

АКТУАЛЬНІ ПИТАННЯ ТЕРЕТИЧНОЇ ТА ПРАКТИЧНОЇ МЕДИЦИНИ

Topical Issues of Clinical and Theoretical Medicine

Збірник тез доповідей

IV Міжнародної науково-практичної конференції Студентів та молодих вчених (Суми, 21-22 квітня 2016 року)

TOM 1

Суми Сумський державний університет 2016 During the microscopic investigation of the experimental animals' thyroid, it was found that the major area of the organ in occupied by lobules of small sizes without clear division into center and periphery. The overall histological picture depicts tiny structure of the follicles. In the lumens of the majority of the follicles, there is an inconsiderable amount of desquamated epithelium. There are some lobules, which are separated from each other by interlayers of the hydropic stroma that has visually empty slots. In-between collagen interlayers of the interfollicular stroma there are found numerous flattened cells. In the other zones there are lobules, which are demarcated by the hydropic stroma to a lesser extent; the follicles vary in size: central lobules are tiny and those at the periphery are larger. In the tiny follicles and cellular aggregates, light-colored cells with vacuolated cytoplasm are prevailing. The inner contours of the tiny follicles are unclear in some areas. The lumen of the tiny follicles is filled with dimly eosinophilic colloid. Large follicles are generally filled with a colloid pink in color, some contain interfollicular epithelial outgrowths.

Thus, on the preparations of the experimental animals, we note evidences of the enhancement of the thyroid functional activity, which is indicated by vacuolated cytoplasm, rarefaction of the colloid and the increased formation of follicles. In the peripheral areas, folliculogenesis is implemented by follicle fragmentation with liberation of smaller "daughter" follicles.

GLOBAL METHYLATION STATUS IN MALIGNANT BRAIN TUMOR TISSUE.

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The DNA methylation is one of the main epigenetic inheritance form, which contributes in the regulation of gene expression. Abnormalities in DNA methylation processes can provide information about many pathophysiological conditions, including tumorigenesis. DNA hypomethylation was the initial epigenetic abnormality recognized in human tumors. Glioblastoma (GB) is the most common and the most aggressive primary brain tumor in adults and therefore is considered one of the major issue in modern medicine. The aim of our study was to compare global methylation status of DNA in peripheral blood cells and in biopsy tumor tissue from patients with diagnosed GB using Imprint® Methylated DNA Quantification Kit. Results of our study show global hypomethylation DNA status of GB tissues compare to global methylation DNA status of peripheral blood cells. Quantification of global methylation status confirm DNA hypomethylation in malignant brain tumor tissue, which may contribute in deregulation process of gene expression and subsequent tumor cell survival. Furthermore, detection of specific DNA methylation changes in GB tissue can subsequently help in understanding of specific genes activation and silencing through epigenetic events in tumor cells.

This work was supported by the Slovak Research and Development Agency under the contract No. APVV-0224-12 and by the project "Biomedical Center Martin" ITMS code: 26220220187, the project is co-financed from EU sources.

STRUCTURAL CHANGES OF RED BLOOD CELLS UNDER THE CONDITIONS OF VIOLATION OF WATER-SALT BALANCE

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Erythrocytes are important cells in your body that travel in the blood. They are involved in a gas exchange that is essential to human life. Erythrocytes – unique cells in our body, which lose their nucleus and other cytoplasmic organelles in the period of growth.