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# T and NK cell subset changes with microbial extracts and human HSP60-derived peptides in Behçet's disease

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**Key words:** Behçet's disease, streptococcus, *Escherichia coli*, human HSP60.

## ABSTRACT

**Objective.** Microorganisms such as streptococcus and autoantigens such as 60 kD heat-shock protein (HSP60) are implicated in the etiopathogenesis of Behçet's disease (BD).

**Methods.** Peripheral blood mononuclear cells from patients with BD (n = 16) and healthy controls (HC) (n = 11) were cultured for 5 days with extracts of *S. sanguis*-KTH-1 (SS), *E. coli* (EC) and a mixed peptide combination from human HSP60 (aa 136-50, 179-97, 224-58 and 336-51) reported to be associated with BD. T and NK cell subset changes were determined by flow cytometry.

**Results.** In unstimulated 5-day cultures  $\gamma\delta^+$  (both  $CD4^+\gamma\delta^+$  and  $CD8^+\gamma\delta^+$ ),  $CD8^+\alpha\beta^+$ ,  $CD4^+CD56^+$  and  $CD8^+CD11b^+$  cells were increased in BD compared to HC. In antigen-stimulated cultures of BD patients  $CD3^+$  and  $\alpha\beta^+$  T cells responded to HSP60 peptides whereas EC stimulated only  $CD16/CD56^+$  NK cells. In the control group, similar to BD,  $\alpha\beta^+$  and  $CD4^+$  T cells responded to HSP60 peptides, however SS and EC mainly activated cytotoxic T cell subsets ( $CD8^+CD11b$  and  $CD4^+CD56^+$  T cells).

**Conclusion.** Significant increases in unstimulated T cell subsets suggest the presence of an in vivo T cell activation in BD. In both patients and controls similar patterns of responses were observed against different microorganisms, however the role of human HSP 60 peptides as immunodominant, cross-reactive antigens could not be demonstrated.

## Introduction

Various microorganisms such as Streptococcus and Herpes Simplex virus are implicated in the etiopathogenesis of Behçet's disease (BD), a systemic vasculitis of unknown etiology (1, 2). Clinical observations such as the presence of oral aphthae as the first manifestation in 70% of the patients, increased oral

manifestations after dental manipulations, hypersensitivity to streptococcal skin tests, dominance of atypical streptococcal species in BD patients' oral flora, and the recent reports of beneficial anti-bacterial therapy put forward the role of streptococcus in BD (3-8).

In elucidating the immune mechanisms of streptococcal pathology, KTH-1 (a crude extract of *Streptococcus sanguis* SSH-83, SS) has been found to stimulate IL-6 and IFN- $\gamma$  secretion by the peripheral blood T cells of BD patients (9). Oral bacterial flora is also shown to activate  $\alpha\beta^+$  T cells in BD (10). KTH-1 upregulates  $\alpha\beta^+$  T cells in short-term T cell cultures and KTH-1 specific  $\alpha\beta^+$  T-cell lines secrete pro-inflammatory mediators IL-6, IL-8 and TNF- $\alpha$  (11,12). The main antigenic stimuli of *S. sanguis* is suggested to be a 95 kD BES-1 gene product consisting of 849 amino acid residues (13).

However, the specificity of the anti-streptococcal response has not been studied in detail in comparison to other microorganisms such as gram negatives in BD (2). Although not significantly different compared to controls, *Escherichia coli* (EC) also caused a higher incidence of skin reactions in BD patients (4). In addition to streptococcal antigens, *E. coli* and *Staphylococcus aureus* also activate BD lymphocytes to release increased amounts of IFN- $\gamma$  and IL-6 (9). Behçet's T lymphocytes secrete IFN- $\gamma$  in response to very low (1-10 pg/ml) doses of Staphylococcal superantigens SEB and SEC1 compared to controls and Hirohata *et al.* proposed a T cell hyperreactivity in BD, unrelated to a species-specific response (14). Heat-shock proteins (HSP) are highly conserved proteins among microorganisms and mammals and HSP60/65 has been suggested as an immunodominant antigen in BD, associated with the "cross-reactivity" of microbial HSP60 with its human homologue (3, 15). Anti-mycobacterial HSP65 antibodies

are shown to be cross-reactive with oral mucosal homogenates and oral streptococcus (16). Four epitopes of mycobacterial HSP65 (amino acid sequences 111-25, 154-72, 219-33 and 311-26) and their human counterparts with 50-80% homology, were recognized to be immuno-dominant antigens for T and B cell responses in BD in studies from UK, Japan and Turkey (15, 17-19). With this background, in this study we aimed to investigate the effects of microbial (*E. coli* and *S. sanguis*) and auto immune stimulation with human HSP-60 peptides on peripheral blood lymphocyte subsets in BD patients to test whether anti-microbial responses against streptococcus are specific and associated with human HSP60 responses.

## Materials and methods

### Patients and controls

Sixteen patients with BD (6 males, 10 females; mean age: 35, range 24-51), diagnosed according to the International Study Group Criteria for BD (20) and followed in the multidisciplinary Behçet's Disease Out-patient Clinic at Marmara University Hospital in Istanbul were studied. At the time of the study, all patients had clinically active disease with at least two clinical manifestations. All patients had a history of oral ulcers, 13 (81%) had folliculitis, 12 (75%) had genital ulcers, 10 (63%) had erythema nodosum, 5 (31%) had venous thrombosis and 4 each (25%) had arthritis or ocular involvement. The pathergy test was positive in 9 (56%) of the patients. Nine patients were HLA-B51-positive. Eleven patients were treated with colchicine (1-2 mg/day), whereas no patients were using immunosuppressive treatment at the time of sampling. Eleven healthy volunteers (6 males, 5 females; mean age: 33, range 21-46) with no signs or symptoms of BD or any other disease, were studied as normal controls. The study was approved by the Institutional Review Board and informed consent was obtained from all patients.

### Preparation of antigenic stimulants

*Streptococcus sanguis* SSH-83 (KTH-1) were kindly provided by F. Kaneko, Japan. Bacteria were washed 3 times

by centrifugation, re-suspended in PBS, and incubated at 42°C for 30 minutes to stimulate HSP expression. After killing at 85°C for 20 minutes, whole bacterial cells were subjected to sonification for obtaining their soluble extracts. Extracts were then centrifuged and washed twice with PBS. Protein contents of the extracts were estimated according to Lowry method. In each experiment, 10 mg/ml of soluble bacterial extracts were used.

Human HSP60 derived peptides 136-150, 179-197, 224-258 and 336-351, previously demonstrated to be immunodominant in BD (17,19), were synthesized (Albachem, Ireland). As T-cell responses to single peptides were low in frequency and magnitude (19), we mixed the four peptides together in equal concentrations of 50 mgr each in serum-free RPMI-1640 medium.

### Cell separation and culture

Blood was taken from the patients and controls into heparinized tubes and peripheral blood mononuclear cells (PBMNCs) were separated by standard Hypaque/Ficoll density gradient centrifugation. Cell counts were determined by using automatic blood count analyser (Coulter, USA). Serum samples were also isolated using heparin-free tubes and inactivated at 56°C for 30 minutes. All cultures were carried out in RPMI-1640 medium supplemented with 1% Penicillin-Streptomycin, 1% Sodium pyruvate, 0.3 mg/ml L-Glutamine and 10% freshly prepared autologous serum. 3-4 x 10<sup>6</sup> T-cells and monocytes were cultured in flat-bottom 6-well plates (Greiner, Germany) in a total volume of 3 ml in the presence or absence of antigenic stimulators. The cells were incubated for 5 days until the immunophenotypical analysis at 37°C in a humidified atmosphere of 5% CO<sub>2</sub>.

### Immunophenotyping

After 5-day incubation period, non-adherent cells were harvested and washed twice with PBS. Cells were resuspended in PBS and incubated with saturating amount of appropriate antibodies for 30 min at room temperature. Flow cytometric analysis were performed using a FACSort instrument (Becton Dick-

inson, Mountainview, CA). Following MoAbs were used for double stainings; anti-CD3, anti-CD4, anti-CD8, anti-CD11b, anti-CD16, anti-CD56, anti-TCR and anti- $\gamma$  TCR. Background fluorescence was eliminated by running isotype matched PE- and FITC-conjugated mouse IgGs as negative controls (Pharmingen, San Diego, CA). Data were acquired and analyzed using the Cell Quest software. The lymphocyte population was identified on a scatter plot, based on forward angle light scatter and 90° light scatter. To eliminate cell debris and electronic noise contaminating the lymphocyte gated population, a leucocyte common antigen, anti-CD45, was used. An anti-CD14 was used to examine monocyte contamination which was consistently less than 1.0%. For dual color fluorescence a cytogram was generated and 20000 cells were analyzed.

### Statistical analysis

Kruskal-Wallis and Mann Whitney-U tests are used for statistical analysis. P < 0.05 was considered as significant.

## Results

In unstimulated cultures,  $\gamma$  T cells, (both CD4<sup>+</sup>  $\gamma$  and CD8<sup>+</sup>  $\gamma$  subsets), CD8<sup>+</sup>  $\gamma$ , CD4<sup>+</sup>CD56<sup>+</sup> and CD8<sup>+</sup>CD11b<sup>+</sup> T lymphocyte subsets were found to be upregulated significantly in BD compared to HC after 5-day incubations (Table I). Only  $\gamma$ /CD4<sup>+</sup> T cells were significantly higher in HC compared to BD (p=0.02). Although not reaching statistical significance, CD16<sup>+</sup>/CD56<sup>+</sup> NK subset were also elevated in HC (p = 0.08).

Although mild changes were also observed in  $\gamma$ ,  $\gamma$  and CD8<sup>+</sup>CD11b<sup>+</sup> T cell subsets, only CD16<sup>+</sup>CD56<sup>+</sup> NK cells increased significantly with *E. coli* stimulation in BD (Table II). A similar trend was also observed with *S. sanguis*, without reaching statistical significance. On the other hand, *E. coli* and *S. sanguis* both increased CD56<sup>+</sup> (especially of CD4<sup>+</sup>CD56<sup>+</sup>T cells) and CD8<sup>+</sup>CD11b<sup>+</sup> cytotoxic T cell subsets in HC (Table III).  $\gamma$ CD8<sup>+</sup> T cells are also increased significantly with *S. sanguis* in HC, with a similar but not significant trend with *E. coli*.

**Table I.** T and NK cell subsets in BD patients and controls after unstimulated (medium only) 5-day cultures (mean ± SD).

Cell subsets	Behçet's disease (n = 16)	Healthy controls (n = 11)	p
CD3	79.1 ± 9.2	71.0 ± 12.1	
CD4	50.7 ± 10.9	51.8 ± 2.8	
CD8	35.0 ± 5.7	30.1 ± 6.5	
CD11b	11.4 ± 8.6	4.7 ± 3.3	0.01
CD16	11.7 ± 7.0	14.3 ± 7.4	
CD56	12.6 ± 3.7	12.9 ± 5.5	
TCR	73.5 ± 11.5	70.5 ± 4.6	
TCR	5.9 ± 3.9	2.8 ± 1.4	0.01
/ CD4	39.8 ± 11.9	48.1 ± 5.8	0.02
/ CD4	1.9 ± 0.7	0.7 ± 0.6	0.04
/ CD8	27.8 ± 6.9	22.5 ± 4.0	0.03
/ CD8	5.6 ± 4.4	1.4 ± 1.1	0.002
CD16 / CD56	6.2 ± 2.6	8.5 ± 3.6	
CD4 / CD16	0.9 ± 0.8	0.9 ± 0.3	
CD4 / CD56	2.7 ± 1.7	1.3 ± 0.4	0.03
CD8 / CD11b	7.1 ± 5.1	2.1 ± 1.5	0.008

**Table II.** T and NK cell subsets after *in vitro* stimulation in patients with Behçet's disease (mean ± SD).

	Medium only	<i>E. coli</i>	<i>S. sanguis</i>	HSPPeptide Mix
CD3	79.1 ± 9.2	81.4 ± 1.3	80.9 ± 10.0	86.9 ± 9.00*
CD4	50.7 ± 10.9	50.6 ± 9.8	51.0 ± 8.2	55.6 ± 11.6
CD8	35.0 ± 5.7	34.5 ± 7.6	36.7 ± 8.6	33.3 ± 7.7
CD11b	11.4 ± 8.6	12.3 ± 7.1	11.6 ± 7.2	9.2 ± 6.9
CD16	11.7 ± 7.0	13.2 ± 6.5	13.9 ± 6.5	10.4 ± 7.7
CD56	12.6 ± 3.7	14.1 ± 3.6	14.1 ± 4.8	11.7 ± 5.5
TCR	73.5 ± 11.5	75.0 ± 12.3	75.3 ± 12.6	80.8 ± 13.1*
TCR	5.9 ± 4.0	7.9 ± 3.7	7.5 ± 4.6	7.7 ± 5.1
/ CD4	39.8 ± 12.0	44.7 ± 11.4	43.0 ± 12.9	48.9 ± 13.5**
/ CD4	1.9 ± 0.7	2.2 ± 2.5	1.3 ± 1.4	1.8 ± 3.1
/ CD8	27.8 ± 6.9	27.2 ± 7.5	29.9 ± 9.1	28.6 ± 8.3
/ CD8	5.6 ± 4.4	7.5 ± 5.5	7.9 ± 5.4	7.6 ± 9.9
CD16 / CD56	6.2 ± 2.6	8.0 ± 3.7 *	7.9 ± 4.0	5.5 ± 3.8
CD4 / CD16	0.9 ± 0.8	0.8 ± 0.4	0.8 ± 0.6	0.9 ± 0.4
CD4 / CD56	2.73 ± 1.73	2.6 ± 1.7	3.1 ± 2.7	2.9 ± 1.9
CD8 / CD11b	7.1 ± 5.1	10.0 ± 7.0	8.7 ± 5.5	6.5 ± 4.3

\*p < 0.05; \*\*p = 0.06 – 0.07.

With HSP60 peptide stimulation, only CD4<sup>+</sup> T cell subset (mainly of CD4<sup>+</sup> T cells) increased significantly in BD. In HC, in addition to CD4<sup>+</sup>CD56<sup>+</sup> T cells are also increased compared to unstimulated cultures. Although mild elevations were observed, changes in CD4<sup>+</sup> T cell subsets did not reach significance in both groups. There was also no correlation between the individual HLA-B51 expressions, colchicine treat-

ment and cell surface antigen expressions after stimulation (data not shown).

**Discussion**

The innate immune system detects infection using a variety of pattern-recognition receptors that recognize pathogen-associated molecular patterns on microorganisms and trigger various effector responses. Interaction with and control of the adaptive immune system

is also a crucial role of the innate immunity. Viral or streptococcal infections have been suggested to be associated with BD and a dysfunction of the innate immune system might predispose to persistent immune activation after infectious stimuli. Abnormal adaptive immune responses, such as tissue lymphocyte/phagocyte infiltrations and a Th1-type, pro-inflammatory cytokine profile with enhanced IL-12 and IFN expressions are then induced (2,15). In this context, lymphocyte and NK-mediated cytotoxic responses are possibly also augmented in BD.

In our study, BD patients had higher levels of both CD4<sup>+</sup> and CD8<sup>+</sup> T cells in addition to cytotoxic-suppressor T cell subsets CD8<sup>+</sup>, CD4<sup>+</sup>CD56<sup>+</sup> and CD8<sup>+</sup>CD11b<sup>+</sup> T cells compared to healthy controls in unstimulated 5-day cultures. These results are in agreement with our previous report of increased CD4<sup>+</sup>CD56<sup>+</sup> T cells in BD in unstimulated, directly analysed PBMCs (21).

Two different cell populations, CD8<sup>+</sup> T cells and NK cells, mediate the cytotoxic function of the immune system. Recent studies have shown that a subset of CD4<sup>+</sup> T cells can also have cytotoxic activity (22). This subset expressing CD56 antigen is generally found in close proximity to inflammatory sites and only a small proportion has been detected in the peripheral blood (23, 24). Therefore, CD4<sup>+</sup>CD56<sup>+</sup> phenotype is regarded to be associated with organ and dermal associated cytotoxic T cells (25).

When we compared responses against two different microorganisms *S. sanguis* and *E. coli*, although similar trends were observed in most subsets, the only subset with a significant response was CD16<sup>+</sup>CD56<sup>+</sup> NK cells with *E. coli*. The role of NK cells is controversial in BD, with normal, decreased or elevated reports in the peripheral blood (21, 26, 27). NK cell activity measured by K562 cytotoxicity was observed to be low, explained by immature NK cells in active patients (27). However, increased levels of NK cells after bacterial stimulation is not previously reported in BD. The similarity of responses with two different microorganisms suggest that this is possibly a general anti-

**Table III.** Tand NK cell subsets after *in vitro* stimulation in healthy controls (mean ± SD).

	Medium only	<i>E. coli</i>	<i>S. sanguis</i>	HSPPeptide Mix
CD3	71.0 ± 12.1	74.2 ± 10.9	76.7 ± 12.5	79.5 ± 9.7*
CD4	51.8 ± 2.8	51.6 ± 4.1	54.9 ± 5.0	56.5 ± 5.7*
CD8	30.1 ± 6.5	31.9 ± 6.8	31.8 ± 6.9	29.2 ± 6.9
CD11b	4.7 ± 3.3	7.0 ± 2.4**	6.8 ± 2.8**	5.3 ± 2.1
CD16	14.3 ± 7.4	16.0 ± 8.9	15.6 ± 7.9	12.2 ± 6.6
CD56	12.9 ± 5.5	18.2 ± 6.2*	18.9 ± 4.3*	13.6 ± 4.4
TCR	70.5 ± 4.6	72.0 ± 7.4	75.5 ± 7.7	77.9 ± 5.5*
TCR	2.8 ± 1.4	4.0 ± 2.3	3.7 ± 2.3	3.91 ± 3.1
/ CD4	48.1 ± 5.8	47.9 ± 5.0	51.6 ± 4.4	47.5 ± 15.5
/ CD4	0.7 ± 0.6	1.1 ± 1.0	1.4 ± 1.5	1.0 ± 0.6
/ CD8	22.5 ± 3.9	22.9 ± 4.3	24.2 ± 5.4	24.7 ± 5.1
/ CD8	1.3 ± 1.1	2.6 ± 2.0	2.5 ± 2.2**	2.0 ± 2.2
CD16 / CD56	8.5 ± 3.6	11.2 ± 5.9	10.8 ± 5.6	6.9 ± 3.6
CD4 / CD16	0.9 ± 0.3	0.9 ± 0.3	0.9 ± 0.3	1.9 ± 1.4
CD4 / CD56	1.3 ± 0.4	2.3 ± 1.7*	2.5 ± 1.1*	2.5 ± 2.4**
CD8 / CD11b	2.1 ± 1.5	3.5 ± 1.3*	3.4 ± 1.6*	2.7 ± 0.9

\*p < 0.05, \*\*p = 0.06 – 0.07.

bacterial response which might be of innate nature.

On the other hand, responses in HC showed a similar pattern for both *S. sanguis* and *E. coli* with increases of mainly cytotoxic subsets CD8<sup>+</sup>CD11b<sup>+</sup>, CD4<sup>+</sup>CD56<sup>+</sup> and CD8<sup>+</sup> T lymphocytes. As an interesting observation, the same subsets are elevated in BD patients' peripheral blood without "stimulation", suggesting an *in vivo* activation of lymphocytes which may limit their proliferation capacity further in *in vitro* conditions.

Elevations of T cells in unstimulated samples in our study are also similar to most, but not all studies done in BD in directly analysed PBMCs (10, 28, 29). However, unlike previous reports, we could not observe increased T cell subsets after *S. sanguis* stimulation (10-12). Similar to a previous study from Japan, we observed mainly TCR-<sup>+</sup> responses against HSP60 peptides in our study, with responses of CD4<sup>+</sup>CD56<sup>+</sup> cytotoxic T cells also in HC. This is in contrast to Hasan *et al.* whom reported significant T cell responses to HSP60 derived peptides in BD patients from UK (17). This discrepancy might be explained with the background level of T cells which is observed to be different according to ethnic origin. A higher percentage of T

cells were reported from the individuals of Mediterranean origin compared to Scandinavians (30). However, we also have not observed a very high level of T cells in our healthy controls and elevated levels of T cells mainly in unstimulated samples in our BD patients suggest that antigens other than HSPs may upregulate this subset in our population. In this respect, non-peptide prenyl pyrophosphates are recently described as the dominant antigens for T cell lines derived from intra-ocular fluid of uveitis patients with BD (31).

Possibly the most important part of our data is the similarity of immune responses against two different (gram positive and negative) microorganisms, revealing the problem of the specificity of anti-streptococcal immunity in BD. As Hirohata *et al.* first proposed (14), antigens possibly common to a variety of different microorganisms might be immuno-dominant in BD. In this respect, highly conserved heat shock proteins are logical candidates. However, the differences between anti-HSP responses and bacterial extracts suggest that antigens other than HSP60 are possibly dominant for immune recognition of *S. sanguis* and *E. coli*. In this respect, further analysis of other bacterial antigens such as BES-1 gene product in

different ethnic populations is required (13).

There are some points in our study which are to be taken into consideration in the interpretation of our data. We have not performed a direct, freshly isolated unstimulated analysis of PBMCs and our 5-day unstimulated cultures might have down- or upregulating effects on certain cell surface expressions. This makes comparing our data difficult with most studies having analysis of unstimulated samples done immediately.

In conclusion, our results suggest that responses to whole bacteria extracts do not correlate with anti-HSP60 peptide responses which are mainly recognized by T cells in BD. We also found no evidence of a species-specific response against *S. sanguis* in our population. Further studies on anti-bacterial immune responses in BD should also include control microorganisms such as *E. coli* to clarify the role of infectious agents in the pathogenesis.

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