Expression of Cytoplasmic 8-oxo-Gsn and MTH1 Correlates with Pathological Grading in Human Gastric Cancer

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Abstract

Background: Cancers have dysfunctional redox regulation resulting in production of reactive oxygen species (ROS), damaging DNA, RNA and free NTPs, and causing the accumulation of oxidative nucleic acids in cytoplasm. The major types are 8-oxo-7,8-dihydroguanine (8-oxoGsn) in RNA and 8-oxo-7,8-dihydro-2’ deoxyguanosine (8-oxodGsn) in Mt-DNA. The MTH1 protein sanitizes oxidized nucleotide pools from NTPs to monophosphates, preventing the occurrence of transversion mutations. This study concerned cytoplasmic 8-oxodGsn/Gsn and MTH1 expression in gastric cancer and para-cancer tissues and elucidated roles of nucleic-acid oxidation and anti-oxidation. Materials and Methods: A polymer HRP detection system was used to detect 8-oxo-Gsn/dGsn and MTH1 expression in 51 gastric cancer and para-cancer tissue samples. Analyses of patient clinical and pathological data were also performed. Results: The expression of MTH1 and the 8-oxo-dGsn/Gsn ratio were significantly higher in cancer tissues than para-cancer tissues (P<0.05). Cytoplasmic 8-oxo-Gsn and MTH1 were both found to positively correlate (P<0.05) with tumor differentiation, while no significant associations were found with gender, age, invasion depth, lymph node metastasis and clinical stage (P>0.05). Conclusions: We found 8-oxo-dGsn/Gsn and MTH1 are both highly expressed in gastric cancer tissues, especially in well differentiated lesions. In addition, oxidated mtDNA is prevalently expressed in gastric cancers, while 8-oxo-Gsn expression in cytoplasmic RNA is a bit lower, but more selectively.

Keywords: Gastric cancer - ROS - MTH1 - nucleic acids - oxidation

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Introduction

The accumulation of oxidative damage in nucleic acids is one of the major causes for mutagenesis and cell death (Nakabeppu et al., 2004). Among the cytoplasmic oxidative nucleic acids, 8-oxo-7, 8-dihydroguanine (8-oxoGsn) of RNA and 8-oxo-7, 8-dihydro-2’ deoxyguanosine (8-oxodGsn) of Mt-DNA are the major types, while oxidative NTP pools, such as 8-oxo-dGTP, which is a product of dGTP oxidation and can be inserted into opposite dA or dC residues of template DNA at almost identical efficiencies, causing G:C to T:A or T:A to G:C transversion mutations. While, MTH1 is an oxidized purine nucleoside triphosphatase that upregulates to prevent cellular DNA or RNA oxidative damage, when 8-oxodGsn/Gsn increase (Shimura-Miura et al., 1999, Kosuke et al., 2006). Recently, Helge et al. (2014) and Huber et al. (2014) reported in nature that cancer cells required MTH1 activity to avoid incorporation of oxidized NTPs, resulting in nucleic-acid damage and cell death.

Overexpression of MTH1 has been reported (Okamoto et al., 1996; Kennedy et al., 2003; Koketsu et al., 2004) in various tumors except gastric cancer. To explore the correlation between oxidation, MTH1 expression and human gastric adenocarcinoma, we investigated the expression pattern of MTH1 and 8-oxoGsn/dGsn in this form of cancer. To the best of our knowledge, this is the first report describing the expression of MTH1 and 8-oxoG in gastric cancer.

Materials and Methods

Clinical data collection

In this study, 51 cases were selected. The patients had been confirmed histologically gastric cancer and received cancer radical operation at the second affiliated hospital of Wenzhou Medical University, during 2009-2013. Histological stages were graded according to the TNM classification (UICC/AJCC 2010).

Experimental reagents

Polyclonal rabbit anti-human MTH1 antibody (dilution 1:200) and monoclonal mouse Anti-8 Hydroxyguanosine antibody (dilution 1:300) were both purchased from Abcam. The corresponding secondary donkey polymer immunohistochemical staining hypersensitivity kit anti-mouse/rabbit antibody were obtained from ZSGB-BIO. (Beijing, China).
Immunohistochemistry experimental procedures.

Resected tumor tissue of all cases were fixed in 4% formaldehyde solution (pH 7.0) for periods not exceeding 24 h. The tissues were processed routinely for paraffin embedding, and 3µl thick sections were cut within a cryostat and placed on cationic anti-slip glass slides (ZSGB BIO., Beijing, China) for immunohistochemistry. Tissue samples were stained with hematoxylin and eosin to determine histological type and grade of tumors.

For the detection of 8-oxo-dGsn and 8-oxoGsn (Zheng et al., 2009; Song et al., 2011). The sections were de-waxed using xylene and gradually dehydrated with gradient ethanol and washed with PBS. Antigen retrieval was performed by microwave at 95°C for 15min with buffer solution of citrate salts (pH 6.0) (Maixin Bio., Beijing, China). For the detection of 8-oxo-dGsn in mitochondria, 5mg/ml RNase A (10mg/ml, TakaRa Biotechnology Co., Ltd, Dalian, China) treated sections were directly subjected to IHC with the 15A3 mAb. While to detect the 8-oxo-Gsn in RNA, the sections were treated with 3N HCl at room temperature (RT) for 30min, thus denaturing the nuclear and mitochondrion DNA, then the sections were treated with 50mM Tris-Base for 5 min, furthermore, the sections were treated with DNase I (1000U/ml, TakaRa Biotechnology Co., Ltd, Dalian, China) to remove nuclear or mitochondria DNA, then subjected to IHC with the 15A3 mAb. Besides, the MTH1 staining was performed according to the instructions provided by manufacturer.

The staining intensity of the cells was graded as follows: 0, negative; 1, light yellow granules; 2, deep yellow granules; and 3, brown granules points. Grading was performed as follows: 0, positive cell ratio ≤5%; 1, 5% < positive cell ratio ≤25%; 2, 25% < positive cell ratio ≤50%; 3, 50% < positive cell ratio ≤75%; and 4, 75%< positive cell ratio ≤100% points. The final points score for expression was determined through the multiplication of the points for staining intensity by the points for positive cell ratio. The expression levels were graded as follows: -, points ≤4; +, 4 < points ≤8; ++, 8 <points ≤12; and ++++, points >12. Points less than 4 was determined as MTH1 and 8-oxo-dGsn negative, whereas +, ++ and +++ were determined as positive. The expression of 8-oxo-Gsn in RNA, was defined as points <3 as negative and ≥3 positive.

Statistical analyses

Data analyses were performed using SPSS software, version 20.0 (SPSS Inc., Chicago, IL, USA). The χ² test was used to evaluate the MTH1 and 8-oxo-dGsn/Gsn expression differences between gastric cancer and para-cancer tissues. The association between MTH1 or 8-oxo-dGsn/Gsn expression and other clinicopathological parameters, including gender, age, clinical stage, histological grade, lymph node metastasis, tumor size and depth of invasion were also analyzed. P values of less than 0.05 were considered statistically significant.

Results

Patient characteristics

The mean age of the 51 patients studied was 63.27 years (range 42-87), including 34 male and 17 female. No significant difference was found in age and gender (P=0.43). Within the total sample, 11 gastric cancer are determined as pathological grade I, 13 as grade II, 5 as grade II-III, 13 as grade III, and 8 grade IV.

Assessment of MTH1 and 8-oxo-dGsn/Gsn expression

Of the 51 gastric cancer cases, 24 (47.1%) exhibited positive MTH1 expression (mainly in the cytoplasm) and 27 (52.9%) exhibited negative MTH1 expression. In the corresponding para-cancer tissue samples, only 6 (11.7%) exhibited positive MTH1 expression. The difference in

Figure 1. MTH1 Expression in Gastric Cancer and Para-Cancer Tissues Detected by Using Polymer Immunohistochemical Staining (Magnification, x200). (A) 8-oxo-dG staining in Moderately differentiated gastric cancer. (B) Negative expression of 8-oxo-Gsn. (C) Positive expression of MTH1. (D)(E)(F) are negative expression of para-cancer tissues of 8-oxo-dG/G and MTH1, respectively.
MTH1 expression between gastric cancer and para-cancer tissues was found to be statistically significant (P<0.05).

As for mitochondrial 8-oxo-dGsn expression, 27 (52.9%) gastric cancer cases exhibited positive expression, and 24 (47.1%) negative. While 12 (19.6%) para-cancer tissue samples exhibited positive 8-oxo-dGsn expression, statistically significant difference (P<0.05) was found between cancer and para-tissue (Table 1).

The similarity was seen in 8-oxo-Gsn expression, 16 (31.4%) cases exhibited positive 8-oxo-Gsn expression, and 35 (68.6%) negative. While 7 (13.7%) para-cancer tissue samples exhibited positive 8-oxo-Gsn expression, statistically significant difference (P<0.05) was found between cancer and para-tissue. The results are shown in Figure 1 and Table 1.

Association between MTH1, 8-oxo-dG/G Expression with other clinicopathological features

There was no significant association between MTH1 or 8-oxo-dGsn/Gsn expression and gender, age, tumor size, tumor location, tumor invasion depth, clinical stage, as well as lymph node metastasis (P>0.05). However, both MTH1 and 8-oxo-Gsn expression exhibited a significant positive correlation with pathological grade (P<0.05). The results are presented in Table 2.

Discussion

Accumulation of ROS is a double-edged sword. On one hand, ROS has been proved to be an endogenous class of carcinogens in the past decades (Guyton et al., 1993; Cerutti et al., 1994; Feig et al., 1994), on another, excessive ROS might induce cancer cells apoptosis, necrosis, or autophagy (Green et al., 1998; Golstein et al., 1993; Cerutti et al., 1994; Feig et al., 1994), on another, excessive ROS might induce cancer cells apoptosis, necrosis, or autophagy (Green et al., 1998; Golstein et al., 1993; Cerutti et al., 1994; Feig et al., 1994). ROS may also be inserted into opposite dA or dC residues of template DNA at almost identical efficiencies, so is the same as 8-oxo-GTP. As a result, G:C to T:A or T:A to G:C transversion mutations occur (Shibutani et al., 1991, Maki et al., 1992; Cai et al., 1997). As to mtDNA, which is more susceptible to mutations or damage induced by high level of ROS production in mitochondria of cancer cells, due to limited repair mechanisms compared to nuclear DNA (Zhu et al., 2005; Saffran et al., 2006; Trachootham et al., 2006; Clay Montier et al., 2009). And now it is clear that mtDNA mutation influences OXPHOS function which associated with tumorogenesis (Woo et al., 2012; Yadav et al., 2013).

In our study, 52.9% (27/51) gastric cancer samples show positive expression of 8-oxo-dGsn in mtDNA, besides, this expression was found to be significantly higher in gastric cancer tissues compared to para-cancer tissues. The positive expression of 8-oxo-dGsn in gastric cancer tissues suggests a role for this oxidative damage in the development and progression of gastric cancer. Further studies are needed to elucidate the biological significance of 8-oxo-dGsn in gastric cancer, and to explore potential therapeutic strategies for targeting this pathway.
no significant difference was found with other parameters, which means that oxidated mtDNA is prevalent among the whole gastric cancer, meanwhile, it is also a metabolic dysfunction signal of cancer cells, or even for normal cells. However, 8-oxo-Gsn in cytoplasmic RNA expression is a bit lower, but more selectively. 8-oxo-Gsn is positive expressed in 31.4% (16/51) of all the gastric cancer samples. In pathological grade I and II cases, the expression ratio are 6/11 and 6/14, while, in grade II-III, III, IV are 1/5, 2/13, 1/8, respectively. Which suggesting that better-differentiated gastric cancer specimen show higher expression of 8-oxo-Gsn in RNA than poorly-differentiated specimen. Interestingly, this just in accordance with the trendency of MTH1 expression, which, was expressed in grade I, II, II-III, III and IV, by 8/11, 8/14, 1/5, 4/13, 3/8, respectively.

The MTH1 protein sanitizes both oxidized dNTP and NTP pools, for example, by converting 8-oxoGTP to monophosphate, thereby preventing incorporation of 8-oxo-7, 8-dihydroguanine into RNA or 8-hydroxy-2-deoxyguanosine into DNA. Recently, Helge et al. (2014) and Huber et al. (2014) reported on Nature that, MTH1 is required for cancer survival, suggesting anticancer strategy by inhibiting MTH1 protein, thus, resulting in excessive oxidative damage of cancer necleic acid, causing a considerable decrease of cancer survival. Our study revealed the high expression of MTH1 protein, which just accorded with the reports of Helge Gad and Huber, moreover, we found MTH1 selectively high expressed in well differentiated gastric cancer.

Therefore, We speculated that the increased ROS level of gastric cancer mainly originated from its dysfunctional mitochondria. The level of ROS might dramatically relevant with mitochondrial capacity (quantity, size and maturity level of mitochondria), and oxidized RNA might produce abnormal protein that, with the dysfunctional mitochondria together, influence cancer differentiation and cancer behavior. However, MTH1 is a momentous protein for cancer cells defending oxidative damage. Thus, a novel anticancer strategy could be actualized through enhancing cancer oxidative stress and inhibiting MTH1 protein, simultaneously.

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