

ETIOPATHOGENETIC MECHANISMS OF CARCINOGENESIS

MARIA RADU¹, MANUELA MIHALACHE², COSMIN MIHALACHE³¹Hospice Palliative Care Centre Sibiu, ^{2,3}"Lucian Blaga" University of Sibiu

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Abstract: The appearance of the neoplastic process is based on a series of complex mechanisms which affect the structure of the DNA and the expression of several genes – genetic and epigenetic mechanisms. The importance of molecular mechanisms of cancerogenesis is multiple and it encounters its applicability in various fields of clinic oncology: diagnosis, prognostic, epidemiologic treatment.

Cuvinte cheie: mecanisme moleculare genetice și epigenetice, oncogene, gene supresoare, protooncogene, virusuri oncogene

Rezumat: Apariția procesului neoplazic are la bază o serie de mecanisme complexe care afectează structura ADN-ului și expresia mai multor gene – mecanisme genetice și epigenetice. Importanța mecanismelor moleculare ale cancerogenezei sunt multiple și își găsesc aplicabilitatea în diverse domenii ale oncologiei clinice: diagnostic, prognostic, tratament, epidemiologic.

The studies in recent years have led to a new formulation of the concept of cancer that claims that at cellular level cancer is a genetic disease of the somatic cells.(18) The transition from a completely normal cell to a malignant cell is done through a series of mutations due to alteration of genes involved in regulating cell growth and intercellular interactions. They can occur during DNA synthesis and cell replication, as a result of exposure to carcinogens in the environment or can be inherited as mutations of germ cells.(18)

GENETIC MECHANISMS, EPIGENETIC, OF CARCINOGENESIS:

The correlation of mutagenic action, of chemical substances, physical, exposure to viral oncogenes infections are arguments in favour of the involvement of the genetic mechanisms, epigenetic, in cancer.(17) The genetic mutational mechanism is reflected in the quantitative abnormalities (changes in the number of chromosomes) or qualitative of the karyotype that characterize tumour cells.(10) The epigenetic mechanisms involved in the process of carcinogenesis can be: DNA hypomethylation and genomic imprinting phenomenon.(13) The main types of genetic mechanisms of carcinogenesis, represented by chromosomal changes, are: nucleotide polymorphism, amplification, modulation, chromosomal translocation, deletion.(11) This occurs at the level of proto-oncogenes, oncogenes, and anti-oncogenes.

Proto-oncogenes: are categories of normal cellular genes that, when suffering structural and functional changes under the action of carcinogens, contribute to the malignant cell transformation.(11) Of the 30,000 human genes about 50 are proto-oncogenes.(2)

Oncogenes: are genes whose expression is intimately associated with the transformation of normal cells into

cancerous cells. Activation of oncogenes causes an increased predisposition to cancer.

Table no. 1. Representative oncogenic genes that activate human tumours (22)

| Oncogenic gens | Cellular function | Activated types of tumours | Mechanism of activation |
|---------------------|----------------------------------|------------------------------------------------------------|-------------------------------|
| EGFR/HER | Growth factor receptor | Glioblastoma, breast and lung cancer | Mutation Amplification |
| HER2/Neu | Growth factor receptor | Gastric, breast, lung cancer | Amplification |
| PRAD1/Cyclin D1 | Cell cycle regulator | Breast and esophageal cancer, lymphoma and parotid adenoma | Amplification Translation |
| K-ras, N-ras, H-ras | Protein Transduction signal | Breast carcinoma, pulmonary, colon, sarcoma | Mutation |
| B-Raf | Transduction signal | Multiple tumour types, melanoma | Mutation |
| src | Adhesion and cytoskeletal signal | Colon, breast, lung cancer and other functions | Unknown |
| | | sarcoma, melanoma | |
| Myc | Transcription factor | Multiple tumour types | Amplification Mutation |
| Myb | Transcription factor | Leukemia | Amplification Over-expression |
| Fas | Transcription factor | Multiple tumour types | Over-expression |
| Int2/FGF3 | Growth factor | Gastric esophageal cancer, head and neck | Amplification |

¹Corresponding author: Manuela Mihalache, Str. Lucian Blaga, Nr. 2A, Sibiu, România, E-mail: manuela.mihalache@ulbsibiu.ro, Tel: +0722 236312
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CLINICAL ASPECTS

There have been identified around 100 tumour suppressor genes and more than 100 dominant oncogenes involved in the carcinogenesis.(19) **Anti-oncogenes or tumour suppressor genes:** are genes that encode the synthesis of proteins involved in maintaining normal cellular function (e.g.: proteins having a role in reorganization and repair of DNA, proteins that regulate the cell cycle, protein that mediate growth factor signal).(22) **Gene p53:** it is a prototype of a suppressor gene; is the molecular link between etiologic agents of cancer (carcinogenesis factors) and the developing process of oncogenesis.(10) In more than half of the cancers appear mutations of p53. In the future determining the type and number of mutations in both p53 and suppressor genes in other tissues of healthy individuals, could identify individuals at risk for cancer.(10,8)

Table no. 2. Representative tumour suppressor genes that inactivate human tumours or hereditary syndromes (22)

| Tumour suppressor genes | Cellular function | Inactivated types of tumours | Mechanism of inactivation | Hereditary syndromes with line of alleles inactivated (14) |
|-------------------------|-------------------------------|-------------------------------------------------------|-------------------------------------|------------------------------------------------------------|
| p53(9) | Cell cycle regulator | Multiple tumour types | Mutation | Li-Fraumeni |
| Rb | Cell cycle regulator | Retinoblastoma, pulmonary cancer, small cell, sarcoma | Deletion | Family retinoblastoma |
| APC | Cellular adhesion | Colon cancer | Deletion | Family adenomatosis |
| PTEN | Transduction signal, Adhesion | Glioblastoma, prostate and breast cancer | Deletion | Cowden's |
| hMSH2 | DNA repair mechanism | Colon, uterine cancer, melanoma | Mutation | Hereditary non-polyposis colon cancer |
| BRCA1 | | Breast and ovarian cancer | Mutation | Hereditary breast and ovarian cancer |
| BRCA2 | | Breast and ovarian cancer | Mutation | Hereditary breast and ovarian cancer |
| WT-1 | Transcription factor | Wilms Tumour | Deletion | Wilms Tumour in children |
| NF-1 | GTP-ase activator | Sarcoma, glioma | Deletion | Neurofibromatosis |
| NF-2 | Cytoskeletal protein | Schwannom | Mutation | Neurofibromatosis |
| p16/CDKN2 | Cell cycle regulator | Melanoma, esophageal, pancreatic cancer | Deletion Methylation Mutation | Family melanoma |

Currently, it is assumed that out of 600 viruses known to be able to infect man, 150 have oncogenic potential. After penetration of the virus into a cell, it is processed and undergoes a series of changes: biological, morphological, the viral nucleic acid is multiplied by the formation of multiple copies of nucleic acid, each copy constituting the nucleic acid of a new virus – viral replication.(3) Oncogenes viruses in association with the mutations of pro-oncogenes, oncogenes and suppressor genes, have a role in carcinogenesis.(12)

Table no. 3. Types of human oncogenes viruses and types of cancer involved (22)

Oncogenes RNA viruses:

| Group and viral family | Cancers type |
|-----------------------------------|----------------------|
| Sarcomas virus | Rous Sarcoma |
| Chronic leukemia viruses HTLV1 | Leukemia or lymphoma |

| | |
|------------------------|-----------------------------------------------------|
| HTLV2 HTLV3 and HIV | with T cells Hairy cell chronic leukemia AIDS |
|------------------------|-----------------------------------------------------|

Table no. 4. Types of human oncogenes viruses and types of cancer involved (22)

Oncogenes DNA viruses:

| Group and viral family | Types of cancer |
|----------------------------------------------|------------------------------------------------------------------------------------|
| Papova A – HPV | Anogenital carcinoma |
| Papova B - polyoma - SV-50 - JCV - BCV | Lymphomas, Kapoṣy Sarcoma Leukemia in children Kapoṣy Sarcoma, Young Sarcoma |
| Adenoviruses | Adenoma |
| Herpesviruses (Epstei-Barr virus) | Nasopharynx cancer, Burkitt lymphoma, Leukemia |
| Hepadnaviruses - HBV | Hepatocellular cancer |

METABOLIC CONSEQUENCES AND PATHOGENETIC IMPLICATIONS OF NEOPLASTIC PROCESS

The main biochemical abnormalities of the oncogenesis process are: changes produced by initiators and tumour progression; decrease in specific activity of oxidases peroxidases;(12) the role of carbohydrate shown through glycan synthesis;(11) disturbance of antioxidant enzyme activities;(12) changes in growth factor activity;(11,21) disturbance of junctional communication; disturbance of cell adhesion and of substrate phenomena;(15) disturbance of topo-inhibiting and inhibition phenomena cell density-dependent.(14)

Some of the pathognomonic tumour growth factors involved in carcinogenesis are: epidermal growth factor (EGFR), tumor angiogenesis factor (TAF), platelet growth factor (PDGF). VEGF (vascular endothelial growth factor) together with EGFR and PDGF are pro-angiogenesis factors.(21) EGFR indicates the number of extracellular connections and, in cooperation with its counterpart HER2, points out the presence of proliferation and apoptosis.(13) Their amplification occurs in more than half of glioblastomas, breast cancer, and lung cancer.(22)

Conclusion: The appearance of the neoplastic process is based on a series of complex mechanisms that affect DNA structure and the expression of several genes, genetic and epigenetic mechanisms. The progresses made in the field of molecular mechanisms of carcinogenesis are multiple, and find their applicability in various fields of clinical oncology:(13) diagnosis (classification of tumours, differential diagnosis between benign and malignant forms, early detection);prognosis (amplifications of oncogenes); treatment (suppression of oncogenes transcription – monoclonal antibodies and radiotherapy); epidemiology (genetic consultation).

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