

Current predictive pathology in the clinical setting of colorectal cancer

Dear Editor,

Colorectal adenocarcinoma (CRC) is the second most common cancer in women and the third most common cancer in men, affecting mainly people older than 65 years¹. The highest incidence is observed in industrialized countries, but in the other countries the incidence is increasing. Consumption of alcohol, processed and red meat, and obesity, are demonstrated risk factors. Also, genetic predisposition plays a role, and some mutations are recognized risk factors. Furthermore, some genetic tumor syndromes were identified².

The efforts to find new targets brought to identify some predictive factors in CRC³. Two genes belong to MAPK pathway, RAS and B-RAF, HER2 is a transmembrane receptor implied in PI3K pathway, and microsatellites are four proteins which forming two heterodimers are designate to correct DNA errors. A recent study on VEGF-A and MMP-7 genes expression could assume a role in the next future if it will be confirmed their predictive role, but further studies are necessary⁴.

RAS belongs to MAPK pathway, regulating cell proliferation. RAS gene mutations are the most prevalent in CRCs, particularly mutations of K-RAS (Kirsten rat sarcoma viral oncogene homolog), followed by N-RAS and, only occasionally, H-RAS mutations.

RAF is part of MAPK signaling pathway; therefore, its activation leads to uncontrolled cellular proliferation and growth. B-RAF mutation occurs in codon 600, almost always V600E and seldom V600K, and is less frequent than RAS mutation in CRCs.

In 2017 it was established by a joint consensus of the American Society for Clinical Pathology, College of American Pathologists, Association for molecular Pathology, and American Society of Clinical Oncology to test these mutations in order to choose the proper therapy⁵. It is already proven that RAS mutations are predictive of resistance to anti-EGFR therapy in metastatic colorectal cancer, while testing B-RAF is recommended for prognostic stratification. In fact, patients with B-RAF V600E mutation are requested to take anti-EGFR antibodies therapy in combination with other pathway inhibitors^{6,7}.

HER2/neu (or erbB-2), human epidermal growth factor receptor², is a transmembrane tyrosine kinase receptor, located on chromosome 17. The dimerization of this receptor activates many signal pathways, including MAPK, PI3K/akt, STAT, regulating cell proliferation and survival. HER2 amplification has been demonstrated in many cancers, mainly in breast and gastrointestinal tract cancers. HER2 status is usually detected on formalin fixed or paraffin-embedded tissues, and has an approved scoring system for CRCs, that needs further confirmation with ISH or FISH^{8,9}. The possibility to treat cancer with specific immunotherapy gives a therapeutic option, although the prognostic value of HER2 in CRCs is still debated.

Microsatellite instability is a condition caused by a DNA mismatch repair defect that leads to a genetic hypermutability status. DNA mismatch repair is constituted of numerous proteins, responsible of recognizing and correct errors in DNA, which occurred during replication and recombination. The inherited DNA mismatch repair defect is implied in some human syndromes, while its epigenetic silencing can be found in sporadic cancers. The cancers with MSI are characterized by a high number of mutations^{10,11}. Microsatellite instability can be revealed

by immunohistochemical technics, determining the expression of four proteins, MSH2, MLH1, MSH6 and PMS2, which are involved in Lynch syndrome. About 75% of the CRCs cases are sporadic¹², but MSI detection is fundamental for cancer classification. The prognostic value is uncertain, but MSI could be a target for new therapies. Furthermore, recent studies are focusing on different forms of MSI expressed in other tumors¹³.

Conflict of Interest

The Authors declare that they have no conflict of interests.

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