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Microsatellite Instability (MSI) Analysis in Patients with Endometrial Cancer*

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Microsatellite instability (MSI) seems to be important for the development of various human cancers including sporadic endometrial cancer. The aim of this study was evaluation of microsatellite instability in 20 postmenopausal women with endometrial adenocarcinoma in DNA samples obtained from cancer tissue and blood of the same patients. Control DNA was obtained from normal endometrial tissue (n=25). MSI was studied at five loci containing single- or dinucleotide repeat sequences and mapping to different chromosomal locations: BAT-25 (at locus 4q12), BAT-26 (2p16), D2S123 D5S346 D17S250 (2p16-p21), (5q21-q22)and (17q11.2-q12). No differences in the MSI frequencies between blood and cancer tissue obtained from patients were detected. The microsatellite instability status was significantly higher in endometrial cancer tissue [5/20 (25%)] as compared to control [3/25 (12%)] (p<0.05). There were no significant differences between MSI presence in the subgroups assigned to the histological grades (p>0.05). The results suggest that the microsatellite instability seems to be important in the development of sporadic endometrial cancer.

Introduction

Microsatellite sequences are short repeat nucleotide sequences disseminated in whole genome in normal conditions. In *Eucaryota* genome there are repeated sequences consisting of one, two, three and four nucleotides. Over 90% of the till this moment studied microsatellite sequences from mononucleotides to tetranucleotides show polymorphism. Small deletions or expansions in tumor DNA, manifested as shifts in allelic electrophoresis mobility characterize microsatellite instability (MSI). Genetic instability is considered to be responsible for a rapid accumulation of somatic mutations in various tumor suppressor genes and oncogenes, thus playing an important role in the initiation and progression of malignant tumors [16, 17].

Previous studies have indicated that MSI seems to be important in the development of various human cancers [2, 25, 33] including sporadic endometrial cancer (EC) [7, 9, 21, 31].

Endometrial cancer is one of the most common malignant neoplasms, which appear in uterine body [26]. About 80% of the cases are diagnosed after menopause. The highest incidence estimated in 57–58 years is moving to 6 and 7 decade of life at present [4]. Endometrial cancer is fourth the most common female carcinoma [29]. Annually 150,000 new cases of this cancer are noted worldwide. Every year 65 new cases of endometrial cancer in age group 65–75 years are diagnosed among every 100,000 women.

Some risk factors have been identified, related to reproduction (such as early age at menarche, late age at menopause and nulliparity) or more immediately estrogen-related (i.e. conditions such as the polycystic ovarian syndrome) [4, 30].

Prognostic factors important in the evaluation of EC are: tumor size, tumor grade, presence or absence and depth of myometrial invasion, myometrial lymphatic/vascular space invasion, steroid hormone receptor status, ploidy and proliferative indices. However, these factors present an incomplete picture of the tumor biology. Therefore, investigation of other prognostic factors is of special clinical relevance, particularly in view of the unexpectedly progressive course of the disease and frequent relapses in some cases.

MSI is characterized by small insertions or deletions within short tandem repeats in tumor DNA when compared to the corresponding normal DNA. MSI was firstly demonstrated in patient with hereditary nonpolyposis colorectal carcinoma (HNPCC), an inherited cancer syndrome that

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predisposes also to endometrial cancer, which represents the second most common malignancy in HNPCC families [14, 24]. Later studies have also demonstrated that MSI is present in about 20% of sporadic colorectal tumors [20]. Endometrial cancer is one of the most common extracolonic tumors associated with HNPCC, and MSI has been reported to be present in 9–45% of sporadic cases [5, 6, 12, 19, 27].

In the present study MSI status in women with sporadic endometrial cancer was investigated.

Material and Methods

Endometrial cancer samples

Twenty patients with histologically-proven diagnosis of endometrial adenocarcinoma were included in the study (mean age \pm SD – 63.75 \pm 4.72 years). Endometrial cancer was classified as sporadic after evaluation of patient's family history in order to exclude the presence of the HNPCC syndrome. Family history for cancer was evaluated by questionnaire interviews in endometrial cancer patients attending the Department of Obstetrics and Gynecology of the Medical University in Łódź after the initial surgical treatment. Tumor tissues and blood were obtained from postmenopausal women with endometrial adenocarcinoma treated at Department of Obstetrics and Gynecology between 2002 and 2004. Tissues were frozen immediately and stored at -70°C. All tumors were staged according to the criteria of the International Federation of Gynecology and Obstetrics (FIGO). There were 7 tumors of I stage, 6 of II stage and 7 of III stage in total. DNA from normal endometrial tissue (n=25) served as a control.

DNA isolation

DNA was extracted from fresh material using commercially available QIAmp Kit (Qiagen GmbH, Hilden, Germany) DNA purification kit according to manufacturer's instruction.

Microsatellite analysis

Tumor DNA and corresponding normal DNA were analyzed using a panel of five microsatellite markers for mononucleotide and dinucleotide repeat sequences: BAT25 (at locus 4q12), BAT26 (2p16), D2S123 (2p16-p21), D5S346 (5q21-q22) and D17S250 (17q11.2-q12) [3]. All primer sequences were as reported in Genome DataBase (GDB, at: http://www.gdb.org).

The PCR was carried out in a Perkin-Elmer/Gene Amp, PCR System 2400 thermal cycler. The thermal cycling conditions were 60s at 94°C, 60s at 60°C, 60s at 72°C, repeated for 30 step cycles. PCR amplification was performed in a final volume of 25 μ l. The reaction mixture contained 5 ng of genomic DNA, 0.2 μ mol of each appropriate primer (ARK Scientific GmbH Biosystems, Darmstad, Germany), 2.5 mM MgCl₂, 1 mM dNTPs and 1 unit of Taq Polymerase (Qiagen GmbH, Hilden, Germany). PCR products were fractionated by denaturing electrophoresis in a 6% polyacrylamide gel (PAGE) and visualized by silver staining.

Statistical analysis

For statistical analysis, the χ^2 test was performed, p<0.05 was considered significant.

Results

A sample was classified as MSI-high (MSI-H) if two or more markers showed instability, MS-stable (MSS) if no instability was noted, and MSI-low (MSI-L) if a single marker revealed novel bands compared with the corresponding normal DNA.

TABLE 1

Number of endometrial cancer patients and control with the presence or absence of microsatellite instability (MSI)

MSI status	Endometrial cancer patients (n=20)		Control (n=25)	
	Number	Frequency	Number	Frequency
MSI+	5	0.25*	3	0.12*
MSI-H	4	0.20	1	0.04
MSI-L	1	0.05	2	0.08
MSS	15	0.75	22	0.88

*p<0.05 as compared to controls



Fig. 1. Number of endometrial cancer patients (n=20) presenting or not microsatellite instability in relation to tumor grade.

MSI was determined in 20 endometrial carcinoma tissues and blood from patients and in 25 control samples. There were no differences in MSI presence between blood and cancer tissue obtained from the same endometrial cancer patients.

Five out of 20 (25%) tumors tested were found to be MSI positive, 4 MSI-high and 1 MSI-low (Table 1). It can be seen from the Table 1 that there were significant differences in MSI frequency between women with endometrial cancer and control. The frequency of MSI presence in cancer samples was higher than in normal samples (p<0.05).

A dependency of the MSI frequency distribution on the tumor grade evaluated according to FIGO criteria in women with endometrial cancer is displayed in Figure 1. The histological analysis of tumor grade showed a lack of association (p>0.05) between tumor grade and microsatellite instability presence.

Discussion

Endometrial tumorigenesis is still poorly understood. The development of endometrial cancer is associated with an accumulation of specific genetic alterations [22]. The activation of oncogenes, the loss or inactivation of repressor genes and impaired mismatch-repair function are known to be involved in the development of tumors. It is well known, that MSI plays an important role in the pathogenesis and disease progression of endometrial cancer [1, 11, 15, 23, 18].

Many studies have documented that MSI may be a marker of a tendency for replication errors in human cancers [13, 23, 31]. Defects in DNA mismatch-repair genes such as MLH1, MSH2 and MSH6 lead to replication errors revealed as instability in microsatellite markers. Several studies have indicated that loss of MLH1 expression can account for a large proportion of sporadic endometrial tumors [8, 10, 28]. In the majority of endometrial tumors with high MSI the loss of MLH1 expression was found, whereas loss of expression was less frequent in tumors with intermediate MSI. Thus, pathological expression of MLH1 does not seem to account for all tumors with a MSI-positive phenotype, indicating that other mismatch repair genes might also be involved [28].

In the light of substantial evidence that the progression of endometrial cancer can be associated with microsatellite instability, it seems reasonable to check a possible correlation between MSI and clinical status of endometrial cancer patients. The patients included in the study received no chemotherapy or hormone therapy. In this work conducted on 20 endometrial carcinoma patients we find a correlation between MSI and occurrence of cancer. Twenty-five per cent of endometrial cancer patients presented MSI, while only 3 normal samples from 25 healthy individuals (12%) showed this pattern. However, the histological analysis of tumor grade showed a lack of correlation between tumor grade and the number of patients presenting MSI.

Recent studies have suggested that genetic alterations, including *p53* mutations, loss of heterozygosity, and *K-ras*, *PTEN* and *Her2/neu* gene amplification, can be detected in endometrial cancer [17]. Our findings provide additional evidence that genetic alterations, including MSI, may occur as relatively early events in the development of endometrial cancer.

The implications of MSI in DNA from patients with endometrial cancer may be associated with different tumorigenic pathways. The genome-wide MSI may be correlated to the existence of pathogenetic mechanisms inducing progressive accumulation of sequence errors and providing a selective advantage during malignant evolution.

Our study implies that it is possible that the MSI process may be involved in the appearance and/or progression of endometrial cancer. Further studies, conducted on a larger population, are required to clarify this point.

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