Identification of chosen apoptotic (TIAR and TIA-1) markers expression in thyroid tissues from adolescents with immune and non-immune thyroid diseases

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Abstract: The aim of this study was to estimate sodium iodide symporter (NIS) and thyroid peroxidase (TPO) expression in thyrocytes from patients with GD and no-toxic multinodular goitre (NTMG) in relationship with apoptotic (TIAR and TIA-1) markers. The investigation was performed on thyroid cells isolated from postoperation thyroid tissues from 15 patients aged 12-21 years old with GD and 15 cases aged 13-21 years old with NTMG. Detection of NIS and TPO was performed by immunohistochemistry. Analysis of apoptotic markers in thyroid tissues was performed using antibodies to TIAR and TIA-1 by Western Blot and immunohistochemistry. Identification of proapoptotic TIAR and TIA-1 molecules in the thyroid tissues revealed a higher expression of both proteins in patients with Graves’ disease (+++; +, respectively) in comparison to patients with NTMG (+; 0). In addition, TIAR expression was detected in three bands [p50, p42, p38 (kDa)] and TIA-1 in two bands [p22, p17 (kDa)], using Western Blot test in patients with thyroid autoimmune diseases. In patients with NTMG expression of both apoptotic proteins was lower and identified in single bands: 42 (kDa) for TIAR and 17 (kDa) for TIA-1. The analysis of expression of NIS and TPO in thyroid follicular cells was higher in patients with Graves’ disease in compared to their detection in patients with NTMG. In addition, degree of thyroid antigen expression positive correlated with amount of proapoptotic markers (TIAR, p<0.001; TIA-1, p=0.025 for NIS; TIAR, p=0.012 for TPO). We conclude that elevated expression of NIS and TPO in Graves’ disease is associated with higher stimulation and activation of apoptosis in thyroid follicular cells during autoimmune process.

Key words: thyrocytes, apoptosis, Graves’ disease, TIAR

Introduction

The course of Graves’ disease (GD) is associated with the inflow of lymphocytes to the thyroid gland and dysregulation of the immune system characterized by reaction to thyroid antigens (peroxidase, thyroglobulin, TSH receptors and Na+/I- symporter). After activation they shift to the inflamed thyroid gland, thus leading to the production of cytokines which can stimulate activity of thyrocytes and increase expression on intracellular proapoptotic markers such as TIAR and TIA-1.

T-cell intracellular antigen 1 (TIA-1) and TIA-1-related protein (TIAR) are the RNA-binding proteins. They consist of three RNA recognition motifs (RRMs) and a glutamine-rich carboxyl-terminal domain [1].

Both proteins play role in nuclear and cytoplasmic RNA metabolism, in pre-mRNA splicing and mRNA translation. TIA-1 has been identified as an important splicing regulator in mammals. It was proven that TIA-1 regulates the alternative pre-mRNA splicing of various human and Drosophila genes (FGFR-2, msl-2,
TIAR, cystic fibrosis transmembrane conductance regulator and Fas) through binding to U-rich stretches, facilitating atypical 5-splice site recognition by U1 small nuclear ribonucleoprotein [2,3].

TIA-1 has also been well characterized as a translational regulator. TIA-1 and TIAR are both able to bind to the 3-untranslated regions of the translational regulatory AU-rich elements of tumor necrosis factor, human matrix metalloproteinases-13, cyclooxygenase-2, 2-adrenergic receptor, mitochondrial cytochrome c, GADD45 and -F1-ATPase mRNAs [4-10].

These both protein may also promote cellular and virus-induced apoptosis, to be implicated in viral replication and to be required for DT40 cell viability. TIA-1 plays also important functions in apoptotic cell death and in adapting the cellular response to metabolic stress and inflammation. TIAR is translocated from the nucleus to the cytoplasm during Fas-mediated apoptosis. TIA-1 is a specific substrate for the Fas-activated protein serine/threonine kinase [10].

The isoforms of both mTIA-1 and mTIAR are predominantly expressed in brain, spleen and testis and mTIAR is also expressed in liver and lung. mTIA-1 and mTIAR are not expressed or only very weakly, in the other tissues tested such as heart, skeletal muscle and kidney [11].

The aim of this study was to estimate sodium iodide symporter (NIS) and thyroid peroxidase (TPO) expression in thyrocytes from patients with GD and no-toxic multinodular goiter (NTMG) in relation to the apoptotic markers.

Material and methods

Patients and study material. The study was performed in a group of 30 adolescent patients (8 boys and 22 girls) aged 8-21 years with GD (n=15, mean age 13.9±3.5 years) and nontoxic nodular goiter (NTNG; n=20, mean age 15.8±3.2 years) hospitalized in the Department of Pediatric Endocrinology and Diabetology, Poznañ (NTNG; n=20, mean age 15.8±3.2 years) hospitalized in the Department of Pediatric Endocrinology and Diabetology, Poznañ University of Medical Sciences. The patients underwent total or subtotal thyroidectomy in the 1st Department of General Surgery, Medical University of Białystok or in the Department of Pediatric Surgery, Poznañ University of Medical Sciences.

The diagnosis was established based on clinical examinations confirmed by laboratory, ultrasonographic and scintigraphic investigations with the use of 131I (in case of nodular goiter with symptoms of hyperthyroidism). Additionally, fine-needle aspiration biopsies of nodular goiter were performed in the Department of Pathological Anatomy, Medical University of Białystok and in the Department of Pediatric Endocrinology and Diabetology, Poznañ University of Medical Sciences. The patients underwent total or subtotal thyroidectomy in the 1st Department of General Surgery, Medical University of Białystok or in the Department of Pediatric Surgery, Poznañ University of Medical Sciences.

The function of the thyroid gland in patients with GD (as well as in nodular goiter) was assessed at the time of diagnosis and prior to surgery. Thyroid function was evaluated based on thyroid hormones and TSH tests performed jointly with the measurement of titers of antithyroid antibodies (ATPO, ATG, TRAK). The expression of proapoptotic proteins was identified in tissue material obtained from patients with immune and nonimmune disorders.

Determination of the antithyroid antibody titers and thyroid hormone concentration. Blood for analysis was collected on empty stomach in the morning hours from the basilic vein and centrifuged for 10 min at 2,000 rotations/min. Sera were stored at -20°C until the required number was collected. Immunodiagnostic test Varelisa (Variable Enzyme Linked Immuno Sorbent Assay, Pharmacia Upjohn Diagnostics, GmbH & Co.KG, Freiburg, Germany) was used to determine anti-peroxidase antibodies (anti-TPO) and anti-TG antibodies in the sera, using human microsomal antigen and human thyroglobulin, respectively. The results were read on a photometer (STAT FAX 303 PLUS, ANALCO-GBC), at 450 mm of light wavelength for which absorption values were proportional to the level of anti-TPO or anti-TG antibodies. The radioreceptor method (TRAK-human, Brahms Diagnostica, GmbH, Berlin, Germany) was employed to assay TRAB in blood serum. The TRAB level was negative at the values <1 U/l, doubtful between 1.0 and 1.5 U/l (grey zone) and positive above 1.5 U/l.

Determination of serum thyrotropin hormone (TSH) was done using a mini-analyzer VIDAS (bioMérieux) and VIDAS TSH test, being a combination of the immunoenzymatic method and final fluorescence measurement (ELFA). In the permanent phase, anti-TSH mouse monoclonal antibodies were used. Normal values for TSH ranged between 0.2 and 5.0 μIU/ml. Serum levels of free thyroxine (fT4) and free triiodothyronine (fT3) were determined using a mini-analyzer VIDAS fT4 and VIDAS fT3 tests that combine the immunoenzymatic method with the final measurement of fluorescence (ELFA).

Normal values ranged between 0.71 and 1.55 ng/dl for fT4 and between 2.6 and 5.4 ng/dl for fT3.

SDS-PAGE and Western Blotting. Tissue samples were homogenized in an ice-cold buffer (250 mM sucrose, 20 mM Tris-HCl, pH 7.4, 1 mM EDTA) containing a cocktail of protease inhibitors (Roche Diagnostics, GmbH, Mannheim, Germany). Homogenates were centrifuged at 1,000 g for 15 min at 4°C, then at 100,000 g for 60 min at 4°C. The resulting pellets containing particulate fractions were recovered, resuspended in 20 mM Tris-HCl, pH 7.4 with 1 μg/ml phenyl-methylsulphonyl fluoride and kept at -80°C. The protein concentrations were evaluated by the bicinchoninic acid protein assay reagent (Pierce Chemical, Co., Rockford, Ill., USA).

A total of 75 μg of crude membrane proteins was mixed with a protein buffer (0.25 mM Tris-HCl, pH 6.8, 20% glycerol, 4% SDS, and 0.1% bromophenol blue) and incubated with 0.125 M dithiothreitol for 30 min at 37°C. Each sample was loaded into individual wells and electrophoresed on a 9% acrylamide, using the SDS-PAGE method. Proteins were electrotransferred to immunoblot PVDF membranes (Bio-Rad Laboratories, Hercules, Calif., USA), which were then saturated with 5% powdered milk in PBS-Tween. Western blotting experiments were subsequently carried out by incubating the blotted membranes with mouse monoclonal anti-human TIAR and TIA-1 antibodies (BD Biosciences Pharmingen) diluted according to the manufacturer’s suggestion overnight at 4°C. After three washings in PBS-Tween for 10 min,
the membranes were incubated with affinity purified anti-mouse antibody labeled with horseradish peroxidase (Jackson ImmunoResearch Laboratories, West Grove, Pa., USA) for 1 h at room temperature under shaking. After extensive washing, membranes were developed with a SuperSignal West Pico (Pierce). The visualization of proteins was performed by chemiluminescence detection procedure – exposure to BioMax Ms film (Sigma-Aldrich, Corp., St. Louis, Mo., USA). The secondary antibody controls were included in the series of Western blots. Immunoblots were reprobed with β-actin antibody 1:10,000 (mouse monoclonal antibody; Sigma) for normalization.

Immunohistochemistry. Tissue sections of 3 μm were mounted on silane-coated glass slides, deparaffinized in xylene, and rehydrated via graded ethanols to water. Then the antigen was heat retrieved at 95-99°C for 20 min in TRS pH 6.0 and after endogenous peroxidase block slides were incubated with anti-NIS, anti-TPO#47, anti-TIAR and anti-TIA-1 (at a concentration according to the manufacture formula) overnight at 4°C; after washing sections were incubated with LSAB+ kit (DAKO). All primary antibodies used were commercially available (BD Biosciences). The reaction was developed with DAB chromogene. The slides were counterstained with Mayer's hematoxylin. The specificity of the immunostaining was checked by omission of single steps in the protocol, replacement of the primary antibody with preimmune serum and peptide competition tests. Slides that showed no staining were considered negative, all other slides with staining were considered positive. Two pathologists independently evaluated reaction on randomly numbered slides. Expression index was created by classifying the samples into three categories based on the percentage of positive cells in the total number of cells counted per field. Grade I (+) included samples with less than 10% positive cells, grade II (++) samples had 10-50% positive cells, and grade III (+++) samples showed more than 50% positive cells.

Ethical issues. Our study was approved by the Committee for Ethics and Supervision on Human and Animal Research of the Medical University of Białystok.

Statistical analysis. The results were analyzed using Statistica 8.0 software. The mean values of immune parameters between groups were evaluated using Student's t test, U Mann-Whitney's test or Fisher's exact probability test. Correlation was assessed by Spearman's signed ranks test. P<0.05 was considered significant.

Results

Table 1 presents the characteristics and laboratory findings of patients with GD (prior to methimazole therapy and during clinical-biochemical euthyreosis before surgery) or with NTMG (before surgery). Patients with nodular lesions above 1 cm in size seen on ultrasonography underwent fine-needle aspiration biopsy, which revealed benign changes in the form of 'colloid nodular goiter'. Scintigraphy additionally performed in 3 patients with nodular goiter and clinical symptoms of hyperthyroidism showed a selective increase in the accumulation of radioactive iodine within the area of single nodular lesions. Postoperative tissue material was used to identify proapoptotic regulatory proteins of the TIAR and TIA family by means of immunohistochemistry and Western blot analysis.

Identification of proapoptotic TIAR (TIA-1 related protein) molecule in the thyroid tissues revealed a higher expression of this protein in patients with Graves' disease (+++) in comparison to patients with NTNG (+) (Figs. 1 and 2). The elevated expression of TIAR in the thyroid tissue of patients with GD was observed mainly in hyperfunctional thyrocytes (with higher epithelium). The expression of TIA-1 molecule was detected only in patients with Graves' disease (+) in lymph nodes with germinal centers.

The tissue material was additionally subjected to Western blot analysis, which in GD patients showed the presence of TIA-1 in the band p22, p17 (kDa)] (Fig. 3a). In group of patient with NTMG presence of TIA-1 in the band 17 (kDa) were observed (Fig. 3b). In GD patients, Western blot analysis showed TIAR expression in band p50, p42, p38 (kDa) (Fig. 4a). In patients with NTNG, the degree of expression of TIAR
proteins was lower and referred to band 42 (kDa) (Fig. 4b). Each thyroid tissue sample obtained from our study patients underwent assessment for the presence of β-actin using the Western blot method, which confirmed that the tissues were not degraded and indicated high-quality analyses (Fig. 5). This study showed increased expression of proapoptotic markers in thyroid tissues from Graves’ patients in comparison to patients with nontoxic nodular goiter.

The relationship between expression of NIS and TPO and the expression of apoptosis regulatory molecules of the TIA-1 family were taken into consideration. The expression of NIS and TPO in thyroid follicular cells was higher in patients with Graves’ disease (+++, ++, respectively) in comparison with their detection in patients with NTMG (+, +, respectively) (Fig. 6 and 7). In addition, the degree of thyroid antigen expression was positive, and correlated with the amount of proapoptotic markers in thyroid tissues. (TIAR, \(p<0.001\); TIA-1, \(p<0.025\) for NIS; TIAR, \(p<0.012\) for TPO). (Table 2). However, such a correlation was not observed in patients with NTNG.

**Discussion**

TIA-1 and TIAR are RNA binding proteins of the RNA recognition motif (RRM)/ribonucleoprotein (RNP) family that have been implicated as effectors of apoptotic cell death. The molecular mechanism by which the PID (protein-interaction domains) of an RNA binding protein might trigger apoptotic cell death is not fully known. Taupin et al proposed that p15-TIA-1 (15-kDa cytotoxic granule-associated protein) might affect the function of TIAR, a TIA-1-related RNA-binding protein what was identified by low-stringency hybridization. Like p40-TIA-1, TIAR possesses three N-terminal RRMs and a C-terminal PID. The RRMs of TIA-1 and TIAR are in 90% identical at
Like TIA-1, TIAR triggers DNA fragmentation in permeabilized thymocytes, suggesting its possible involvement in apoptosis. Results of their study show that TIAR is a ubiquitously expressed nuclear protein that rapidly moves to the cytoplasm in response to exogenous triggers of apoptosis. TIAR is normally confined to the nucleus of cells, but during Fas-mediated apoptosis, it is rapidly translocated to the cytoplasm. Cytoplasmic redistribution precedes the onset of DNA fragmentation and the nuclear architectural changes that facilitate histone extraction [12].

Similar studies were conducted by Tain et al, who proved that TIA-1 itself has been linked to apoptosis.
The serine-threonine kinase FAST is activated during Fas-mediated apoptosis in Jurkat cells and phosphorylates TIA-1 prior to the onset of DNA fragmentation [13]. In the mouse model TIAR were essential for primordial germ cell development, as it appears to be necessary for cell survival [14]. It may be suggest that proapoptotic stimuli modify the activities of TIA-1 and TIAR, leading to changes in the splicing patterns of key pre-mRNAs.

Tsuzuki et al showed that apoptosis triggered by 20-hydroxyecdysone during pupal metamorphosis of the silkworm Bombyx mori is accompanied by an increase in TIAR mRNA levels [15]. Jin K. et al indicated that ischemia-induced apoptosis also leads to an increase in TIAR mRNA levels [16]. These observations also suggest that TIAR could be required for apoptosis. TIA-1 and TIAR may normally work for cell survival, but during apoptosis they are modified, so as to be rendered inactive for cell survival tasks, yet active to participate in new tasks in apoptosis progression [17].

The role of TIA and TIA-1 hasn't been explained in thyroid autoimmunological diseases. Immunohistochemistry revealed that lymphoid cells may be positive for T-cell-restricted intracellular antigen 1 (TIA-1). Okamoto et al. described a case of patient with T-cell lymphoma without clinical or laboratory symptoms of autoimmune thyroiditis. They suspected that TIA-1 expression together with TCRβ gene rearrangement seemed categorized into TCRα/b+ cytotoxic T-cell lymphoma. Although Yamaguchi et al. reported a case of TCRγ/d+ thyroid T-cell lymphoma, no case of thyroid T-cell lymphoma expressing cytotoxic molecules such as TIA-1 has been reported [18,19].

In our studies elevated expression of TIAR/TIA-1 in GD in comparison with its expression in non-immune diseases indicated to enhanced apoptosis activity in immune process. Our findings suggest that significant expression of TIAR may indicate on enhanced activity of egzogenous way of apoptosis in autoimmune diseases, for example in GD. Kawakami et al observed the association between the apoptosis of thyrocytes and the process of autoimmune thyroid diseases and suggested that Fas-mediated apoptosis of human thyrocytes is modulated by thyroid-stimulating antibodies, thyroid stimulation blocking antibodies and cytokines [20]. Similarly, we conclude that elevated expression of NIS and TPO in Graves’ disease is associated with higher stimulation and activation of apoptosis in thyroid follicular cells during autoimmune process. According to our knowledge the correlation between NIS and TPO and apoptosis regulatory molecules of the TIA-1 family in thyroid diseases was not considered in any publication. Therefore more and more observations are needed. The understanding of the apoptotic process at the molecular level may broaden the knowledge of the pathogenesis of autoimmune thyroid disorders and provide new diagnostic, prognostic and therapeutic potentials.

We conclude that elevated expression of NIS and TPO in Graves' disease is associated with higher stimulation and activation of apoptosis in thyroid follicular cells during autoimmun search process.

References


Submitted: 3 January, 2010
Accepted after reviews: 29 March, 2010