

Evolutionarily novel genes at work in tumors

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Scientists from Peter the Great St. Petersburg Polytechnic University studied the evolutionary ages of human genes and identified a new class of them expressed in tumors -- tumor specifically expressed, evolutionarily novel (TSEEN) genes. Credit: Peter the Great St.Petersburg Polytechnic University

A team of scientists from Peter the Great St.Petersburg Polytechnic University (SPbPU) studied the evolutionary ages of human genes and identified a new class of them expressed in tumors—tumor specifically expressed, evolutionarily novel (TSEEN) genes. This confirms the team's earlier theory about the evolutionary role of neoplasms.



A report about the study was published in *Scientific Reports*.

A <u>tumor</u> is a pathological new growth of tissues. Due to genetic changes, it has impaired cellular regulation and therefore defective functionality. Tumors can be benign or malignant. Unlike the latter, the former grow slowly, don't metastasize, and are easy to remove. Malignant tumors (cancer) are one of the primary mortality factors in the world.

A team of scientists from Saint Petersburg discovered a new class of evolutionarily novel genes present in all tumors—the so-called TSEEN (Tumor Specifically Expressed Evolutionarily Novel) genes. "The evolutionary role of these genes is to provide genetic material for the origin of new progressive characteristics. TSEEN genes are expressed in many neoplasms and therefore can be excellent tumor markers," said Prof. Andrei Kozlov, a Ph.D. in Biology, the head of Laboratory "Molecular Virology and Oncology" at Peter the Great St. Petersburg Polytechnic University.

The new research confirms a theory that has been proposed by A. Kozlov earlier. According to it, the number of oncogenes in a <a href="https://human.com/hum

Prof. Kozlov also made a reference to the article "Evolutionarily Novel Genes Are Expressed in Transgenic Fish Tumors and Their Orthologs Are Involved in Development of Progressive Traits in Humans" (2019)



that has recently been published by his laboratory. In this article, the team confirmed their hypothesis using transgenic fish tumors and fish evolutionarily novel genes. The orthologs of such genes are found in the human genome, but in humans they play a role in the development of progressive characteristics not encountered in fish (e.g. lungs, breasts, placenta, ventricular septum in the heart, etc). This confirms the hypothesis about the evolutionary role of tumors. The studies referred to in the article lasted for several years, and their participants used a wide range of methods from the fields of bioinformatics and molecular biology.

"Our work is of great social importance, as the cancer problem hasn't been solved yet. Our theory suggests new prevention and therapy strategies," said Prof. Kozlov. According to him, to fight cancer, a new paradigm should be developed in oncology. TSEEN genes may be used to create new cancer test systems and antitumor vaccines.

More information: A. A. Makashov et al, Oncogenes, tumor suppressor and differentiation genes represent the oldest human gene classes and evolve concurrently, *Scientific Reports* (2019). <u>DOI:</u> 10.1038/s41598-019-52835-w

E. A. Matyunina et al. Evolutionarily novel genes are expressed in transgenic fish tumors and their orthologs are involved in development of progressive traits in humans, *Infectious Agents and Cancer* (2019). DOI: 10.1186/s13027-019-0262-5

Provided by Peter the Great Saint-Petersburg Polytechnic University

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