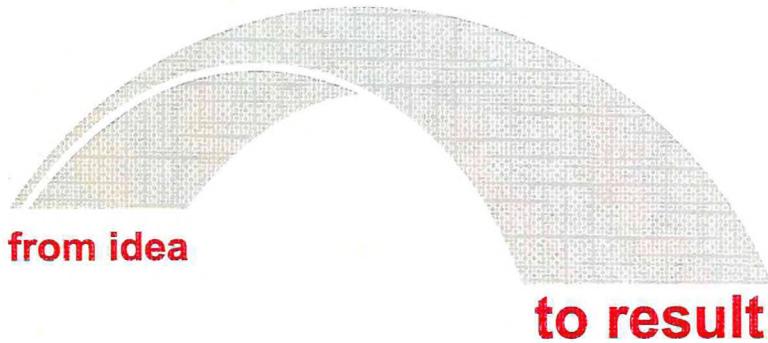


**International Conference
“Physics of Cancer:
Interdisciplinary Problems and Clinical Applications”**

Book of Abstracts

Tomsk, Russia, May 23-26, 2017



Institute of Strength Physics and Materials Science
of Siberian Branch Russian Academy of Sciences

INTERNATIONAL CONFERENCE
**Physics of cancer:
interdisciplinary problems and clinical applica-
tions**

May 23-26, 2017
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Book of Abstracts

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of cancer cells are much larger than the fluctuations of healthy cells, that corresponds to the data processing of cell dynamics by atomic force microscopy.



DYNAMIC CHANGES IN RETROTRANSPOSONS METHYLATION STATUS IN RESPONSE OF ANTITUMOR THERAPY OF LUNG CANCER

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Malignant cell transformation is accompanied by two processes of DNA methylation changes: promoter hypermethylation of single-copy genes and hypomethylation of retrotransposons. The composition of circulating DNA (cirDNA) from plasma and cell-surface-bound circulating DNA (csb-cirDNA) was shown earlier to be altered in the blood of cancer patients due to accumulation of tumor-specific aberrantly methylated DNA fragments, which are currently considered valuable cancer markers. The present study compared LINE-1 retrotransposon methylation patterns in plasma cirDNA and csb-cirDNA from healthy donors and lung cancer (LC) patients, and also from LC patients during the post-treatment follow-up period. It was shown that methylation level determination in csb-cirDNA from blood samples of LC patients appeared to be more informative than analysis of plasma cirDNA. Our data provide evidence that quantitative analysis of LINE-1 retrotransposon methylation status in cirDNA is a useful tool for lung cancer diagnostics, evaluation of cancer treatment efficiency and post-treatment monitoring.