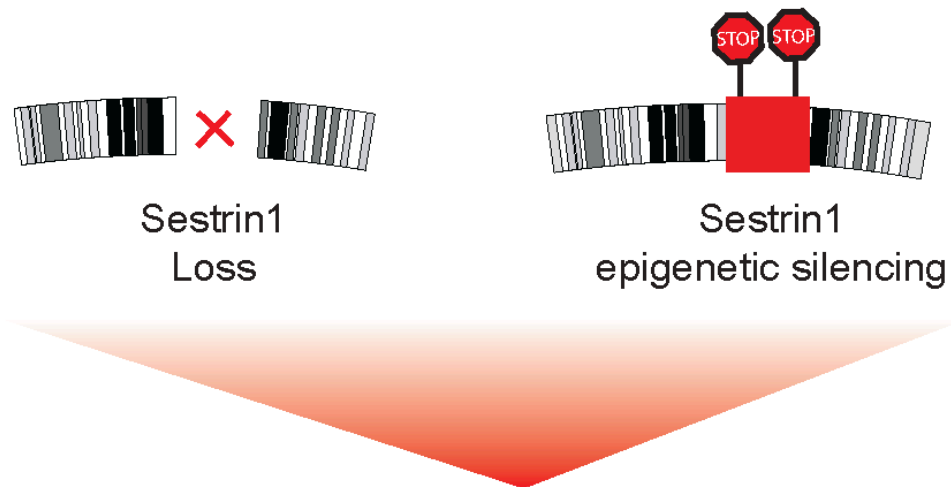
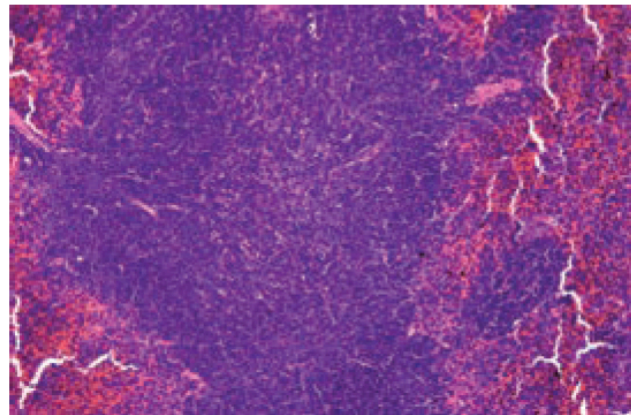


The gene behind follicular lymphoma

June 28 2017



Follicular lymphoma



Disruption of a region in chromosome 6 or epigenetic modifications of the DNA block Sestrin1 expression and these contribute to the development of follicular

lymphoma. Credit: Elisa Oricchio/Natalya Katanayeva/EPFL

Follicular lymphoma is an incurable cancer that affects over 200,000 people worldwide every year. A form of non-Hodgkin lymphoma, follicular lymphoma develops when the body starts making abnormal B-cells, which are white blood cells that in normal conditions fight infections. This cancer is associated with several alterations of the cell's DNA, but it has been unclear which gene or genes are involved in its development. EPFL scientists have now analyzed the genomes of more than 200 patients with follicular lymphoma, and they discover that a gene, *Sestrin1*, is frequently missing or malfunctioning in FL patients. The discovery opens to new treatment options and it is now published in *Science Translational Medicine*.

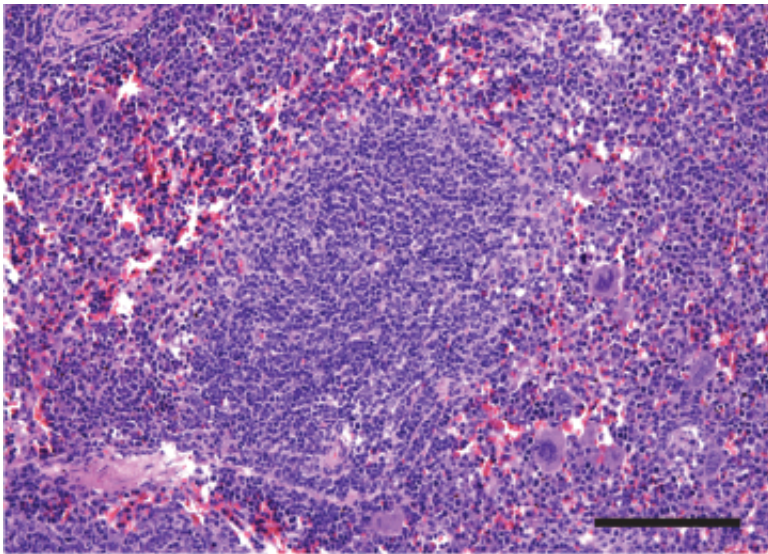
One of the common features of [follicular lymphoma](#) is a genetic abnormality between two chromosomes (14 and 18). In an event known as "[chromosomal translocation](#)" the two chromosomes "swap" certain parts with each other. This triggers the activation of a gene that protects cells from dying, making cells virtually immortal—the hallmark of a tumor.

Moreover, approximately 30% of follicular lymphoma patients lose also a portion of chromosome 6, affecting multiple genes involved in suppressing the emergence of a tumor. These patients typically have poor prognosis. Another 20 % of patients have alterations causing chromosomal disorganization and the consequent malfunctioning of several genes and proteins. The bottom line is that for both group of patients it is very difficult to pinpoint which of all the affected [genes](#) are actually causing the disease.

The lab of Elisa Oricchio at EPFL, with colleagues from the US and

Canada, analyzed the genomes of over 200 follicular lymphoma patients. Their analyses revealed that a specific gene, *Sestrin1*, can be harmed by both loss of chromosome 6 and silenced in patients.

Sestrin1 helps the cell defending itself against DNA damage—for example after exposure to radiation—and oxidative stress. In fact, *Sestrin1* is part of the cell's anti-tumor mechanism that stops potentially cancerous cells from growing.



Follicular lymphoma is the most common type of slow-growing blood cancer. Credit: E. Oricchio et al., *Science Translational Medicine* (2017)

Disruption of a region in chromosome 6 or epigenetic modifications of the DNA block *Sestrin1* expression and these contribute to the development of Follicular Lymphoma.

Beyond identifying the *Sestrin1* gene as frequently altered in FL patients, the scientists demonstrated that *Sestrin1* is able to suppress tumors in vivo. They showed that *Sestrin1* exerts its anti-tumor effects

by blocking the activity of a protein complex called mTORC1, which is well known for controlling protein synthesis as well as acting as a sensor for nutrient or energy changes in the cell.

Finally, the identification of loss of Sestrin1 as a key event behind the development of follicular lymphoma is particularly important because it helps identifying [patients](#) that will benefit from new therapies. Indeed, this study shows that the therapeutic efficacy of a new drug that is currently in clinical trial depends on Sestrin1. Importantly, this dependency can be extended beyond follicular lymphoma to other tumor types.

More information: E. Oricchio et al., "Genetic and epigenetic inactivation of SESTRIN1 controls mTORC1 and response to EZH2 inhibition in follicular lymphoma," *Science Translational Medicine* (2017). stm.sciencemag.org/lookup/doi/10.1126/scitranslmed.aak9969

Provided by Ecole Polytechnique Federale de Lausanne

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