Association of LCT-13910 C/T Polymorphism and Colorectal Cancer

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Purpose: The activity of epithelial lactase (LCT) is associated with a polymorphism 13910 bp upstream in the lactase encoding gene. Because the association between the LCT-13910 polymorphism and the risk for colorectal cancer is not clear, we investigated the role of the LCT-13910 polymorphism as a potential risk factor for colorectal cancer and colorectal polyps in the Turkish population.

Methods: One hundred sixty-six subjects (74 with polyps, 44 with colorectal cancer, 48 controls), who had undergone a total colonoscopy between January 2012 and November 2012 in our endoscopy unit were genotyped for the LCT-13910 polymorphism by using the polymerase chain reaction and minisequencing.

Results: The CC genotype in the lactose gene 13910 locus, which is accepted as the genetic indicator of lactase deficiency, was determined as 83.7%. The CC genotype rate was determined as 89.1% in patients who had a history of lactose intolerance and 81.5% in those without a history of lactose intolerance (P = 0.236). No difference was detected between the patients who had colorectal polyp(s) and/or cancer and the controls with regard to the LCT-13910 polymorphism. No differences were determined between groups when they were compared with regard to the C or the T allele.

Conclusion: No differences were detected between the patients who had colorectal polyp(s) and/or cancer and those with normal colonoscopy findings with regard to lactase gene polymorphisms. No differences were determined between the groups when they were compared with regard to the C or the T allele.

Keywords: Adenocarcinoma; Colorectal neoplasms; LCT gene; Lactose intolerance

INTRODUCTION

Colorectal cancer (CRC) is the third most common cancer in the world. The effect of milk and dairy products on CRC is controversial [1]. Several cohort studies have shown that consumption of milk reduces the risk of CRC. This positive effect is known to be due to the components of milk, including calcium, vitamin D, lactoferrin, and conjugated linoleic acid. In the epithelial cells of the intestine, calcium and vitamin D regulate cell growth and promote cell differentiation by stimulating calcium-sensing receptors.

Also, in the intestinal lumen, calcium may bind complex fatty acids and secondary bile acids, thereby reducing their cytotoxic effects [2-4].

Lactose is the main sugar of milk, and lactose intolerance (LI) is common. Symptoms of LI include diarrhea, abdominal pain, and flatulence after drinking or eating milk or products that contains milk. These symptoms are caused by low levels of small intestinal lactase due to mucosal injury or, more often, due to reduced genetic expression of the enzyme lactase-phlorizin hydrolase. Lactase deficiency is defined as an intestinal brush border lactase enzyme activity that is lower than that of normal infants. Lactose malabsorption means the failure of the small bowel to absorb a sizable fraction of ingested lactose. LI is a clinical syndrome that includes many symptoms (e.g., abdominal pain, bloating, flatulence, and diarrhea) due to lactose malabsorption [5]. Lactose malabsorbers may consume small amounts of milk and dairy products, which may result in low calcium intake, a risk factor for the development of CRC.

The lactase gene codes an enzyme called lactase-phlorizin hydrolase that helps the digestion of milk. The level of lactase en-
zyme remains high until weaning, after which it falls to lower levels in the adult population where it is called adult-type hypolactasia. In some populations like northern Europe, high levels of lactase enzyme are maintained in adulthood, which is called lactase persistence (LP) [6]. Multiple single nucleotide polymorphisms (SNPs) in the DNA sequence of the coding region and the regulatory region of the lactase gene on 2q21 have been identified. The SNP LCT-13910C > T (rs4988235), located in the minichromosome maintenance-6 gene upstream from the LCT-encoding gene, was found to be associated with adult-type hypolactasia. Genetic analysis has established that LP in adulthood is inherited as an autosomal dominant trait. The presence of the T allele of the SNP located at -13.9-kb upstream of the lactase gene has been strongly correlated with LP. The CC genotype defines lactose malabsorbers; CT and TT genotypes define lactose absorbers [7-12]. Recent studies showed a significant association between the CC genotype and an elevated CRC risk among members of the Finnish population, but it was not confirmed in British, Spanish, Hungarian, or Italian patients [13-15]. For those reasons, in this study, we aimed to investigate the role of LCT-13910 polymorphisms as a potential risk factor for CRC and colorectal polyps in the Turkish population.

METHODS

We performed a total colonoscopy on all patients between January 2012 and November 2012 in our endoscopy unit and screened all of them for LCT-13910 polymorphisms. Inclusion criteria were Turkish ethnicity, age > 25 years, and informed consent to be interviewed. A partial colonoscopy, inflammatory bowel disease, familial polyposis, non-CRC-related previous bowel surgery, history of a neoplastic disease other than CRC or adenoma served as exclusion criteria. Seventy-seven of 92 patients with adenomatous polyps, 1 with a colon carcinoma, and in 2 controls; then, 74, 44, and 48 patients were analyzed in the polyp, cancer, and control groups, respectively. Approval for this research was obtained from the local Ethics Committee.

DNA was extracted from mononuclear cells of the peripheral blood. We used Invitrogen Blood DNA Isolation Kits (Lot#: 1388070, Purelink Genomic DNA Kit, Life Technologies, Carmaillo, CA, USA) for extraction of DNA. Under the conditions recommended by the manufacturer (95°C for 10 minutes to activate the AmpliTaq Gold enzyme, followed by 40 cycles of 15 seconds at 95°C and 60 seconds at 60°C), the 25-µL polymerase chain reaction, with 2x TaqMan Genotyping Master Mix (Applied Biosystems, Waltham, MA, USA), 20x probe and primers assay mix, and 5 ng of genomic DNA, was performed in 96-well plates in a real time-PCR System (7900 HT, Applied Biosystems). Both primers and probes were obtained from Applied Biosystems. The data were collected and analyzed using the SPSS ver. 15.0 (SPSS Inc., Chicago, IL, USA). The associations of age, sex, job, smoking, alcohol use, history of LI, presence of diabetes, marital status, the LCT-13910 genotype (CC, CT, TT), and allele frequency with colon cancer, colon polyps or normal colonoscopy were investigated. The categorical variables of colon cancer were compared using the chi-square test, and the continuous variables were analyzed using the analysis of variance test or the t-test. The patients with colon cancer and colon polyp were gathered in a group as colon neoplasms, and the associations were investigated.

RESULTS

One hundred sixty-six patients were included in the study (male, 62%; mean age, 60.7 ± 10.4 years). No differences with regard to age, presence of diabetes, alcohol use, smoking, and history of LI were found between the patients who had an adenomatous polypl...
or cancer and those whose colon was normal (Table 1). The percent of the male sex was lower in the controls. When patients with polyps and with cancer were approached as a single group and compared with the controls, the percents of the male sex (71.2%, P < 0.0001) and of alcohol use (21.2%, P = 0.048) were found to be higher in the neoplasia group.

Although a history consistent with LI was seen at a rate of 27.7%, the CC genotype in the lactose gene 13910 locus, which was accepted as the genetic indicator of lactase deficiency, was determined as 83.7%. The CC genotype rates were determined as 89.1% in patients who had a history of LI and 81.5% in those without a history of LI (P = 0.236). No differences with regard to lactase gene polymorphisms were detected between the patients who had a colorectal polyp or cancer and those with normal colonoscopy findings. No differences were found between the groups when they were compared with regard to the C or the T allele.

**DISCUSSION**

The main mechanism of lactose malabsorption in healthy individuals is a genetically regulated reduction in the lactase enzyme activity, which is caused by racial or ethnic factors. Most of the world's populations have low intestinal lactase levels during childhood. This characteristic is most common in the Asian and the African populations; in contrast, most Caucasians, particularly of northern European background, have elevated lactase activity into adulthood. LP has also been observed in populations that historically domesticated cows or other milk providers and consumed products of milk into adulthood. Evidence exists for a convergent evolution of LP in some of these groups. LP in adulthood is inherited as an autosomal dominant trait, but the molecular basis is not sufficiently clear [16-18].

Genetic polymorphisms were found to be related with lactase deficiency and are being widely used to determine that condition. In this study, a CT polymorphism located 13.9-kb upstream of exon 1 of the lactase gene, which is the most commonly polymorphism used worldwide and shows about 90% concordance with lactase deficiency in the white population, was used. This polymorphism is closely related with LP in the Finnish population, where it was first determined, and in the German, English, Italian, and even Indian populations [19-21]. However, no relationship could be found between lactase deficiency and the 13910 CT polymorphism in non-White populations such as Africans, the Chinese, and the Japanese [21-25]. The relationship between lactase deficiency and the 13910 CT polymorphism in the Turkish population is not known. In 2 studies in the Turkish population, the rates of lactase deficiency diagnosed with the H2 breath test, the lactose tolerance test, and the urine galactose/creatinine ratio were found to fall in the range from 71% to 85% [26, 27]. LP seems to be 6% according to the Global Lactase persistence Association Database. Namely, lactase deficiency is seen in 94% of the adult population [28].

We investigated if differences with regard to lactose gene polymorphism could be found between the 44 patients with CRC, the 74 patients with an adenomatous polyp of the colon, and the 48 patients with normal colonoscopy findings. The 13910 CC genotype, which was found to be related with lactose nonpersistence, was identified in more than 80% of the patients in each subgroup, and no differences were found between the groups. No differences were found between the groups when they were compared with regard to the C or the T allele. The CC genotype in the lactose gene 13910 locus, which was accepted as the genetic indicator of lactase deficiency, was determined as 83.7%. The CC genotype rates were determined as 89.1% in patients who had a history of LI and 81.5% in those without a history of LI (P = 0.236).

A literature search about ‘LCT-13910 polymorphisms and colorectal cancer,’ revealed a few studies. Firstly, Rasinperä et al. [13] found that the CC genotype occurred more frequently in the patient group in the Finnish population (CRC patients, 23.5%; controls, 18%), but in the same study, these findings were not supported in the British (CRC patients, 9.2%; controls, 8.8%) and the Spanish (CRC patients, 31.9%; controls, 36.7%) populations. Among Hungarians, the prevalence of the CC genotype was studied by Bácsi et al. [14]; they found the frequency of the CC genotype to be higher in the Hungarian population (patients, 41%; controls, 36%) than in the Finnish population, but they found no statistically significant association between CRC and the CC genotype. In the Italian population, Tarabara et al. [15] reported that the LCT-13910C > T polymorphism was not associated with an increased risk for CRC or polyps (CC genotype: CRC patients, 62.09%; patients with polyp, 64.77%; controls, 63.02%). Our results were compatible with those found in studies done on the Italian and the Hungarian populations (CC genotype: CRC patients, 86.4%; patients with polyp, 82%; controls, 83%).

Our study does have some limitations. Firstly, the small number of the subjects in this study may have hindered the demonstration of the relationship between the LCT-13910 polymorphism and an increased risk for colorectal polyps and cancer. We did not use a method such as lactase expression in biopsy to show directly lactase deficiency; this could have masked a real relationship. According to the results of this study, the LCT-13910 polymorphisms (CC/CT/TT) are not associated with an increased risk for development of CRC in the study population. Further studies including larger populations are needed to confirm our findings.

**CONFLICT OF INTEREST**

No potential conflict of interest relevant to this article was reported.

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