Variances in the Expression of mRNAs Related to the Histaminergic System in Endometrioid Endometrial Cancer

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Research has indicated higher concentrations of histamine and polyamine in endometrioid tissue in comparison with healthy tissue. The aim of this study was to evaluate changes in the expression patterns of messenger RNA (mRNAs) related to the histaminergic system in endometrial samples and whole blood in women with endometrioid endometrial cancer. The study group consisted of 30 women with endometrioid endometrial cancer qualified for hysterectomy (G1 well-differentiated, 15 cases; G2 moderately differentiated, 8 cases; and G3 poorly differentiated, 7 cases). The control group included 30 women with no neoplastic changes during routine gynecological examinations. The molecular analysis consisted of the microarray analysis of mRNAs related to the histaminergic system, and reverse-transcription quantitative polymerase chain reaction (RTqPCR) Out of 65 mRNAs connected with the histaminergic system, 10 differentiate the samples of tissue and blood obtained from patients with endometrial endometrial cancer in comparison with the control group (p < 0.05). mRNA histamine receptor 1.3 (HRH1, HRH3), and solute carrier family 22 member 3 (SLC23A2) differentiating samples of endometrioid endometrial cancer independent of either G or control. The selected mRNA transcripts seem to be promising for molecularly targeted therapies in the context of endometrioid endometrial cancer.

Keywords: endometrioid endometrial cancer, grading, histaminergic system, molecular marker