

Effects of thyroxine on adhesion molecules and proinflammatory cytokines secretion on human umbilical vein endothelial cells

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Abstract

Thyroid dysfunction is associated with elevated cardiovascular risk factors and atherosclerosis. It could be suggested that, hyperthyroidism is related to a higher prevalence of arterial abnormalities. Therefore, evaluating the endothelial dysfunction (ED) related biomarkers seem to be an important issue. It is not clear whether endothelial cells are biologically responsive to thyroid hormones (THs) or how THs induces the production of endothelial cells (EC)-derived proinflammatory mediators. Hence, in this study the effects of thyroxine (T₄) on ED and inflammatory related mediators were evaluated. Human umbilical vein endothelial cells was used as endothelial cell model which was treated with concentrations of 50, 100, 200 nmol/L of T₄ in various exposure times. In the following, gene and protein expression levels of EC-related markers including intercellular adhesion molecule-1 (ICAM-1), vascular endothelial growth factor (VEGF), and E-selectin were determined using real time polymerase chain reaction (RT-PCR) and western blotting methods. Also, interleukin-6 (IL-6) and tumor necrosis factor (TNF- α) protein levels as proinflammatory cytokines were determined by enzyme linked immunosorbent assay (ELISA) method. Gene and protein expression analysis revealed that T₄ treatments up regulated the levels of VEGF, ICAM-1, and E-selectin as ED markers. In addition, T₄-treated cells had higher significant levels of IL-6 and TNF- α versus untreated cells in different incubation times. This study proposed the atherosclerotic effects of thyroid hormone. Based on our findings, T₄ had strong effects on the gene and protein expression levels of pro-inflammatory, angiogenesis, and ED major mediators associated with atherosclerosis development.

Keywords: E-selectin; ICAM-1; Interleukin-6; Thyroxine; TNF- α ; VEGF.

INTRODUCTION

Thyroid hormones (THs) exert numerous effects on the cardiovascular system. It has been indicated that thyroid dysfunction is related to cardiovascular diseases, and both hyperthyroidism and hypothyroidism accelerate cardiovascular disorders. Although with the advances in the prevention and treatment of atherosclerosis, this disorder still stands as one of the main causes of morbidity and mortality worldwide (1). THs have been related to several pro-atherogenic condition, but the relationship between thyroid function with atherosclerosis has not been fully determined (2). It could be suggested that, hyperthyroidism is associated with a higher prevalence of arterial abnormalities (3). A low thyroid-stimulating hormone (TSH) level and increased levels of

L-thyroxine (T₄), triiodothyronine (T₃) or both, is recognized as hyperthyroidism. THs activate vascular endothelium and decrease the metabolism of adhesion molecules, which lead to increase of their circulating levels. This theory that vascular endothelium are targets of THs are endorsed by increased endothelial function during hyperthyroidism (4,5). Previous studies revealed that hyperthyroidism elevated the circulating levels of biological mediators, which had major roles in hemostasis, vascular tone regulation, and permeability (3,4). Indeed, vascular endothelial tissues have receptors for THs and respond to changes in the concentrations of circulating THs (6).

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DOI: 10.4103/1735-5362.258490

In fact, disruption of coronary endothelial function is related to increased risk of cardiovascular disorders (7). Therefore, evaluating the endothelial dysfunction (ED) related biomarkers such as proinflammatory mediators including interleukin-6 (IL-6) and tumor necrosis factor (TNF)- α are notable issues. Studies have shown that the increased circulating levels of IL-6 and TNF- α have main roles in peripheral endothelial dysfunction (7-10). Definitely, vascular endothelium is responsive in inflammatory reactions, which is engaged in atherosclerosis. In addition, cytokines play an important role in endothelial disruption, which is related to inflammation (11). In this regards, IL-6 as an inflammatory mediator was synthesized by various cells, including endothelial cells (ECs) and also, inflammatory cytokine TNF- α had important function in the impairment of macrovascular and microvascular circulation (11,12). Activation of EC is defined as endothelial expression of cell-surface adhesion molecules, including intercellular adhesion molecule-1 (ICAM-1) and endothelial leukocyte adhesion molecule E-selectin. This event is characteristically prompted by proinflammatory cytokines such as IL-6 and TNF- α which cause the leukocytes to be attached to the vascular wall (13).

On the other hand, vascular endothelial growth factor (VEGF) is considered as the main mediator in the regulation of vascular growth, vasculature hemostasis (14) and acts as potent mitogen for ECs. In addition, VEGF is related to increased vascular permeability. This angiogenesis related mediator is detectable in many tumors such as thyroid cancerous cells (15). On the other hand, reports have shown the pro-angiogenic role of THs analogues, including T₄ (16). However, it is not clear whether endothelial cells are biologically responsive to THs or how THs induce the production of EC-derived proinflammatory mediators. The present study aimed to investigate the impacts of T₄ on EDs with insight to atherosclerosis induction related pathways and EC activation.

MATERIALS AND METHODS

Chemicals

L-Thyroxine powder was obtained from Sigma-Aldrich Company (Cat No: T1775, St. Louis, Missouri, USA) and liquidized in ethanol solution. Anti-E-selectin, -ICAM-1, -VEGF, and - β -actin were purchased from Santa Cruz Biotechnology, Inc (California, USA). The RNA Extraction and cDNA synthesis Kits were purchased from GeneAll Company (South Korea). A real time polymerase chain reaction (RT-PCR) master mix was obtained from Ampliqon (Herlev, Denmark). Western blotting materials were purchased from Bio-Rad Laboratories Inc (Hercules, California, USA). TNF- α (BE55181) and IL-6 (BE53061) enzyme linked immunosorbent assay (ELISA) kit was prepared from IBL Company (IBL International GmbH, Hamburg, Germany).

Cell culture and treatments

Human umbilical vein endothelial cells (HUVECs, NCBI code: C554, Pasteur Institute of Iran, Tehran, I.R. Iran) were cultured in DMEM/F12 supplemented with 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin in 5% CO₂ at 37 °C. For analytical examinations, 70% confluent cells were incubated 24 h before treatment in a serum-free medium, then medium removed and cells washed with PBS. In following, the HUVECs cells were treated with different doses of T₄ at 50, 100, and 200 nmol/L for 24 and 48 h, respectively.

ELISA

The effect of T₄ on proinflammatory cytokines was evaluated by determining the IL-6 and TNF- α levels using ELISA method. In this regards, 5×10^3 cells/well of HUVECs cells were seeded overnight and then treated with T₄ (50, 100, and 200 nmol/L) for 12, 24, and 48 h, respectively. After treatment times, the cell culture plates were centrifuged in 1500 g for 5 min and then the cell supernatant was collected. This was followed by protein detection based on ELISA kit protocols. The absorbance of wells were read using ELISA reader at 450 nm and then quantified.

Real-time PCR

Cells (10⁶ cells/well) were seeded in culture plates and then treated with T₄ (50, 100, 200 nmol/L) (17) for 24 and 48 h respectively. In the following, total RNA was extracted according to the manufacturer's guidelines. Total RNA was reverse transcribed into cDNA using hyperscript reagent kit and RT-PCR was performed. All the quantitative RT-PCR measurements were carried out using a Mic qPCR (Bio Molecular Systems, Australia) with mastermix. The designed Primer pairs for ICAM-1, E-selectin, VEGF, and β -actin are presented in Table 1. Second-strand synthesis and RT-PCR amplification were done for 40 cycles with denaturation at 95 °C for 15 s, annealing at 62 °C for 30 s, and extension at 72 °C for 30 s, with final extension at 72 °C for 5 min after completion of all cycles. The results were measured by the 2^{- $\Delta\Delta$ Ct} method (18).

Western blotting analysis

HUVECs (10⁶ cells/well) were treated with 50, 100, and 200 nmol/L concentration of T₄ for 24 and 48 h, respectively, then trypsinized (19) and collected. Radio immunoprecipitation assay lysis buffer containing complete protease inhibitors was added to cell pellet and maintained on ice for 20 min. The mixture was then centrifuged at 12000 g for 15 min at 4 °C. The collected supernatant was used for protein assay according to Bradford method protocol (20). Proteins were separated on SDS polyacrylamide gel (21) and then transferred onto a polyvinylidene fluoride membrane. The membrane was blocked using 5% skimmed milk solution at room temperature followed by 24 h incubation

at 4 °C with primary antibodies (anti-ICAM-1, -E-selectin, -VEGF, and - β -Actin). The membrane was then washed with TBST and incubated with secondary antibody (Abcam, USA) for 3 h, while the polyvinylidene fluoride was washed with TBST three times. The membrane was incubated with enhanced chemiluminescence western blotting detection reagent (GE Healthcare, Boston, MA, USA) and exposed to X-ray film. Protein bonds were quantified using the image J software and any differences in protein expression levels were identified by normalization to the levels of the β -actin protein.

Statistical analysis

The data were analyzed using Prism software, version 6.1 and were presented as mean \pm standard deviation (SD). For multiple comparisons between groups, one-way analysis of variance (ANOVA) was utilized and *P* values < 0.05 were considered as significant.

RESULTS**Effect of thyroxine on interleukin-6 and tumor necrosis factor**

Significant elevated IL-6 levels in T₄ treated cases (50, 100, and 200 nmol/L after 12, 24, and 48 h) were obtained when compared with the untreated cells (*P* < 0.05, 0.01, and 0.001, respectively; Fig. 1). Further analysis showed that there were no significant differences between various T₄ treated concentrations in all examined treatment times (*P* > 0.05).

Table 1. The designed primer pairs for ICAM-1, E-selectin, VEGF, and β -actin for RT-PCR.

Target gene	Primer Sequence (5'→3')		Product Size (bp)
ICAM-1	Forward	AGACATAGCCCCACCATGAG	121
	Reverse	CAAGGGTTGGGGTCAGTAGA	
E-selectin	Forward	ATGTGAAGCTGTGAGATGCG	152
	Reverse	CCACTGCAGCTCATGTTGAT	
VEGF	Forward	AGGAGGAGGGCAGAATCATC	90
	Reverse	GGCACACAGGATGGCTTGAA	
β -actin	Forward	CTGGAACGGTGAAGGTGACA	161
	Reverse	TGGGGTGGCTTTTAGGATGG	

ICAM-1, intercellular adhesion molecule-1; VEGF, vascular endothelial growth factor; RT-PCR, real time polymerase chain reaction.

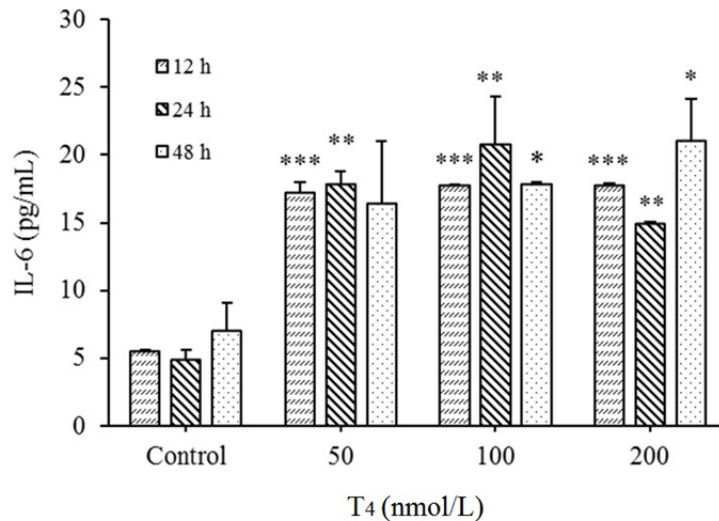


Fig. 1. Up regulation of interleukin-6 (IL-6) levels in HUVECs by T₄ exposure. After treatments with T₄ (50, 100 and 200 nmol/L) for 12, 24, and 48 h, protein expression was evaluated by ELISA method. **P* < 0.05, ** *P* < 0.01, and *** *P* < 0.001, indicate significant differences in comparison with control group. Data are presented as mean ± SD. Values derived from three independent experiments. T₄, thyroxine.

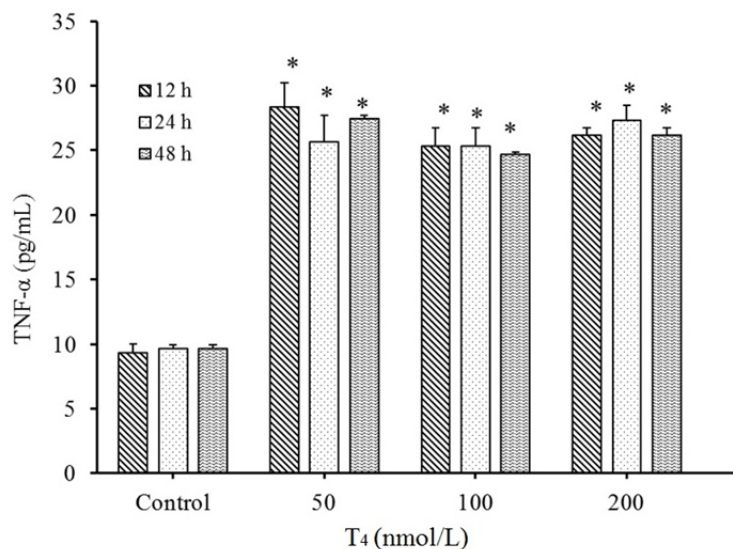


Fig. 2. Up regulation of tumor necrosis factor-α (TNF-α) level in HUVECs by T₄ exposure. After treatments with T₄ (50, 100, and 200 nmol/L) for 12, 24, and 48 h, protein expression evaluated by ELISA method. **P* < 0.001 indicates significant differences compared with control group. Data are presented as mean ± SD. Values derived from three independent experiments.

TNF-α levels of T₄ treatments (50, 100, and 200 nmol/L) were increased in all studied times compared to the controls (*P* < 0.001, Fig. 2) whereas no significant differences in the levels of TNF-α between different concentrations of T₄ were found (in all examined treatment times) (*P* > 0.05).

On the other hand, serum IL-6 and TNF-α level increased after T₄ treatments in all tested

concentrations, but there were no significant differences between various treatment times (*P* > 0.05).

Effects of thyroxine on the VEGF, ICAM-1, and E-selectin gene expression

mRNA levels of ICAM-1, E-selectin, and VEGF were detected by RT-PCR and the Ct values of each samples were converted to fold changes using $2^{-\Delta\Delta Ct}$ formula and then

compared with the fold changes of the control group. Based on our results, E-selectin mRNA was significantly increased in all the examined concentrations after 24 and 48 h compared to the control cells (except 200 nmol/L). However, as presented in Fig. 3A, the gene expression levels of E-selectin were increased more significantly at lower T_4 concentrations. Indeed higher

E-selectin levels were observed at lower T_4 concentrations in both times.

After 24 and 48 h, ICAM-1 gene expression levels increased in all examined T_4 concentrations compared to the control ($P < 0.05$ and 0.01 , respectively) (Fig. 4A and 4B). In addition, the gene expression levels of ICAM-1 had highest levels at 100 nmol/L of T_4 compared with other T_4 concentrations in both times.

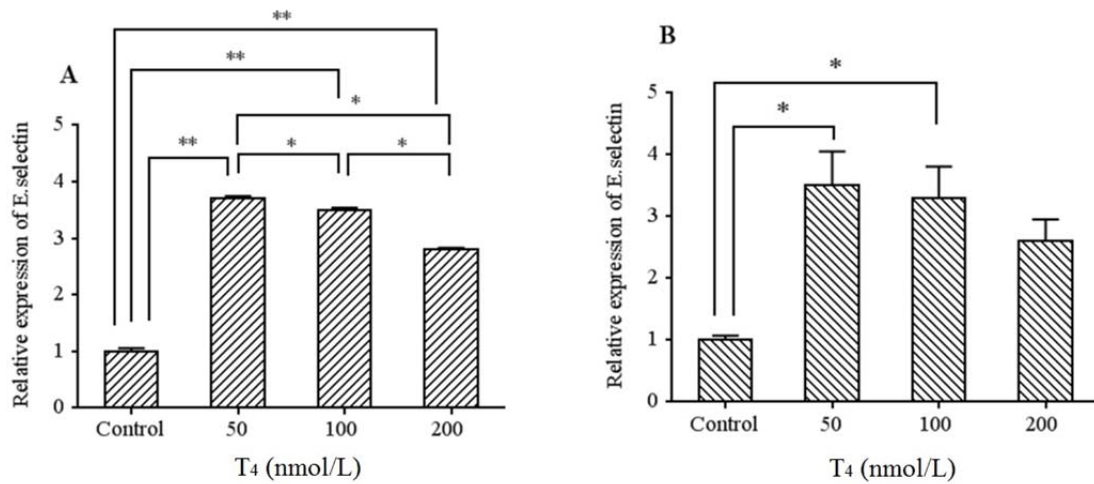


Fig. 3. mRNA levels of E-selectin in HUVECs cells treated by T_4 (50, 100, and 200 nmol/L) for (A) 24 h and (B) 48 h exposure times. Data presented as mean fold change \pm SD. Values derived from three independent experiments. * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$ indicate significant differences in comparison with control group.

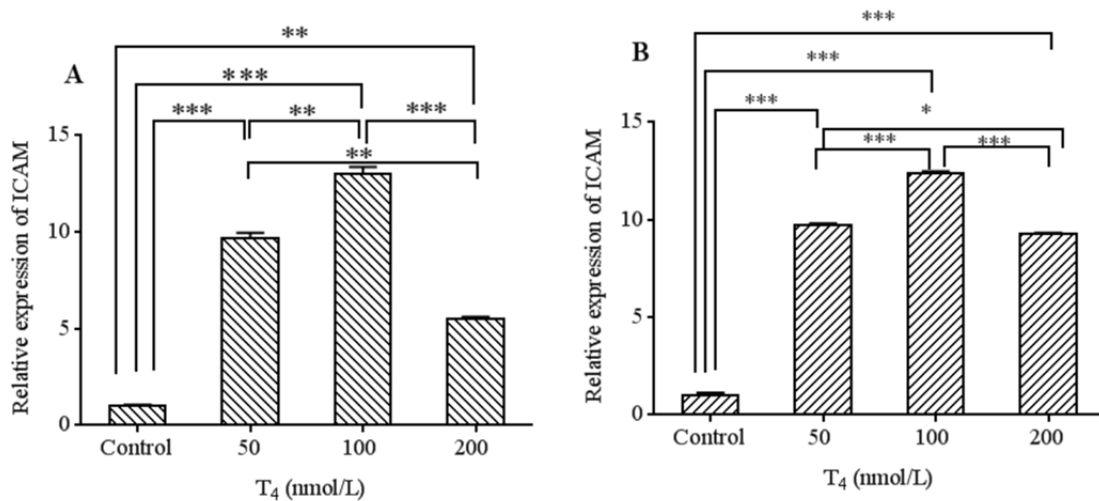


Fig. 4. mRNA levels of ICAM-1 in HUVECs cells treated by T_4 (50, 100, and 200 nmol/L) for (A) 24 h and (B) 48 h exposure times. Data presented as mean fold change \pm SD. Values derived from three independent experiments. * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$ indicate significant differences in comparison with control group. ICAM-1, intercellular adhesion molecule 1.

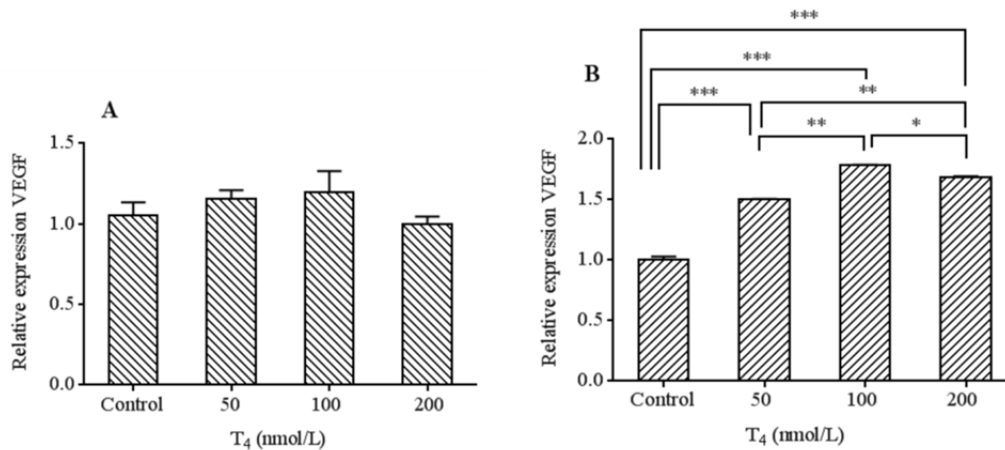


Fig. 5. mRNA levels of VEGF in HUVECs cells treated by T₄ (50, 100 and 200 nmol/L) for (A) 24 h and (B) 48 h exposure times. Data presented as mean fold change ± SD. Values derived from three independent experiments. **P* < 0.05, ***P* < 0.01, and ****P* < 0.001 indicate significant differences in comparison with control group. VEGF, Vascular endothelial growth factor.

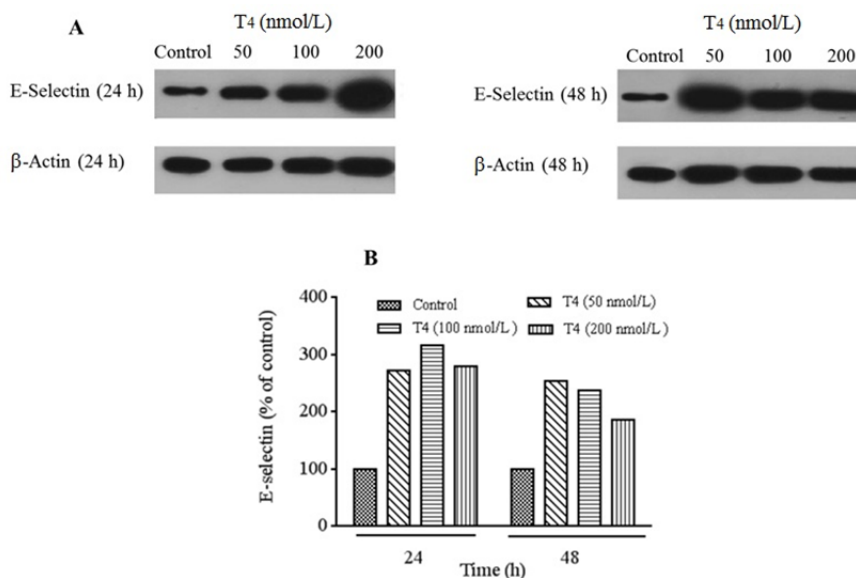


Fig. 6. Protein detection for evaluation of E-selectin expression performed using western blotting technique. (A) Protein expression levels of E-selectin in HUVECs cells treated by T₄ (50, 100, and 200 nmol/L) for 24 and 48 h and (B) T₄ treatment significantly increased relative expression of E-selectin compared with control group. For analysis protein's bands densities, Image J software was used to analyze the densities. Each protein bands normalized to β-actin bands.

Based on our findings VEGF mRNA levels elevated in all T₄ treatment compared to the control group (Fig. 5) which in the cases of 48 h incubation time, the extent of increase was more noticeable and reached significant levels (*P* < 0.001, Fig. 5B).

No significant differences in the levels of VEGF mRNA between different concentrations of T₄ after 24 h were found (Fig. 5A). There were significant differences in the levels of VEGF between different concentrations of T₄ after 48 h (*P* < 0.05).

Additionally, higher VEGF levels were presented at 100 nmol/L in all exposure times.

Effects of thyroxine on the E-selectin, ICAM-1, and VEGF protein expression

To identify, E-selectin, ICAM-1, and VEGF protein expression levels in HUVECs cells, protein bindings were assessed by western blotting and band intensities were quantified using the Image J software. As presented in Fig. 6A and 6B, E-selectin protein levels increased after 24 and 48 h of treatments (50, 100, and 200 nmol/L of T₄) compared to

untreated cells. The extent of increase after 24 h was more noticeable than 48 h treatment times. The protein expression level of E-selectin was in accordance with the RT-PCR gene expression results in all selected doses.

ICAM-1 protein band assessment showed that, this protein levels was increased after 24 and 48 h of treatments in all examined T₄ concentrations (Fig. 7A and 7B). Furthermore, T₄ at 100 nmol/L exhibited highest impact compared to other tested concentrations. The results of protein expression assay of ICAM-1 were parallel to RT-PCR gene expressions.

Analysis of western blotting protein bands showed that, VEGF protein levels were elevated over 24 and 48 h of treatment with 50, 100, and 200 nmol/L of T₄ compared to the untreated cells (Fig. 8A and 8B). According to our findings, VEGF protein expression at 100 nmol/L had the highest effect compared to the other examined T₄ concentrations after 48 h treatment whereas after 24 h exposure, higher protein expression was present in 50 nmol/L of

T₄. VEGF protein expression was in agreement with RT-PCR gene expressions.

DISCUSSION

It has been reported that THs have noticeable effects on cardiovascular system, so thyroid abnormalities have led to abnormal cardiovascular functions. Indeed, endothelial cells are the possible targets for THs (22-24). In the present study we evaluated the possible effects of THs on ED related mediators.

Our findings confirmed the association between T₄ treatments and ED in HUVEC. Furthermore, our findings may suggest that T₄ administration is associated with ED, which is reflected by increased proinflammatory cytokines (IL-6 and TNF- α) and higher levels of ED markers including VEGF, ICAM-1, and E-selectin. Moreover, we found that T₄ treatment could enhance ED in these cellular models, at different treatment times.

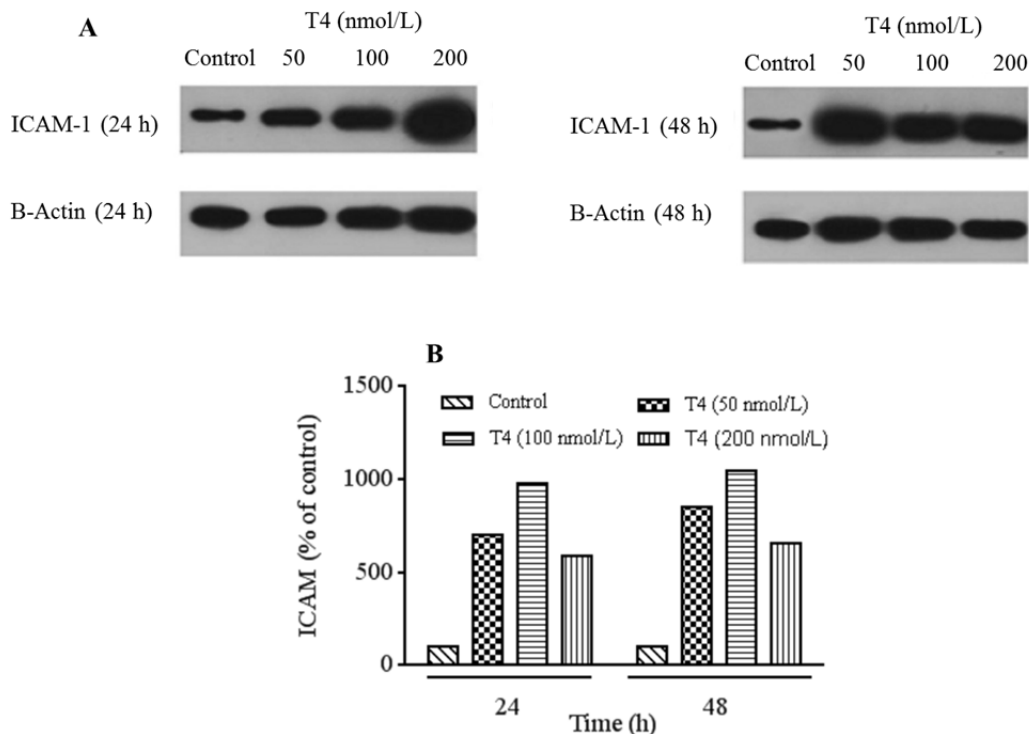


Fig. 7. Protein detection for evaluation of ICAM-1 expression performed using western blotting technique. (A) Protein expression levels of ICAM-1 in HUVECs cells treated by T₄ (50, 100, and 200 nmol/L) for 24 and 48 h and (B) T₄ treatment significantly increased relative expression of ICAM-1 compared with control group. For analysis protein's bands densities, Image J software was used to analyze the densities. Each protein bands normalized to β -actin bands. ICAM-1, Intercellular adhesion molecule 1.

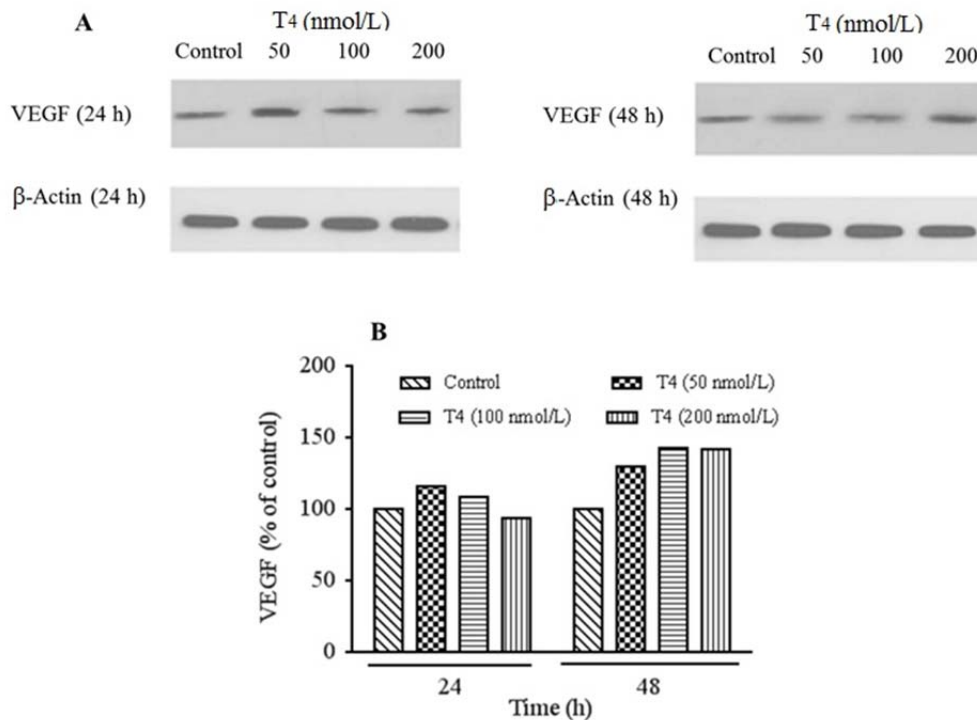


Fig. 8. Protein detection for evaluation of VEGF expression performed using western blotting technique. (A) Protein expression levels of VEGF in HUVECs cells treated by T₄ (50, 100, and 200 nmol/L) for 24 and 48 hand (B) T₄ treatment significantly increased relative expression of VEGF compared with control group. For analysis of protein's band densities, Image J software was used to analyze the densities. Each protein bands normalized to β-actin bands. VEGF, Vascular endothelial growth factor

However, the impairment of endothelial function could be explained by the increased level of potential ED markers as cardiovascular risk factors, thus it could be suggested that T₄ exerted atherogenic activity or acted as a major determinant in vascular homeostasis regulation. In agreement with our results, Popławska-Kita *et al.* indicated that both subclinical hyperthyroidism and overt hyperthyroidism may be accompanied with ED, including elevated levels of IL-6 (4). In another survey, Burggraaf *et al.* reported that increased THs was related to increased levels of several endothelium-associated proteins. They did not find any relationships between hyperthyroidism and levels of the inflammation markers IL-6, TNF-α. This discordance in comparison with our results may be related to the fact that Burggraf *et al.* evaluated the circulating levels of these parameters, while we detected ED mediators in endothelial cell levels (3). In another survey, hyperthyroid rat had elevated circulating levels of TNF-α. These researchers suggested that altered levels of

TNF-α resulted from thyroid effects on oxidative stress in rat liver (25). Overall, it could be suggested that inflammatory response against ED is dependent on the functions of IL-6 and enormous proinflammatory cytokines including TNF-α (11). Previous studies reported that ICAM-1 not only acts as leukocyte adhesion molecule, but also participates in inflammatory responses in vascular wall by induction of ECs activation, which result in atherosclerotic plaque developments (26). In this regards, Hara *et al.* reported that THs increases the expression of E-Selectin from EC in patients with Graves' disorder, which is comparable with our results (27). Other reports proposed that, ICAM-1 and IL-6 might be considered as major element in the assessment of the inflammatory events in the thyroid gland throughout radioiodine treatments, particularly Graves' disease (28). Recently, numerous studies evaluated the T₄-induced angiogenesis (17). In these ways, VEGF as a noticeable inducer of angiogenesis and important mitogen for EC, activates specific signal transduction

pathways, which causes EC proliferation and angiogenesis (29). In addition, in our study, THs administration stimulated the expression of VEGF in gene and protein levels, which shows the antigenic effects of T₄.

In agreement with our findings, Liu *et al.* demonstrated that T₄ induced the migration and formation of tube-like structures in HUVECs. In addition, they indicated that T₄ up regulated the expression of basic fibroblast growth factor mRNA via effects on integrin avb3/PKD/HDAC5 signaling pathway, which had a main role in angiogenesis (17).

On the other hand, we confirmed the atherogenic role of T₄ on HUVECs cells. Indeed T₄-induced inflammation and atherogenesis may be due to these facts, which have been described by Szmítko *et al.* After ECs activation due to inflammatory events, the elevated levels of selectins, VCAM-1, and ICAM-1 stimulate the monocytes adherence (30) and also adhesive molecules are regulated by proinflammatory cytokines including IL-1 and TNF- α , which consequently lead to monocytes migration and related stigma (31). To the best of our knowledge, this is the first survey that compares the effects of T₄ on proinflammatory mediators (IL-6 and TNF- α) and ED related factors. In summary, our results confirmed the relationship between T₄ and endothelium function in human cellular model, which was found to be able to predict the atherosclerosis risk factors.

CONCLUSIONS

Overall, our study suggests the atherosclerotic effects of THs and the relationships of hyperthyroidism with atherogenesis. According to the present study, T₄ had strong effects on the gene and protein expression levels of ICAM-1 and E-selectin as leukocyte adhesion molecules and VEGF as angiogenesis main mediator. In this regards, increased expression of proinflammatory and EC related mediators were associated with augmentation of atherosclerosis risks due to THs. Further studies should be conducted to confirm the impacts of T₄ on the function of

endothelial cells and inflammatory mediators in order to obtain more reliable data.

ACKNOWLEDGMENT

This research was financially supported (Grant No. 95-01-43-2787) by The Urmia University of Medical Sciences, Urmia, I.R. Iran. We are thankful to the technicians of Cellular and Molecular Research Center, Urmia University of Medical Sciences, Urmia, I.R. Iran for their technical helps.

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