T and B lymphocyte subpopulations and activation/differentiation markers in patients with selective IgA deficiency

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Summary

Selective deficiency of immunoglobulin A (IgAD) and common variable immunodeficiency (CVID) are genetically closely related diseases, both of unknown pathogenesis. A plethora of abnormalities in lymphocyte subpopulations and expression of activation markers were repeatedly documented in CVID patients, while almost no data are available about lymphocyte subpopulations in IgAD patients. We determined basic lymphocyte subpopulations and those subpopulations that were reported to be abnormal in CVID patients (CD25, human leucocyte antigen (HLA)-DR CD45RA, CD45RO, CD27, CD28 and CD29 on both CD4⁺ and CD8⁺ cells, CD57 and CD38 on CD8⁺ cells, CD21, CD27, IgM, IgD on B lymphocytes) in 85 patients with IgAD, 47 patients with CVID and in 65 healthy controls. Statistical analysis was performed by the Mann-Whitney U-test; significant P-values were determined by means of Bonferoni's correction. Our results showed an increase in the relative number of CD8⁺ cells and a decrease in the absolute number of CD4⁺ cells compared to healthy people, but similar abnormalities in CVID patients were much more expressed. IgAD patients had significantly decreased expression of HLA-DR and increased expression of CD25 on CD4⁺ lymphocytes, also CD29 expression was decreased on CD8+ cells, while other activation/differentiation markers on T cells (including the expression of CD45RA and CD45RO antigens) were not changed. There were no statistically significant abnormalities in B lymphocyte developmental stages in IgAD patients compared to healthy controls. Our observation showed that the majority of T and B lymphocyte subpopulation abnormalities described previously in CVID are not present in IgAD patients.

Keywords: common variable immunodeficiency, B lymphocyte subsets, differentiation antigens, IgA deficiency, T lymphocyte subsets

Introduction

Selective deficiency of immunoglobulin A (IgAD) is the most frequent, but usually asymptomatic, abnormality of immunoglobulin production, with a frequency ranging from 1:278 to 1:3040 in Caucasians [1]. Despite its high frequency, very little is known about the aetiopathogenic mechanisms leading to this immunological abnormality.

IgAD is linked to a more severe immunoglobulin production disease – common variable immunodeficiency (CVID), which is characterized by decreased IgG and IgA levels, while IgM levels are variable. Clinically significant immunodeficiency evolves at any age after apparently normal function of the immune system [2]. Familial occurrence of both diseases has been documented repeatedly [3,4]. Deficiency of transmembrane activator and CALM interactor (TACI) was observed both in CVID and IgAD patients [5,6]. All these results show that IgAD and CVID may be different phenotypic manifestations of the same, or similar, genetic background. Repeatedly documented progression of IgAD into CVID [7] shows that IgAD may be a developmental stage in evolving CVID.

Although the pathogenesis of CVID has not yet been elucidated, many abnormalities in the number or function of immune cells were described in CVID patients. Great attention has been given to abnormalities in lymphocyte subpopulations. Defective activation of T lymphocytes [8–10] and impaired cytokine production [11–13] were observed in CVID patients. Various T subpopulation abnormalities including a decreased number of CD4⁺ cells [14–16], reduced expression of CD45RA [15] and CD62L [17] or increased expression of CD45RO [15] on CD4⁺ cells, human leucocyte antigen (HLA)-DR [16] and CD57 on CD8⁺ [18] were reported. Also, decreased expression of CD27, CD28, CD29 on both CD4⁺ and CD8⁺ cells and increased expression of CD38 and CD57 on CD8⁺ cells were observed [19]. When evaluating B lymphocyte subpopulations, a decrease in the number of memory CD27⁺ IgD⁻ IgM B cells has been described repeatedly [20–24].

Despite the assumed relation of CVID and IgAD, there are almost no data available about T and B lymphocyte subpopulations in IgAD. Previous studies performed on limited numbers of patients showed a possible decrease of the proportion of CD4⁺ cells and increase of proportion of CD8⁺ cells [25–27], while Klemola *et al.* [28] did not observe any significant abnormalities in relative numbers of CD3⁺, CD4⁺ and CD8⁺ cells. Because of the above-mentioned similar genetic background of CVID and IgAD, assessing those abnormalities in IgAD may help to elucidate whether lymphocyte abnormalities in CVID are primary, inherited or whether they evolve secondarily during the development of the diseases, or are even a consequence of immunodeficiency in CVID.

In our study we have evaluated lymphocyte subpopulations most frequently known to be abnormal in CVID patients in a group of 85 patients with IgAD, 65 controls and 47 patients with CVID.

Patients and methods

Eighty-five patients with IgAD fulfilling diagnostic criteria of IgAD [29] (48 females, 37 males) aged 18-72 years were included in the study. These patients were referred most frequently for immunological investigation because of apparently increased frequency of mild respiratory tract infections; none of them suffered from clinically significant immunodeficiency or clinically apparent autoimmune disease. Lymphocyte subpopulations were also determined in 47 patients with CVID (30 females, 17 males) aged 10-75 years. A detailed analysis of relations of lymphocyte subpopulations in the CVID subgroup has been published previously [19]. As a control group, 65 healthy people with normal serum IgA levels (35 females, 30 males) aged 22-66 years were used. Controls were recruited from hospital staff and medical students. All blood samples were obtained in an acute infection-free period; in CVID patients on IVIG treatment, blood samples were obtained prior to immunoglobulin infusion.

The study was approved by the Ethics Committee of the Faculty of Medicine of Masaryk University in Brno. Informed consent was obtained from all investigated individuals before the samples were drawn.

Lymphocyte subpopulation determination

All blood samples were taken between 8 and 11 a.m. to exclude diurnal variation of lymphocyte subsets. B lymphocyte subpopulations were analysed from density gradient separated cells while T lymphocyte subpopulations were determined from non-separated blood, as described previously [19]. Immunophenotyping of lymphocytes was performed by four-colour cytometry Epics MCL (Beckman Coulter Miami, FL, USA).

Major lymphocyte populations were determined by two cocktails of monoclonal antibodies (MoAb): 1, anti-CD45 fluorescein isothiocyanate (FITC), anti-CD4 phycoerythrin (PE), anti-CD8 phycoerythrin-Texas red X (ECD) and anti-CD3PC5; 2, anti-CD45FITC, anti-CD56PE, anti-CD19ECD, anti-CD3 r-phycoerythrin-cyanine 5 (PC5); (Cyto-Stat tetraCHROME; Beckman Coulter, Miami, FL, USA) and anti-CD16PE (Immunotech, Marseille, France). In addition, the following MoAbs were used: PE-conjugated anti-CD38, CD45RA, CD45RO, FITC-conjugated anti-CD25, CD27, CD28, CD29, CD57, CD62L, PC5-conjugated anti-CD3 (all from Dako A/S, Glostrup, Denmark); PC5-conjugated anti-CD27, ECD-conjugated anti-CD4, CD8, CD19 (all from Beckman Coulter); FITC-conjugated anti-CD38, PE-conjugated anti-IgD and anti-CD21, PC5-conjugated anti-IgM (all from Pharmingen International, San Diego, CA, USA) and PE-conjugated anti-HLA-DR (Becton Dickinson, San Jose, CA, USA).

Statistical analysis

Normality of the obtained lymphocyte subpopulations was assessed by the Kolmogorov–Smirnov test. Because the majority of the obtained values showed a distribution different from normal, a non-parametric Mann–Whitney *U*-test was used. Critical values for statistical significance were determined by Bonferroni's correction [30] for each group of lymphocytes: absolute and relative numbers of major lymphocyte subpopulations, CD4⁺ activation/differentiation markers, CD8⁺ activation/differentiation markers, absolute and relative numbers of B cell activation states. Statistical package STATISTICA (StatSoft, Inc.) version 7 was used.

Results

The absolute and relative numbers of CD3⁺, CD4⁺ and CD8⁺ T lymphocytes in IgAD patients, healthy controls and CVID patients are shown in Tables 1 and 2. As can be seen, the absolute number of CD4⁺ cells was decreased and the relative number of CD8⁺ cells was increased in IgAD patients compared to healthy controls. Comparing CVID patients with IgAD patients, the relative numbers of CD3⁺, CD8⁺ cells were increased in CVID patients while CD4⁺ and natural killer (NK) (CD16/56⁺ CD3⁻) cells were decreased compared to IgAD patients. Evaluating absolute numbers, the number of

(s.d.). Lymphocytes are expressed as percentage of CD45 cens, lymphocyte subpopulations as percentage of lymphocytes.						
	Controls $(n = 65)$	IgAD (<i>n</i> = 85)	CVID (<i>n</i> = 47)	P, IgAD controls	P, IgAD CVID	
Lymphocytes (%)	31.0 ± 7.8	29.0 ± 7.5	$24{\cdot}0\pm10{\cdot}7$	0.084	0.007*	
CD3 ⁺ (%)	74.0 ± 7.5	73.0 ± 7.1	79.0 ± 9.1	0.799	< 0.001*	
CD4 ⁺ (%)	45.9 ± 8.5	42.7 ± 8.2	37.8 ± 9.5	0.015	< 0.001*	
CD8 ⁺ (%)	21.8 ± 7.4	25.9 ± 5.8	36.0 ± 11.4	< 0.001*	< 0.001*	
CD19 ⁺ (%)	11.09 ± 3.1	12.0 ± 4.0	10.0 ± 7.3	0.094	0.099	
CD16/56 ⁺ CD3 ⁻ (%)	15.0 ± 7.8	13.0 ± 6.4	8.0 ± 7.1	0.422	0.004*	

 Table 1. Relative numbers of basic lymphocyte subpopulations in IgAD, CVID and controls. Results are expressed as median \pm standard deviation (s.d.). Lymphocytes are expressed as percentage of CD45⁺ cells, lymphocyte subpopulations as percentage of lymphocytes.

*Indicates statistically significant difference using critical value after Bonferroni's correction: 0-008. CVID: common variable immunodeficiency; IgAD: selective deficiency of immunoglobulin A.

CD4 cells and NK cells was also decreased significantly in CVID patients compared to IgAD patients.

Assessing the expression of activation/differentiation markers on CD4⁺ cells, a decreased expression of HLA-DR on CD4⁺ cells and an increased expression of CD25 on CD4⁺ cells was observed in IgAD patients compared to controls; this increase was also significant when comparing IgAD patients with CVID patients. No other significant differences were observed when comparing IgAD and controls. When comparing IgAD patients with CVID patients, an increase in the expression of CD45RA and a decrease in the expression of CD45RO and HLA-DR was observed in IgAD patients. Also, the number of CD28⁺ cells was increased and that of CD29⁺ cells decreased in IgAD compared to CVID patients (see Table 3).

When assessing activation/differentiation markers on CD8⁺ cells, only a decrease in CD29⁺ expression was observed comparing IgAD and healthy people or CVID patients, while almost in all parameters significant differences were observed between CVID and IgAD patients (decrease in CD45RO, CD57, CD38, HLA-DR and increase in CD25, CD27 and CD28 when comparing IgAD to CVID patients) (see Table 4).

There were no significant abnormalities in the absolute and relative numbers of B lymphocyte differentiation stages when comparing IgAD and controls, while in CVID patients typical abnormalities described previously by others [20–24]

Table 2. Absolute numbers of basic lymphocyte subpopulations in IgAD, CVID and controls. Results are expressed as median \pm standard deviation(s.d.).

	Controls $(n = 65)$	IgAD (<i>n</i> = 85)	CVID (<i>n</i> = 47)	P, IgAD controls	<i>P</i> , IgAD CVID
Leucocytes (10 ⁹ /l)	6.1 ± 1.5	6.1 ± 1.6	6.3 ± 2.3	0.962	0.483
Lymphocytes (10 ⁹ /l)	1.89 ± 0.12	1.77 ± 0.12	1.61 ± 0.25	0.170	0.186
CD3 ⁺ (10 ⁹ /l)	1.32 ± 0.49	1.25 ± 0.43	1.22 ± 0.77	0.231	0.483
CD4 ⁺ (10 ⁹ /l)	0.89 ± 0.36	0.75 ± 0.28	0.54 ± 0.32	0.006*	< 0.001*
CD8 ⁺ (10 ⁹ /l)	0.39 ± 0.32	0.43 ± 0.20	0.58 ± 0.50	0.143	0.023
CD19 ⁺ (10 ⁹ /l)	0.21 ± 0.09	0.20 ± 0.11	0.15 ± 0.18	0.905	0.023
CD16/56 ⁺ CD3 ⁻ (10 ⁹ /l)	0.24 ± 9.20	0.22 ± 0.13	0.14 ± 0.11	0.117	0.003*

*Indicates statistically significant difference using critical value after Bonferroni's correction: 0-007. CVID: common variable immunodeficiency; IgAD: selective deficiency of immunoglobulin A.

Table 3. Activation/differentiation markers on CD4⁺ cells on IgAD, CVID and controls. Results are expressed as median ± standard deviation (s.d.).

	Controls $(n = 65)$	IgAD (<i>n</i> = 85)	CVID $(n = 47)$	P, IgAD controls	P, IgAD CVID
CD4 ⁺ CD45RA ⁺ /CD4 ⁺ (%)	44·9 ± 16·2	45.0 ± 14.4	24.3 ± 16.3	0.775	< 0.001*
CD4 ⁺ CD45RO ⁺ /CD4 ⁺ (%)	49.7 ± 15.1	54.1 ± 15.1	75.4 ± 18.9	0.597	< 0.001*
CD4 ⁺ HLA-DR ⁺ /CD4 ⁺ (%)	6.9 ± 6.4	3.9 ± 3.6	10.2 ± 8.2	< 0.001*	< 0.001*
CD4 ⁺ CD25 ⁺ /CD4 ⁺ (%)	3.5 ± 3.1	9.7 ± 4.0	6.5 ± 7.9	< 0.001*	< 0.001*
CD62L ⁺ CD4 ⁺ /CD4 ⁺ (%)	75.3 ± 11.8	74.6 ± 11.2	71.9 ± 18.4	0.887	0.131
CD4 ⁺ CD27 ⁺ /CD4 ⁺ (%)	89.0 ± 8.5	86.3 ± 7.9	85·9 ± 12·8	0.039	0.314
CD4 ⁺ CD28 ⁺ /CD4 ⁺ (%)	98.3 ± 4.6	98.3 ± 3.6	95.6 ± 5.9	0.868	< 0.001*
CD4 ⁺ CD29 ⁺ /CD4 ⁺ (%)	50.4 ± 12.5	47.2 ± 13.4	65.3 ± 16.6	0.197	< 0.001*

*Indicates statistically significant difference using critical value after Bonferroni's correction: 0.006. CVID: common variable immunodeficiency; IgAD: selective deficiency of immunoglobulin A.

	Table 4.	Activation/differentiation ma	rkers on CD8 ⁺ cell	ls of IgAD, CVID	and control. Results are	expressed as median	\pm standard deviation	(s.d.).
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	Controls	IgAD	CVID	P, IgAD	P, IgAD
	(<i>n</i> = 65)	(n = 85)	(<i>n</i> = 47)	controls	CVID
CD8 ⁺ CD45RA ⁺ /CD8 ⁺ (%)	62·4 ± 13·5	58.5 ± 14.8	54·8 ± 16·2	0.197	0.131
CD8 ⁺ CD45RO ⁺ /CD8 ⁺ (%)	34.5 ± 13.6	28.9 ± 13.1	$41\cdot2 \pm 14\cdot2$	0.010	< 0.001*
CD8 ⁺ CD57/CD8 ⁺ (%)	17.3 ± 15.0	18.5 ± 12.9	34.7 ± 14.7	0.965	< 0.001*
CD8 ⁺ CD38/CD8 ⁺ (%)	13.3 ± 8.3	10.0 ± 7.1	23.8 ± 17.0	0.014	0.001*
CD8 ⁺ CD25 ⁺ /CD8 ⁺ (%)	1.1 ± 3.2	1.7 ± 5.6	0.6 ± 5.4	0.346	< 0.001*
CD8 ⁺ HLA-DR ⁺ /CD8 ⁺ (%)	6.2 ± 6.2	$2 \cdot 4 \pm 4 \cdot 6$	14.5 ± 15.8	0.280	0.005*
CD62L ⁺ CD8 ⁺ /CD8 ⁺ (%)	40.3 ± 17.5	39.2 ± 15.5	34.5 ± 19.1	0.603	0.151
CD8 ⁺ CD27 ⁺ /CD8 ⁺ (%)	69.2 ± 19.2	58.0 ± 15.6	42.5 ± 22.4	0.022	0.001*
CD8 ⁺ CD28 ⁺ /CD8 ⁺ (%)	65.7 ± 20.4	58·8 ± 17·9	35.6 ± 20.4	0.030	0.002*
CD8 ⁺ CD29 ⁺ /CD8 ⁺ (%)	93.4 ± 7.4	$79{\cdot}1 \pm 12{\cdot}8$	$92{\cdot}4\pm21{\cdot}7$	< 0.001*	< 0.001*

*Indicates statistically significant difference using critical value after Bonferroni's correction: 0-005. CVID: common variable immunodeficiency; IgAD: selective deficiency of immunoglobulin A.

were observed when compared to IgAD patients (Table 5). When assessing B lymphocyte numbers according to the Freiburg [22] classification of CVID, no major abnormalities were observed. Only one IgAD patients and one control person fell into group Ib.

Discussion

The goal of our study was to determine whether lymphocyte subpopulation abnormalities observed in CVID patients were also present in a genetically related disorder – IgAD. The results showed that the longest known abnormalities in CVID – the decrease in CD4⁺ cells and the increase in the relative number of CD8⁺ cells – can also be documented in IgAD patients although, in our results, these abnormalities were demonstrated only in absolute numbers of CD4⁺ cells and in relative numbers of CD8⁺ cells. The decrease in CD4⁺ cells and in relative numbers of CD8⁺ cells. The decrease in CD4⁺ cells was ascribed to the decrease in CD45RA⁺ cells [15], but we could not detect this abnormality in our IgAD patients; the percentage of CD45RA and also CD45RO was normal both in CD4⁺ and CD8⁺ cells.

A surprising observation was that of a decrease in the expression of HLA-DR on CD4+ cells of IgAD patients compared to both CVID patients and healthy controls, and an increase in the CD25 expression on CD4⁺ cells, again compared to both CVID and IgAD patients. Razziudin et al. [25] also described increased expression of Tac (CD25) antigen on peripheral blood lymphocytes in two of three IgAD patients; these individuals had an increased proportion of OKDR (HLA-DR) and OKIa1 (HLA-DR)-positive peripheral lymphocytes. HLA-DR and CD25 are both activation markers of T lymphocytes; CD25 is an earlier activation marker compared to HLA-DR [31]. We cannot explain the cause of possible activation of CD4⁺ cells in our IgAD patients because they were investigated in an infection-free period and none of them had clinically apparent autoimmune disease at the time of investigation. Surprisingly, no other activation abnormalities were observed in IgAD patients compared to controls. As well as on the activated lymphocytes, CD25 is also expressed on T regulated (T_{reg}) cells; we cannot exclude the possibility that this subpopulation abnormality influences the above-mentioned difference

Table 5. Absolute and relative numbers of differentiation stages of B cells. CVID patients with B lymphocytes < 1% were excluded. Naive B lymphocytes were defined as: $CD19^+ CD27^- IgM^+ IgD^+$ cells, IgM memory as: $CD19^+ CD27^+ IgM^+ IgD^+$ cells, switched memory as: $CD19^+ CD27^+ IgM^- IgD^-$ cells, immature as $CD19^+ IgM^+ CD21^-$.

	Controls	IgAD	CVID	P, IgAD	P, IgAD
	(n = 65)	(n = 85)	(n = 40)	controls	CVID
Naive (10 ⁹ /l)	0.099 ± 0.053	0.111 ± 0.072	0.142 ± 0.139	0.308	0.271
IgM memory (10 ⁹ /l)	0.046 ± 0.025	0.043 ± 0.031	0.015 ± 0.043	0.902	< 0.001*
Switched memory (109/l)	0.034 ± 0.038	0.028 ± 0.024	0.004 ± 0.012	0.108	< 0.001*
IgM ⁺ CD21 ⁻ (10 ⁹ /l)	0.009 ± 0.011	0.012 ± 0.015	0.029 ± 0.064	0.034	< 0.001*
Naive/B cells (%)	52.8 ± 12.5	56.8 ± 14.5	83.3 ± 15.8	0.135	< 0.001*
Naive/PBL (%)	1.8 ± 1.1	1.6 ± 1.1	0.2 ± 0.6	0.203	< 0.001*
IgM memory/B cells (%)	22.6 ± 7.4	20.5 ± 10.4	8.0 ± 12.3	0.520	< 0.001*
Switched memory/B cells (%)	19.6 ± 7.5	14.9 ± 7.5	1.8 ± 3.8	0.039	< 0.001*
IgM ⁺ CD21 ⁻ /B cells (%)	4.8 ± 4.3	6.7 ± 4.2	14.8 ± 14.8	0.025	< 0.001*

*Indicates statistically significant difference using critical value after Bonferroni's correction: 0.006. CVID: common variable immunodeficiency; IgAD: selective deficiency of immunoglobulin A.

Unlike in CVID, we could not document any statistically significant abnormalities in the differentiation stages of B cells, only a tendency to a decreased number of switched memory cells and a mild increase in IgM⁺ CD21⁻ immature cells was present, but these abnormalities were much less extensive than those observed in CVID patients. In IgAD patients, a reduced but still significant number of IgA+ B cells can be documented [28]. On a molecular level, an Ia germline transcript expression was reduced and α circle transcripts were not detected in three IgAD patients [32]. Another study showed normal Ia germline transcription in four IgAD patients, but in two of them the Sµ–S α switch recombination was impaired while in the remaining two decreased secreted and membrane Ca mRNA was markedly decreased in IgA switched B cells [33]. All these results show a probable heterogeneity in the B cell defect of IgAD patients, but according to our results, unlike the situation in CVID, such abnormalities do not affect maturation of the majority of B cells, as reflected by lymphocyte activation marker determination in blood.

In conclusion, our study showed that, with the exception of abnormalities in the numbers of CD4⁺ and CD8⁺ cells, no major abnormalities of lymphocyte subpopulations or expression of activation/differentiation markers were observed in IgAD patients. These results may show that lymphocyte subpopulation abnormalities seen in CVID are probably not a consequence of an inborn genetic defect but that they evolve as a part, or as a consequence, of immunodeficiency in CVID.

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