

## Increased levels of antibodies against heat shock proteins in stroke patients

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Ischemic stroke is the second leading cause of death worldwide. One of the main risk factors of the ischemic stroke is atherosclerosis which is a chronic inflammatory and immune-mediated disease. Bacterial infections generate specific human antibodies against various antigens, including Hsps. It has been demonstrated that Hsps are selectively overexpressed in the atherosclerotic lesions. The amino acid sequence homology between human and bacterial Hsps may lead to an autoimmune response by immunological cross-reaction. Such immune response against Hsps overexpressed in the blood vessels under stressful conditions may contribute to inflammatory processes and subsequent development of atherosclerosis. In this study we determined the antibody levels against bacterial and human Hsp by ELISA in blood plasma obtained from stroke patients. Using ANOVA we analyzed levels of Hsp-antibodies in control and patient groups and correlate them with several stroke risk factors. The group of stroke patients had elevated levels of anti-Hsp antibodies compared to the control group. We also discovered an antibody level increase in patients that previously underwent another stroke. Our data provide evidence that autoimmunity could underlie formation of atherosclerosis plaque leading to stroke.

**Key words:** HSP, Heat Shock Proteins, Stroke, DnaK, DnaJ, GroEL, Hsp70, ELISA

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### INTRODUCTION

Heat shock proteins are a family of highly conserved proteins with well-studied structures and functions. Under physiological conditions, they constitute 5–10% of the total protein level in the cell. Their synthesis is constitutive or induced under stress conditions, i.e. heat shock, oxidative shock, viral infection, NO, UV, ethanol, certain chemicals or exposure to cytokines (Georgopoulos & Welch, 1993; Pockley, 2001; Xu, 2002; Mehta *et al.*, 2005). The basic functions of Hsp may include protective activity and weakening of the intracellular and extracellular stressors. They are also involved in the synthesis and folding of newly formed proteins and they are involved in many metabolic pathways (Hartl, 1996). A multitude of heat shock protein functions is related to their ability to form complexes with many proteins, abil-

ity to induce conformational changes and aid in crossing cellular membranes. Hsp also take part in regulation of apoptosis in the cell (Beere, 2005; Kaźmierczuk *et al.*, 2009). Based on molecular weight, heat shock proteins can be divided into several groups based on molecular weight: small Hsp, Hsp40, Hsp60, Hsp70 and Hsp90 (Kaźmierczuk *et al.*, 2009).

Representatives of small Hsp family are IbpA and IbpB. A characteristic feature of these enzymes is relatively low molecular weight (12–43 kDa) and the presence of a conserved  $\alpha$  — crystalline domain. In addition, proteins belonging to this family have oligomeric structure. Small Hsp expression is induced under the influence of stress, such as elevated temperatures. Their function is to prevent irreversible aggregation of denatured proteins (Kuczyńska-Wiśnik *et al.*, 2010).

An example of Hsp40 family member is DnaJ, with a mass 41 kDa. DnaJ also belongs to a family of -JDP, i.e. proteins that have a 70 amino acid motif J. DnaJ domain is primarily responsible for the stimulation of ATP hydrolysis of DnaK (Banecki *et al.*, 1996; Nakamoto *et al.*, 2014).

On the other hand, GroEL is a representative of Hsp60 family. Its main function is chaperone activity in the protein synthesis process. Only 84 proteins are predicted to absolutely depend upon GroEL to fold correctly, but these include at least 13 essential proteins.

DnaK is the main protein in the Hsp70 system, displaying ATPase activity. This protein is expressed under physiological growth conditions, but it is largely induced in response to heat shock, oxidative or osmotic stress, and limiting nutrient levels (Dahiya *et al.*, 2014). Proteins belonging to the Hsp70 family are characterized by their N-terminal ATPase domain, a substrate binding domain, and a short C-terminal domain, which could interact with various protein partners, thereby regulating the function of a given chaperone (Rosenzweig *et al.*, 2013; Nakamoto *et al.*, 2014).

Hsp90 family members have over a hundred identified substrates, but they also facilitate steroid hormones' receptor signaling. Activity of Hsp90 affects expression of a number of genes through the activation of protein kinases. Hsp90 level is kept constant under physiological conditions, however, it increases significantly under stress conditions. The main inducer of overproduction

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**Abbreviations:** ANOVA, analysis of variance statistical test; ELISA, enzyme-linked immunosorbent assay; Hsp, heat shock protein; JDP, J-domain protein

of Hsp90 is oxidative stress that occurs for example in reperfusion (Richter & Buchner, 2001; Picard, 2005; Caplan *et al.*, 2007; Nakamoto *et al.*, 2014). A bacterial homologue of human Hsp90 is HtpG.

Involvement of heat shock proteins in the pathogenesis and progression of atherosclerosis is undoubtedly complex. Traditional risk factors, such as metabolic disorders or hypertension, increase expression of the Hsp on the surface of endothelial cells in blood vessels. Hsp expression may in turn promote production of cytokines, increased expression of adhesion molecules, and the creation and promotion of the inflammatory response (Berberian *et al.*, 1990; Xu, 2002). The inflammatory response can also be induced due to cross reactivity between human and bacterial Hsp. Hsps also influence activity of the major enzymes of homocysteine metabolism — one of the main risk factor of stroke (Grabowski *et al.*, 2012). Induction of immune response against autologous or bacterial heat shock proteins on the surface of the vascular endothelium may lead to endothelial damage, followed by the development of atherosclerotic plaque, which is a cause of serious diseases — heart attack and stroke. They are also involved in development of auto-immunological diseases like rheumatoid arthritis, juvenile idiopathic arthritis and paraneoplastic syndrome (Nishizawa *et al.*, 1996).

## MATERIALS AND METHODS

**Specimen collection and preparation.** Study population included seventy four stroke patients hospitalized in the Clinic of Adult Neurology of the Medical University of Gdańsk (25 women and 50 men, mean age  $69 \pm 15$  years) and twenty one non-stroke members of the control group (25 cases, mean age  $43 \pm 12$  years). Information about presence of risk factors was collected from records and questionnaire based on interview with a given individual or closest relatives. Blood was collected in the Clinic of the Department of Neurology of the Medical University of Gdańsk (Sawula *et al.*, 2009). Plasma from each patient was obtained by centrifugation ( $2000 \times g$  for 15 min). Samples were frozen and stored at  $-70^\circ\text{C}$ .

**Protein purification.** Proteins were purified as described (Spence *et al.*, 1989; Woo *et al.*, 1992; Wawrzynów *et al.*, 1995; Banecki *et al.*, 1998).

**Enzyme-linked immunosorbent assay.** Antibody levels against particular proteins were measured by the ELISA test. Ninety-six-well microtiter plates were coated with GroEL, HtpG, DnaJ, DnaK and Hsp72. After 60 min of incubation at  $37^\circ\text{C}$ , plates were washed with PBS containing 0.05% Triton X-100. Non-specific binding was blocked with 1% BSA (Sigma, CAS Number 9048-46-8) in PBS. After another wash, 50  $\mu\text{l}$  of serum sample was added and incubated for 1 hour at  $37^\circ\text{C}$ . After washing with PBS containing 0.05% Triton X-100, the wells were incubated with secondary antibodies conjugated with horseradish peroxidase (Sigma, CAS Number 9003-99-0) for 1 h at  $37^\circ\text{C}$ . Then the plates were washed and developed with 50  $\mu\text{l}$  TMB and incubated for 10 min. The reaction was terminated by adding 50  $\mu\text{l}$  1M  $\text{H}_2\text{SO}_4$  and optical density was measured at 450 nm using an ELISA plate reader (Wallac Victor<sup>2</sup> 1420).

**Statistical analysis.** Data was analyzed using STATISTICA software for Windows v. 10 (StatSoft, USA) using one-way ANOVA statistics. A  $p$ -value of  $<0.05$  was estimated as statistically significant.

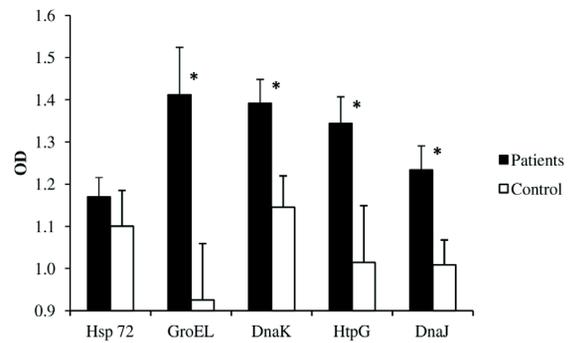


Figure 1. The relative antibody levels against Hsps in patients and control group.

\*Indicate  $p < 0.05$  compared to the control group (ANOVA).

## RESULTS

In our study we have demonstrated an increase in the level of anti-Hsp antibodies against bacterial heat shock proteins, however there was no statistically significant difference in anti-Hsp72 antibodies level between groups ( $p > 0.05$ ) (Fig. 1). Antibody level against Hsp72 was 1.10 in the control and 1.17 in the patient group. Interestingly, only for human recombinant Hsp72 the level of antibodies was not elevated. For all other studied bacterial proteins we have observed higher level of antibodies in stroke patient group. The mean of antibody level against DnaJ reached 1.01 in the control group and 1.23 in the patient group and it is a statistically significant difference ( $p = 0.047$ ). Anti-GroEL antibody level was 0.93 in the control group and 1.41 in the patient group. The group of patients had higher level of antibodies compared to the control group ( $p = 0.030$ ). The mean values of antibody level against DnaK in control and patient groups were 1.15 and 1.39 respectively, and by one-way ANOVA we determined that it is a statistically significant difference ( $p = 0.041$ ). Anti-HtpG antibody level was 1.01 in the control group and 1.34 in the patient group. The difference is statistically significant ( $p = 0.022$ ).

Our results confirm the widely accepted hypothesis that the Hsp-antibodies have an essential role in pathological mechanism of the atherosclerosis and stroke. We assume that previous bacterial infections and subsequent increase in antibody level are a risk factor of stroke. We have analyzed the correlation between risk factors of stroke and antibody levels (Table 1). Interestingly, the level of antibodies against all studied proteins increased in the group of patients that previously underwent an-

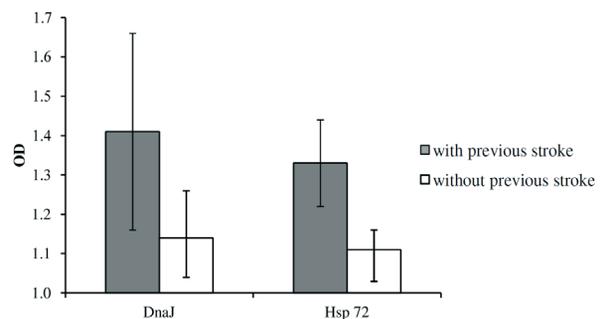


Figure 2. The relative antibody levels against Hsps in group of patients with and without previous stroke.

**Table 1. Relation between antibody level against Hsps and selected risk factors of stroke (*p*-values for ANOVA).**

Risk factors	Hsp72	DnaJ	HtpG	GroEL	DnaK
Previous stroke	0.046	<u>0.022</u>	0.326	0.145	0.145
Carotid artery stenosis	0.936	0.735	0.091	0.369	0.094
Myocardial infarction	0.904	0.828	0.062	0.562	0.395
Smoking	0.738	0.609	0.076	0.438	0.742
Stress	0.076	0.455	0.235	0.457	0.492
Depression	0.250	0.617	0.170	0.862	0.081
Migraine	0.162	0.338	0.479	0.101	0.743
Hypercholesterolemia	0.232	0.958	0.705	0.685	0.947
Coronary artery disease	0.238	0.974	0.255	0.612	0.678
Hypertension	0.428	0.506	0.615	0.849	0.109
Heart defects	0.489	0.253	0.902	0.434	0.356
Vascular diseases	0.532	0.795	0.675	0.642	0.225
Parkinson's disease	0.618	0.977	0.303	0.791	0.171
Atrial fibrillation	0.713	0.560	0.758	0.249	0.956
TIA	0.717	0.111	0.493	0.735	0.131
Kidney disease	0.769	0.335	0.880	0.139	0.924
Diabetes mellitus	0.826	0.117	0.325	0.627	0.348
Arrhythmia	0.866	0.915	0.728	0.107	0.633
Coffee consumption	0.873	0.982	0.486	0.244	0.485

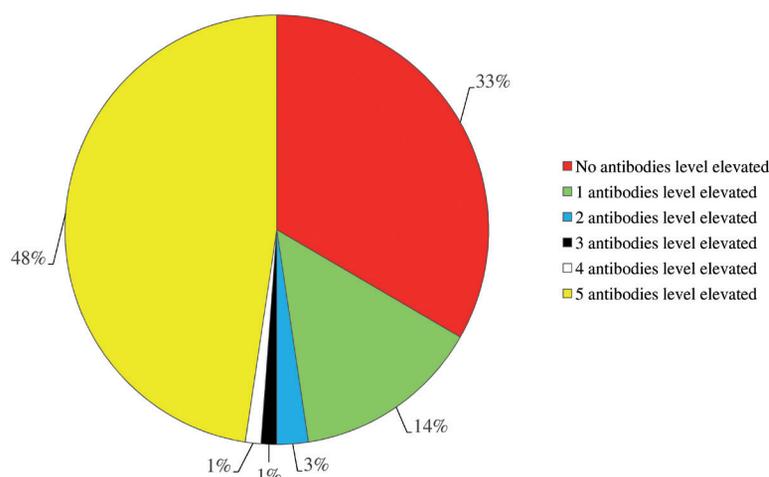
other stroke. Statistically significant difference was observed only in the level of antibodies against Hsp72 ( $p=0.046$ ) and DnaJ ( $p=0.022$ ) in the group of patients that did and did not undergo a previous stroke (Fig. 2). That may also indicate a significant role of antibodies against bacterial Hsps in stroke and after stroke patient recovery.

By comparing the distribution of the antibody level in patients, we can notice an interesting correlation (Fig. 3). Only 33 percent of subjects in the patient group did not show elevated levels of antibodies against the proteins

examined here, while 48 percent of patients have elevated levels of antibodies against all five studied proteins. Explanation of this result may be a prevalence of bacterial infections, leading by cross-reaction to formation of an autoimmune reaction directed against Hsp.

## DISCUSSION

Early stage of atherosclerosis is characterized by an immune-mediated reaction, which may be caused by local autoantigens (Sigal, 2007). It has been described in the literature that Hsp and the oxidized low-density-lipoproteins (oxLDL) are the main autoantigens involved in the pathogenesis of atherosclerosis (Blasi, 2008). The role of Hsps in atherogenesis is not clear, although previous studies have demonstrated protection of endothelial cells from stress by Hsps (Johnson *et al.*, 1993). Also, it has been shown that Hsps can stimulate plaque formation (Seitz *et al.*, 1996). Presently it is assumed that the antibodies against microbial Hsps can be misdirected towards host Hsps overexpressed in the stressed cells of the vascular endothelium. In consequence, that may promote the plaque formation. This autoimmune reaction is possible because of the amino acid sequence similarity between microbial and human Hsp. While the bacterial Hsps have 97% ho-



**Figure 3. The percent of patients with elevated levels of antibodies anti-Hsp.**

mology between species, bacterial and human Hsps have more than 50% homology (Young, 1992).

The DnaJ protein (Hsp40) is a co-chaperone for DnaK (Hsp70). It has been shown that the level of human Hsp40-HDJ-2 is significantly increased in atherosclerotic carotid artery and correlates with luminal stenosis in ulcerated atherosclerosis. It has been also shown that HDJ-2 and Hsp70 are present in atherosclerotic carotid artery plaques and expressed independently. Since bacterial DnaJ's N-terminal region is highly conserved between species, it is possible that an immunological cross-reaction between human Hsps and bacterial DnaJ-DnaK system takes place by shared B- and T-cell epitopes.

Another well-studied bacterial protein, GroEL, belongs to the Hsp60 family. The Hsp60 family is one of the most interesting groups of Hsps because of its wide range of effects on the development of autoimmune diseases. The Hsp60 family has been demonstrated to be involved in the development of rheumatoid arthritis in humans, adjuvant arthritis and insulin-dependent diabetes mellitus in rodents. Also, microbial Hsp60s are involved in pathogenesis of systemic sclerosis, psoriasis, Kawasaki disease, and Behcet's disease (Perschinka *et al.*, 2003). It has been also shown that Hsp60 has a pathogenic role in cellular immune reactions in early human atherosclerosis (Knoflach *et al.*, 2007). Bacterial Hsp60 stimulates the release of pro-inflammatory cytokines and cell adhesion in atherosclerotic lesions (Rupinder *et al.*, 2001). It has been shown as well that antibodies that display cross-reactivity between human and microbial Hsp60 are related to atherosclerosis (Xu *et al.*, 1999). Hsp60 localizes in atherosclerotic lesions of the arterial wall, whereas in non-atherosclerotic regions Hsp60 is not observed (Kol *et al.*, 1998). Also, Hsp60 reactive T-lymphocytes promote atherosclerosis development (Almanzar *et al.*, 2012). It was demonstrated that bacterial Hsp60 is a major epitope that could cause autoimmunity for producing antibodies reactive to their human counterpart (Okada *et al.*, 2007). Also, periodontopathic bacterial infection elevates levels of cross reactive antibodies between human Hsp60 and bacterial GroEL (Tabeta *et al.*, 2000). In another study it was shown that there is a correlation between level of antibodies against Hsp60 and prevalence of coronary artery disease (Knowlton *et al.*, 2008).

The presence of antibodies against GroEL in the group of stroke patients could be caused by previous bacterial infections. The acute phase of stroke induces stressful conditions with subsequent increase of Hsp's expression. Because of high concentration of Hsps in plasma, the antibodies can act less specifically. Therefore, given the similarity of amino acid sequence between bacterial and human Hsps, it is highly possible that the immune response to Hsps derived from pathogens might cross-react with host Hsps.

Similar assumption as we described above for the GroEL protein can be also made in relation to DnaK because of the close structural homology with human Hsp. It has been shown that *E. coli* DnaK and human Hsp70 are 50% identical (Man-Un Ung *et al.*, 2013) at the amino acid sequence level, so the immunological cross-reactivity is also possible in this instance. Hsp70 may play a dual function in atherosclerosis. On one hand Hsp70 may act as a cytoprotector, a chaperone and play a role as an anti-apoptotic protein, reducing the level of Bax and AIF proteins synthesized in the cells of hypoxic tissue. (Goel *et al.*, 2010). On the other, may stimulate the innate immune response and promote inflammation (Dvorianchikova *et al.*, 2014).

It has been shown that there is a correlation between anti-Hsp70 antibody level and the presence of vascular diseases such as claudication, critical ischemia and aneurysmal disease (Chan *et al.*, 1999). It is interesting that HSP70 mRNA expression during heat shock was higher than during postischemic reperfusion (Nishizawa *et al.*, 1996). Hsp72 overexpression during stroke and its similarity to prokaryotic Hsp70 can also lead to immunological cross reaction, but levels of antibodies against Hsp72 display no differences between the patient and control groups.

We also measured antibodies level in plasma against the HspG protein which is a distant eukaryotic relative of the human Hsp90 and actually is not well characterized. HspG has about 40% sequence identity and 55% similarity to its eukaryotic counterparts (Qing Huai *et al.*, 2005). Human Hsp90 regulates the conformation, activation and function of over 100 cellular proteins. It has been shown that Hsp90 expression level is higher in atherosclerotic plaques compared with expression in normal artery. Hsp90 induces an immune response which makes Hsp90 a target autoantigen in the pathogenesis of atherosclerosis (Businaro *et al.*, 2009). It is supposed that Hsp90, like other Hsps, is sustaining inflammatory mechanisms underlying pathogenesis of atherosclerosis (Benagiano *et al.*, 2003). Reperfusion induces expression of Hsp90 in the heart, brain and kidney. Hsp90 overexpression in the ischemic region can preserve the tissue from damage through an exquisite stimulation of the endothelial NO pathway (Kupatt *et al.*, 2004) and maintain activity of the proteasome in hypoxic cells (Hota *et al.*, 2010). It has been observed that the HSP90 mRNA expression was significantly higher in reperfused hearts than in heat-shocked hearts (Nishizawa *et al.*, 1996). It is possible that after stroke the level of Hsp90 suddenly increases and antigen-presenting cells produced during the earlier bacterial infection can redirect immune response against native Hsp90.

Possibly the high concentration of anti-Hsp72, anti-DnaJ and other antibodies in plasma after the first stroke cause increased immune response against these proteins during another stroke. The more intensified immune response against Hsp72 may be caused by increased expression against this protein during the first stroke and next autoimmunization. Increased level of antibodies against DnaJ in the group of stroke patients may be related to its participation in Hsp40/Hsp70 co-chaperone system. Increased expression of human Hsp40/Hsp70 co-chaperone system during the stroke may lead to autoimmunization against human Hsp40 and in consequence may cause the immunological cross-reaction against bacterial DnaJ.

Our results suggest an essential role of Hsps in development of the atherosclerosis and stroke. Still, the exact role of Hsps in autoimmune diseases and stroke remains unclear and requires further studies.

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