The molecular basis for the presence of two autoimmune diseases occurring simultaneously – Preliminary observations based on computer analysis

LABIB R. ZAKKA¹, PEDRO A. RECHE², & A. RAZZAQUE AHMED¹

¹Center for Blistering Diseases, Boston, Massachusetts, USA, and ²Immunomedicine Group, Department of Immunology, Facultad de Medicina, Universidad Complutense de Madrid, Madrid, Spain

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Abstract

Specific Human Leukocyte Antigen Class II (HLA II) molecules associated with pemphigus vulgaris (PV), mucous membraine pemphigoid (MMP), and mixed connective tissue disease (MCTD) may react with multiple T cell epitopes within desmoglein 3 (Dsg 3), bullous pemphigoid antigen 2 (BPAG 2), and 70 kDa polypeptide small nuclear ribonucleoproteins (snRNP70) in autoantibody production. We report a group of patients with simultaneous occurrences of PV with MCTD, and MMP with MCTD. In one patient group, we performed serological studies to show presence of antibodies to Dsg 3, Dsg 1, and snRNP70 simultaneously. In the second group, we performed serological studies to show presence of antibodies to BPAG 1, BPAG 2, β 4 integrin, and snRNP70 simultaneously. In both groups, HLA II genes were analyzed and the observations were consistent with previously described associations with PV, MMP, and MCTD. It is possible that HLA-DQ β 1*0301 allele, present in 10 of 17 patients and DR β 1*04 in some of the others, may have the ability to bind to several relevant T cell epitopes in the snRNP70 molecule. We have utilized a computer model to demonstrate that HLA II-restricted T cell epitopes present within the known autoantigens may be capable of eliciting an immune response. While other explanations and mechanisms exist, the authors suggest that epitope spreading may be one possible mechanism, amongst others, that may result in the simultaneous presence of two separate pathogenic autoantibodies.

Keywords: MHC Class II Genes, Epitope Spreading, Pemphigus Vulgaris, Mucous Membrane Pemphigoid, Mixed Connective Tissue Disease

Introduction

Pemphigus Vulgaris (PV) is a potentially fatal autoimmune mucocutaneous blistering disease (AMBD). Patients with PV present with flaccid blisters, affecting the skin as well as the mucosal surfaces, that rupture easily leaving denuded epithelium [1-4]. Histology shows an intraepidermal vesicle, and keratinocyte cell surface antibody deposition is seen on immunopathology [1-7]. The target antigens in PV are Desmoglein 1 (Dsg 1), Desmoglein 3 (Dsg 3) [8], and possibly the acetylcholine receptor [9]. Disease severity and activity may correlate with serum autoantibody titers [8,10–12].

Mucous Membrane Pemphigoid (MMP), Also known as Cicatricial Pemphigoid (CP), is an AMBD that affects the elderly. Patients present with a mean age of onset of 73 years [13]. MMP affects the mucous membranes of the conjunctiva, nose, oral mucosa, oesophagus, larynx, pharynx, genitalia, anal canal, and the skin [2-4,6,7,14,15]. Upon healing, the lesions produce irreversible scaring [2-4,6,7,14,15]. The antigens targeted by autoantibodies in MMP are a 230 kDa protein called Desmoplakin, also referred to as Bullous Pemphigoid Antigen 1 (BPAG 1), a 180 kDa hemidesmosome protein, also referred to as Bullous Pemphigoid Antigen 2 (BPAG 2), and human β 4 integrin subunit [16,17].

Mixed Connective Tissue Disease (MCTD) is a systemic autoimmune disease with overlapping features of systemic lupus erythematosus (SLE),

Correspondence: A. Razzaque Ahmed, Center for Blistering Diseases, 70 Parker Hill Avenue, Boston, MA, USA 02120, E-mail: arahmedmd@msn.com

dermatomyositis, rheumatoid arthritis, scleroderma, and polymyositis [18–21]. Primary symptoms of MCTD include arthralgias, Reynaud's Phenomenon, esophageal reflux or dysmotility, and finger swelling or diffuse hand swelling among others [19–32] (Table I). The primary cause of death in patients with MCTD is pulmonary disease [19,22,23]. The target antigen is a complex of small nuclear ribonucleoproteins (snRNP), including a 70 kDa polypeptide (snRNP70), that are non-covalently linked with the U1 ribonucleoprotein (U1-RNP) of the spliceosome complex [19,33,34]. It has been reported that the snRNP70 contains the major antigenic epitopes to which MCTD sera react [35–37].

Antibodies directed against protein antigens are produced by B cells engaging CD4 helper T cells. The mechanism occurs as follows: Protein antigens recognized by B cells through their B cell receptor (BCR) are internalized and processed [38]. Peptide antigens resulting from such processing are loaded onto Major Histocompatibility Complex Class II (MHC II) molecules, Human Leukocyte Antigen Class II (HLA II) molecules in humans, and expressed on the cell surface of B cells [38].

CD4 T cells bearing the appropriate T cell receptors (TCR) can then recognize these cell surface expressed HLA II-peptide antigens [38]. Only those B cells displaying CD4 T cell epitopes (HLA II-peptide antigens that are recognized by CD4 T cells) are subsequently capable of producing antibodies against the original antigen [38]. By this same mechanism, CD4 helper T cells lead to the production of pathogenic autoantibodies after recognizing T cell epitopes from self-antigens [39].

Earlier clinical studies have demonstrated that certain patients who have clinical, histological, and immunopathological features of PV have MCTD present simultaneously [40]. Likewise, certain patients clinically presenting as MMP have MCTD simultaneously [18]. The treatment and the clinical

Table I. Major clinical features of MCTD*.

	% involvement
Arthritis/arthralgia	95
Raynauds	85
Oesophageal involvement	67
Impaired lung diffusion	67
Swollen hands	66
Myositis	63
Scleroderma	33
Serositis	27
Renal disease	10
Cerebral involvement	10

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outcomes of these patients have been described in those publications [18,40]. The purpose of this study was to identify potential autoreactive HLA II-restricted helper T cell epitopes shared among PV, MMP, and MCTD that could explain the simultaneous occurrence of these relatively rare three diseases.

Methods

Patients

This study reports on two groups of patients. These patients presented to the Center for Blistering Diseases (CBD) in Boston between March 2001 and November 2009. An Institutional Review Board (IRB) approval was obtained as well as a written consent from each patient. It is important to note that the patients were diagnosed prior to their evaluation at the CBD and were referred to the CBD for the management of their complex disease. During the diagnostic work-up, a thorough serological evaluation was done. During this evaluation, it was observed that the patients had antibodies to snRNP70. In addition, in a review of systems, patients were found to have symptoms of MCTD. Since two relatively rare diseases were present in a group of patients, HLA typing for HLA Class II genes was performed.

Inclusion criteria

- (1) Upon their initial diagnosis, the patients had histology, confirmed by direct immunofluores-cence, for MMP and PV.
- (2) Careful history of medications was taken and patients were not included if they were on medications known to be associated with inducing either MCTD, MMP or PV [41–45].
- (3) All these patients with PV had antibodies to keratinocyte cell surface as determined by indirect immunofluorescence (IIF) using monkey esophagus as substrate. In addition, on Enzyme-Linked Immunosorbent Assay (ELISA), patients demonstrated high levels of antibodies to Dsg 1 and Dsg 3.
- (4) Patients with MMP had antibodies to human β4 integrin subunit as determined by an immunoblot assay using Bovine Gingival Lysate as substrate. The presence of a 205 kDa band was considered the binding of the antibody in the patient sera to human β4 integrin subunit. The positive control was UMA6 antibody. The sera was negative for presence of antibody to human α6 integrin subunit. In addition, IIF was done to determine binding of patient sera to BPAG1 and BPAG2 were determined by a standard ELISA.
- (5) The patients met the criteria set by Alarcon-Segovia [46,47] to be considered as having MCTD.

Autoimmunity Downloaded from informahealthcare.com by 108.7.175.111 on 05/29/12 For personal use only. (6) The presence of antibodies to snRNP70 in the sera of the patients tested on at least three different days at monthly intervals.

Eleven patients with extensive mucocutaneous PV were studied. Each patient had minimal 25–30% or more of body surface involvement with at least 2 mucosae involved. These patients had not responded to a mean dose of 125 mg/day of Prednisone and were non-responsive to at least 2 more immunosuppressive therapies. One feature that distinguishes these patients from other PV patients is that their disease was considered severe, widespread and recalcitrant to conventional immunosuppressive therapy.

Six patients with MMP are presented. These 6 patients all had extensive erosions in the oral cavity, pharynx, and nasal disease. Three patients had Ocular Cicatricial Pemphigoid (OCP) and 2 had vaginal disease. Each of the 6 patients had been treated intermittently with systemic corticosteroids and at least 3 or more immunosuppressive agents. All had received Dapsone therapy. Like the PV subsets, these MMP patients had progressive worsening disease with minimal benefit from conventional immunosuppressive therapy.

Serological analysis

In both groups of patients, antibodies to Dsg 1, Dsg 3, BPAG 1, and BPAG 2 were measured using an ELISA [12,48–50]. The index values of the ELISAs for BPAG 1 and BPAG 2 are identical: fewer than 9 is negative and greater than 9 is positive. The Dsg 3 ELISA index values are: Fewer than nine is negative, 9–20 is intermediate, and greater than 20 is positive. The Dsg 1 ELISA index values are: Less than 14 is negative, 9–20 is intermediate, and more than 20 is positive. Antibodies to human β 4 integrin subunit and α 6 integrin subunit were done by an immunoblot using bovine gingival lysate as substrate [51].

Patient sera in both groups was evaluated by IIF using monkey esophagus as substrates which measured the titers of the antibodies to intercellular cement substance (ICS) and BMZ proteins. The serological tests for antibodies to Dsg 3, Dsg 1, BPAG 1, BPAG 2, and snRNP70 were performed by laboratories at hospitals from where the patients were referred, and some by Beutner Laboratories, Buffalo, NY.

HLA Class II genes

Peripheral blood leukocytes obtained from 11 patients with PV and MCTD, and six patients with MMP and MCTD were drawn during the active stage of their diseases and studied for HLA II genes. We used the Polymerase Chain Reaction with Sequence Specific Primers (PCR-SSP) to identify HLA II genes encoded by the DR β 1 and DQ β 1 loci as described [52]. The PCR-SSP was performed by the referring institutions, and some by the American Red Cross, HLA Lab in Dedham, MA.

Prediction of HLA II-restricted CD4 T cell epitopes. CD4 helper T cell cooperation can only be delivered upon the recognition of peptides bound to HLA II molecules (CD4 T cell epitopes). Therefore, we used peptide binding predictions to HLA II molecules as the basis for anticipating CD4 T cell epitopes [53]. Thus, we anticipated HLA II-restricted CD4 T cell epitopes in Dsg 3, BPAG 2, and snRNP70 using RANKPEP (http://imed.med.ucm.es/Tools/rankpep. html) [54,55]. RANKPEP implements motif profiles (position specific scoring matrix) derived from peptides known to bind to MHC as the basis to predict peptide-MHC binding [54–56] and, subsequently, T cell epitopes.

Specifically, we selected the relevant motif profiles to predict HLA-DQ7 (HLA-DRQ1*0301) and HLA-DR4 (HLA-DRB1*0402) restricted CD4 T cell epitopes in Dsg 3, BPAG 2, and snRNP70. Only those peptides that had a binding score above the binding threshold (BT) were considered as CD4 T cell epitopes. The BT is a valuable index in determining the potential capability of the T cell epitope in triggering T cell response in-vivo (details elsewhere in Reche et al. [54–56]).

Results

High resolution HLA II gene analysis By PCR-SSP

Only patients were studied without their families. However, based on linkage disequilibrium studies from HLA international workshops, the data and their known associations have been written as haplotypic rather than phenotypic.

Group 1: Patients simultaneously having pemphigus vulgaris and mixed connective tissue disease. Data on HLA typing is presented in Table I. Patients in this group had a diagnosis of PV and MCTD. This group consisted of 11 patients.

Serology results. All 11 patients had high antibody titers to Dsg 1 and Dsg 3, as measured by ELISA, that were high. The IIF titers of anti-ICS antibodies were also very high. All 11 patients had presence of antibodies to snRNP70. This data is presented in Table II.

High resolution HLA II gene analysis by PCR-SSP demonstrated that 6 patients had HLA-DR β 1*0402 and 8 patients carried DQ β 1*0503. Five patients did not carry the HLA-DR4 molecule. Also, 6 patients carried HLA-DQ β 1*0301 and of those patients, 4 did

					Group 1							
		HL	A II									
	1st Haplotype		2nd Haplotype		Serology							
Patient	DRβ1	DQβ1	DR _{β1}	DQβ1	Anti-Dsg 1	Anti-Dsg 3	IIF anti-ICS antibodies	Anti-snRNP70				
1	402	302	1401	503	110	177	640	+				
2	402	302	1401	503	168	335	320	+				
3	402	302	1401	503	110	201	160	+				
4	402	302	1101	301	88	143	160	+				
5	402	302	1104	301	41	171	320	+				
6	402	302	1301	603	77	127	160	+				
7	1101	301	1401	503	81	156	320	+				
8	701	301	1401	503	79	148	320	+				
9	1103	301	1401	503	82	63	640	+				
10	1101	301	1401	503	107	189	80	+				
11	1501	609	1401	503	121	233	160	+				

Table II. Human Leukocyte Antigen Class II gene analysis of patients with *Pemphigus Vulgaris* (PV) and Mixed Connective Tissue Disease (MCTD).

(HLA II = Human Leukocyte Antigen Class II; Dsg = Desmoglein; IIF = Indirect immunofluorescence; ICS = Intercellular Cement Substance; RNP = Ribonucleoprotein; Dsg 1 ELISA Index Values: <14 = Negative; 14-20 = Indeterminate; >20 = Positive [12,45]; Dsg 3 ELISA Index Values: <9 = Negative; 9-20 = Indeterminate; >20 = Positive [12,45]).

not carry the HLA-DR4 molecule. Also, four patients carried HLA-DQ β 1*0302 and 1 patient carried the HLA-DQ β 1*0603. One patient did not carry neither HLA-DR4 nor HLA-DQ β 1*0301.

Group 2: Patients simultaneously having mucous membrane pemphigoid and mixed connective tissue disease. Data on HLA typing is presented in Table II. Patients in this group had a diagnosis of MMP and MCTD. This group consisted of 6 patients.

Serology results. All 6 patients had antibody titers to BPAG 1 and BPAG 2, as measured by ELISA, that were high. The IIF titers for anti-BMZ antibodies were also very high. Antibodies to the subunits of human $\alpha 6$ were negative, and all 6 patients had antibodies to the subunit of human $\beta 4$ integrin. All 6 patients had presence of antibodies to snRNP70. This data is presented in Table III.

High resolution HLA II gene analysis by PCR-SSP demonstrated that 3 patients carried the HLA-DR4 molecule. Four patients carried HLA-DQ β 1*0301, and of those patients 2 were homozygous for HLA-DQ β 1*0301, and 2 did not carry the HLA-DR4 molecule. One patient carried HLA-DQ β 1*0302 allele with no HLA-DR4 molecule, and one patient carried both HLA-DQ β 1*0302 and HLA-DQ β 1*0603.

CD4 T cell epitopes associated with HLA Class II DRβ1*0402 and DQβ1*0301-genes

The HLA II genes associated with PV, MMP, and MCTD and the relevant antigens, when subjected to

the RANKPEP program, demonstrate that there are potential T cell epitopes within snRNP70 (Figure 1), Dsg 3 (Figure 2), and BPAG 2 (Figure 3) that are predicted to bind to DR β 1*0402. Also, the RANK-PEP program demonstrates potential T cell epitopes within snRNP70 (Figure 1), Dsg 3 (Figure 2), and BPAG 2 (Figure 3) that are predicted to bind to DQ β 1*0301.

One study reported increased binding of T cells within amino acid sequence 374-437 of the snRNP70 in patients with MCTD [37]. However, the study did not report the HLA II allele of the patients studied. The RANKPEP program shows binding of snRNP70 to HLA-DQ7 on amino acid 373-384 (Figure 1). Also, the RANKPEP program showed binding of BP180 to HLA-DQ7 on amino acid 505-513, 635-643, 765-773, 841-849, 1055-1063, 1192-1210,1283-1291, and 1320-1328 among others (Figure 3), in accordance with the literature [57-60]. Moreover, the RANKPEP program shows binding of Dsg 3 to HLA (DR4)DRβ1*0402 on amino acid 193-200, 217-225, 765-774, 605-613 in accordance with the literature [61-63]. Therefore, it should be noted that the RANKPEP program is reliable in predicting antigen-MCH II binding.

Discussion and conclusion

In this report we are presenting 11 patients who simultaneously had the clinical and serological diagnosis of PV and MCTD. We also report 6 patients who simultaneously had the clinical and serological features of MMP and MCTD. Clinical data on some Human Leukocyte Antigen Class II gene analysis of patients with mucous membrane pemphigoid (MMP) and Mixed Connective Tissue Disease (MCTD). ((HLA II = Human Leukocyte = Ribonucleoprotein; BPAG1 and BPAG2 Index = Basement Membrane Zone; RNP Antigen Class II; BPAG = Bullous Pemphigoid Antigen; IIF = Indirect immunofluorescence; BMZ Values: <9 = Negative; >9 = Positive [43, 44]) Table III.

						Group 2				
		HL	HLA II							
	1st Ha	lst Haplotype	2nd Ha	2nd Haplotype			Serology	Sy		
Patient	DRβ1	DQB1	DRβ1	DQβ1	Anti-BPAG 1	Anti-BPAG 2	IIF anti-BMZ antibodies	Anti-β4 antibodies	Anti-α6 antibodies	Anti-snRNP70
1	701	501	101	302	27	149	160	+	I	+
2	408	301	408	301	47	130	320	+	Ι	+
3	402	302	1104	603	88	320	640	+	Ι	+
4	301	201	1104	301	121	132	80	+	Ι	+
2	407	302	4011	301	76	87	160	+	Ι	+
6	1101	301	1101	301	74	82	320	+	I	+

of these patients have been reported previously [18,40].

All the patients in Group 1 had either the HLA II alleles DRB1*0402 or DQB1*0503 that have been reported in patients with PV in several studies [64-80]. However, only six patients had HLA-DR4 molecule, while the remaining 5 patients did not carry any allele known to be associated with MCTD [23,32,81–88]. Moreover, ten patients carried either DQ\\beta1*0301, DQ\\beta1*0302, or DQ\\beta1*0603 alleles, four of whom did not carry HLA-DR4 molecule. Nonetheless, these 4 patients had the clinical features and serological markers typically associated with MCTD. It has been reported by Delgado et al. that DQβ1*0302, *0303, *0305, *0602 and *0603 allele share amino acid sequences 71-77 (Thr-Arg-Ala-Glu-Leu-Val-Thr) with *0301 [89]. Therefore, it is possible that these alleles have peptide binding specificities similar to $DQ\beta1*0301$.

All of the patients in group 2 carried either the HLA II alleles $DQ\beta1*0301$, *0302, or *0603 that have been reported in patients with MMP [17,89–92]. However, only three of six patients also carried the HLA-DR4 molecule, while the remaining 3 patients did not carry any allele known to be associated with MCTD [23,32,81–88]. Nonetheless, these 3 patients had clinical features and serological markers typically associated with MCTD.

The authors suggest that an immunogenetic basis may exist to explain the unique observations presented, while highlighting that there could be several other explanations. Although no other studies exist showing the presence of MMP and MCTD or PV and MCTD simultaneously in a patient, MMP and PV have been reported with many other autoimmune diseases. MMP has been reported in patients with SLE and Rheumatoid Arthritis [93-96]. Also, PV has been reported with SLE, Sjogren's Syndrome, and Systemic Sclerosis [97-105]. Moreover, there are multiple reports in the literature describing patients with PV and MMP simultaneously [8]. One study reported that DQβ1*0301 homozygous typing T cell lines from patients with PV proliferated in reaction to Dsg 3 peptide [106].

In the opinion of the authors, based on the computer modeling and several reports in the literature documenting the presence of these 2 diseases simultaneously in one patient, the phenomenon of epitope spreading may provide a possible explanation for the simultaneous presence of 2 autoimmune diseases in these 2 groups of patients presented in this report. Four possible scenarios of epitope spreading exist. The first scenario may be that one large antigen molecule may contain at least 2 similar epitopes with possible sequence overlap, and when 2 different HLA II molecules present them, they stimulate two different T cells [107].

Homo sapiens small nuclear ribonucleoprotein 70kDa polypeptide (RNP antigen) (SNRP70) Protein sequence:	RNP binding to DQβ*0301 Matrix: HLA (DQ7)DQβ1*0301, Consensus sequence: IWHAVHAWH All rows highlighted in red represent predicted binders.										
Epitopes predicted to bind to DQβ1*0301 are represented as yellow. Epitopes predicted to bind DRβ1*0402 are	RANK	POS.	N	SEQUENCE	С	MW (Da)	SCORE	% OPT.			
highlighted as green.	1	137	MVY	SKRSGKPRG	YAF	954.1	15.071	33.00 %			
Predicted epitope that overlaps and likely to be presented by both MHC molecules is represented as red.	2	187	VKG	WRPRRLGGG	LGG	1013.21	12.75	27.92 %			
	3	373	RDR	DRDREHKRG	ERG	1150.23	11.87	25.99 %			
39 CGIAPYIREFEDPRDAPPPTRAETREERMERKRREKI	RNP bi	aing	to DR	0402							
76 ERRQQEVETE KMMTPHNIPNAQGDAFKTLFVAR 110 VNYDTTESKLRREFEVYGPIKRIHMVY <mark>BKRSGKPR</mark> 145 <mark>GYAFIEYEHFEDW HSAYKHADGKKIJGRRVIVDVE</mark> 180 RGRTVMCWRPREGGGLGGTRRGGADVNIRHSGR 214 DDTSRYDERPGPSPLPHRDRDRDRERERERSRER 248 DKERERRSRSRDRRRSRSRDKEERRRSRERSKD 263 KDRDRKRRSSRERARREERKEELRGGGGDMA	TUP	380 24-98 380 26-98		, Consensus seque n red represent pre							
10 VNYDTTESKLRREFEVYGPIKRIHMYYSKRSGKPR 45 GYAFTEVEHERDTHSAYKHADGKKIDGRRVLVDVE 80 RGRTKKGNERERINGGGGLGCTREGGGDVNIKHSGR 214 DDTSRYDERPGPSPLPHRDRDRDRERERRERSRER 248 DKERERRSRSRORRRSRSRDKEERRRSRESKD 248 KERDRKRSSRSRERARRERERKELELRGGGDMA 218 EPEEAGDAPPDDGPGELGEDGPDGPEEKGRDRDR	TUP	highlig	phted in	, Consensus sequ				% OPT			
110 VNYDTTESKLRREFEVYGPIKRIHMYY <mark>SKRSGKPR</mark> 145 GYAF EYEHERDMISAYKHADGKKIDGRRULVDVE 180 RGRTVKUWRPRREGGGLGGTRRGGADVNIRHSGR 14 DDTSRYDERFCPSPLPHRDRDRDRERERRERSRER 248 DKERERRSRSRDRRRSRSRDKEERRRSRERSKD 158 EFSEAGDAPPDDGPPGELGPDGPDGPEEKGRDRDR 153 ERRSHRSERERRDRDRDRDRDRDREKKGERGSER 168 GRDEARGGGGGQDNGLEGLGNDSRDMYMESEGG	All rows	highlig	phted in	, Consensus seque	dicted	binders.		% OPT 55.68 %			
110 VNYDTTESKLRREFEVYGPIKRIHMYY <mark>SKRSGKPR</mark> 145 GYAFIEVEHERDENSAYKHADGKKIDGRRVLVDVE 180 RGRTIKGWRPREGGGLGGTRRGGADVNIRHSGR 14 DDTSRYDERFGPSPLPHRDRDRDRERERRERSRER 248 DKERERRSRSRDRRRSRSRDKEERRSRERSKD 181 EPSERGDAPPDDGPPGLGEDGPGDEEKGRDRDR 181 EPSERGDAPPDDGPPGLGEDGPGDEEKRBRDR 181 EPSERGDAPPDDGPPGLGEDGPGDEEKRBRDR 181 EPSERGDAPPDDGPPGLGEDGPGDEEKRBRDR	All rows	highlig POS.	phted in	, Consensus sequenting of the sequence of the	dicted C	binders. MW (Da)	SCORE	The second second second			

Figure 1. Predicted CD4 T cell epitopes in small nuclear ribonucleoproteins 70kDa polypeptide (snRNP70). Potential snRNP70-specific CD4 T cell epitopes restricted by HLA-DQ7(DQB1*0301) and HLA-DR β 1*0402 are shown in yellow and green, respectively. T cell epitopes that are predicted to bind both HLA-DR β 1*0402 and HLA-DQ7(DQB1*0301), are show in red. All shown peptides exceed the Binding Threshold (see Methods for details). POS. = Position; N = N-terminal; C = C-terminal; MW = Molecular Weight (Daltons); %OPT. = Optimal Score.

Homo sapiens desmoglein 3 (pemphigus vulgaris antigen)	Matrix:	HLA (I	DQ7)DO	oinding to D Ωβ1*0301, Cons	ensus	sequence:	IWHAVH4	WH,	
Epitopes predicted to bind to DQβ1*0301 are represented as yellow. Epitopes predicted to bind DRβ1*0402 are highlighted as green. Predicted epitope that overlaps and likely to be presented by both	Optimal Score: 45.671, Binding Threshold: 11.70 All rows highlighted in red represent predicted binders.								
MHC molecules is represented as red.	RANK	POS.	N	SEQUENCE	C	MW (Da)	SCORE	% OPT.	
	1	727	LGA	ATESGGAAG	FAT	701.69	17.045	37.32 %	
MMGLFPRTTGALAIFVVVILVHGELRIETKGQYDEEEMTMQQAKR	2	911	ALS	ASGSVQPAV	SIP	796.88	15.559	34.07 %	
ROKREWVKFAKPCREGEDNSKRNPIAKITSDYOATOKITYRISGVGI	3	400	SKK	LVDYILGTY	QAI	1038.21	15.092	33.05 %	
DOPPFGIFVVDKNTGDINITAIVDREETPSFLITCRALNAOGLDVEKP	4	73	IAK	ITSDYQATQ	KIT	1008.05	14.73	32.25 %	
LILTVKILDINDNPPVFSQQIFMGEIEENSASNSLVMILNATDADEPN	5	232	ASS	YRLVVSGAD	KDG	961.09	13.563	29.70 %	
ILNSK <mark>IAFKIVSQ</mark> EPAGTPMFLLSRNTGE <mark>VRTLTNSLD</mark> REQASS <mark>YRL</mark>		27	2 (Sala	197 G 19 192	1.01010				
VVSGAD KDGEGLSTQCECNIKVKDVNDNFPMFRDSQYSARIEENIL	Desm	ogle	in 3 I	binding to D	Rβ1*	0402			
SSELLRFQVTDLDEEYTDNWLAVYFFTSGNEGNWFEIQTDPRTNEG		10.1			- C.				
ILKVVKALDYEQLQSVKLSIAVKNKAEFHQSVISRYRVQSTPVTIQV				Rβ1*0402, Conse			ICFWHNH	NM,	
INVREGIAFRPASKTFTVQKGISSKKLVDYILGTYQAIDEDTNKAAS				04, Binding Thre					
NV <mark>KYVMGRNDG</mark> GYLMIDSKTAE <mark>IKFVKNMNR</mark> DSTFIVNKTITAEV	All row	s highl	ighted		4 mma di				
LAIDEYTGKTSTGTVYVRVPDFNDNCPTAVLEKDAVCSSSPSVVVS				in red represen	t predi	cted binde	rs.		
		1.2.1	-	8					
ARTLNNRYTGPYTFALEDQPVKLPAVWSITTLNATSALLRAQEQIP	RANK		N	SEQUENCE	c	MW (Da)	SCORE		
	1	605	N SPG	SEQUENCE TRYGRPHSG	C RLG	MW (Da) 1012.1	SCORE 17.99	40.33 %	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG		605 765	N SPG SGT	SEQUENCE TRYGRPHSG MRTRHSTGG	C RLG TNK	MW (Da) 1012.1 984.09	SCORE 17.99 17.387	40.33 9	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSCRLGPAAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV	1	605	N SPG	SEQUENCE TRYGRPHSG	C RLG	MW (Da) 1012.1	SCORE 17.99	40.33 % 38.98 %	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSCRLGPAAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME	1 2	605 765	N SPG SGT	SEQUENCE TRYGRPHSG MRTRHSTGG	C RLG TNK	MW (Da) 1012.1 984.09	SCORE 17.99 17.387	40.33 9 38.98 9 33.96 9	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSCRLGPAAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SEVCTNTYARGTAVEGTSGMEMTTKLGA <mark>ATESGGAAG</mark> FATGTVS	1 2 3	605 765 193	N SPG SGT NSK	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE	C RLG TNK PAG	MW (Da) 1012.1 984.09 1016.21	SCORE 17.99 17.387 15.149	40.33 % 38.98 % 33.96 % 33.73 %	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSGRLGPAAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SEVCTNTYARGTAVEGTSGMEMTTKLGA <mark>ATESGGAAG</mark> FATGTVS AASGFGAATGVGICSSGQSGT <mark>MRTRHSTGG</mark> TNKDYADGAISMNF	1 2 3 4	605 765 193 443	N SPG SGT NSK TAE	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE IKFVKNMNR	C RLG TNK PAG DST	MW (Da) 1012.1 984.09 1016.21 1131.39	SCORE 17.99 17.387 15.149 15.046	40.33 % 38.98 % 33.96 % 33.73 % 32.63 %	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSGRLGPAAIGLLLLGLLLLAPLLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SSEVCTNTYARGTAVEGTSGMEMTTKLGAATESGGAAGFATGTVS SAASGFGAATGVGICSSGQSGTMTKTHISTGGTNKDYADGAISMNF DSYFSQKAFACAEEDDGQEANDCLLIYDNEGADATGSPVGSVGC	1 2 3 4 5	605 765 193 443 422	N SPG SGT NSK TAE ASN	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE IKFVKNMNR VKYVMGRND	C RLG TNK PAG DST GGY	MW (Da) 1012.1 984.09 1016.21 1131.39 1063.23	SCORE 17.99 17.387 15.149 15.046 14.554	40.33 % 38.98 % 33.96 % 33.73 % 32.63 % 30.82 %	
PGVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSGRLGPÄAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV IGGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SSEVCTNTYARGTAVEGTSGMEMTTKLGAATESGGAAGFATGTVS GAASGFGAATGVGICSSGQSGTMRTRHSTGGTNKDYADGAISMNF DSYFSQKAFACAEEDDGQEANDCLLIYDNEGADATGSPVGSVGC CSFIADDLDDSFLDSLGPKFKKLAEISLGVDGEGKEVQPPSKDSGYG	1 2 3 4 5 6	605 765 193 443 422 83	N SPG SGT NSK TAE ASN TQK	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE IKFVKNMNR VKYVMGRND ITYRISGVG	C RLG TNK PAG DST GGY IDQ	MW (Da) 1012.1 984.09 1016.21 1131.39 1063.23 947.1	SCORE 17.99 17.387 15.149 15.046 14.554 13.746	% OPT 40.33 % 38.98 % 33.96 % 33.73 % 32.63 % 30.82 % 28.19 % 27.89 %	
GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG RYGRPHSGRLGPAAIGLLLLGLLLLLAPLLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SEVCTNTYARGTAVEGTSGMEMTTKLGAATESGGAAGFATGTVS GAASGFGAATGVGICSSGQSGTMRTRHSTGGTNKDYADGAISMNF DSYFSQKAFACAEEDDGQEANDCLLIYDNEGADATGSPVGSVGC SFIADDLDDSFLDSLGPKFKKLAEISLGVDGEGKEVQPPSKDSGYG ESCGHPIEVQQTGFVKCQTLSGSQGASALS <mark>ASGSVQPAV</mark> SIPDPLQ	1 2 3 4 5 6 7	605 765 193 443 422 83 812	N SPG SGT NSK TAE ASN TQK NDC	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE IKFVKNMNR VKYVMGRND ITYRISGVG LLIYDNEGA	C RLG TNK PAG DST GGY IDQ DAT	MW (Da) 1012.1 984.09 1016.21 1131.39 1063.23 947.1 989.1	SCORE 17.99 17.387 15.149 15.046 14.554 13.746 12.572	40.33 % 38.98 % 33.96 % 33.73 % 32.63 % 30.82 % 28.19 %	
ARTLNNRYTGPYTFALEDQPVKLPAVWSITTLNATSALLRAQEQIP GVYHISLVLTDSQNNRCEMPRSLTLEVCQCDNRGICGTSYPTTSPG IRYGRPISG RLGPAAIGLLLLGLLLLLAPLLLTCDCGAGSTGGV GGFIPVPDGSEGTIHQWGIEGAHPEDKEITNICVPPVTANGADFME SSEVCTNTYARGTAVEGTSGMEMTTKLGAATESGGAAGFATGTVS GAASGFGAATGVGICSSGQSGTMRTRHSTGGTNKDYADGAISMNF .DSYFSQKAFACAEEDDGQEANDCLLIYDNEGADATGSPVGSVGC CSFIADDLDDSFLDSLGPKFKKLAEISLGVDGEGKEVQPPSKDSGYG ESCGHPIEVQQTGFVKCQTLSGSQGASALSASGSVQPAVSIPDPLQ 4GNYLVTETYSASGSLVQPSTAGFDPLLTQNVIVTERVICPISSVPG VLAGPTQLRGSHTMLCTEDPCSRLI	1 2 3 4 5 6 7 8	605 765 193 443 422 83 812 70	N SPG SGT NSK TAE ASN TQK NDC RNP	SEQUENCE TRYGRPHSG MRTRHSTGG IAFKIVSQE IKFVKNMNR VKYVMGRND ITYRISGVG LLIYDNEGA IAKITSDYQ	C RLG TNK PAG DST GGY IDQ DAT ATQ	MW (Da) 1012.1 984.09 1016.21 1131.39 1063.23 947.1 989.1 1020.15	SCORE 17.99 17.387 15.149 15.046 14.554 13.746 12.572 12.438	40.33 9 38.98 9 33.96 9 33.73 9 32.63 9 30.82 9 28.19 9 27.89 9	

Figure 2. Predicted CD4 T cell epitopes in Desmoglein 3 (Dsg 3). Potential Dsg 3-specific CD4 T cell epitopes restricted by HLA-DQ7(DQB1*0301) and HLA-DR β 1*0402 are shown in yellow and green, respectively. T cell epitopes that are predicted to bind both HLA-DR β 1*0402 and HLA-DQ7(DQB1*0301), are show in red. All shown peptides exceed the Binding Threshold (see Methods for details). POS. = Position; N = N-terminal; C = C-terminal; MW = Molecular Weight (Daltons); %OPT. = Optimal Score.

00485.3 alpha 1 type XVII collagen us pemphigoid autoantigen BP180	Bullo to DO			higoid auto	pantig	gen BP	180 bin	ding	
	Optima	I Scor	e: 45.6	Qβ1*0301, Con 71, Binding Th d in red represe	reshol	d: 11.70		/HAWH,	
We show the state of the sta	RANK	POS.	N	SEQUENCE	С	MW (Da)	SCORE	% OPT.	
kknkrdgtevterivtetvttrltslppkggtsngyaktaslgggsrlekqslthg	1	505	LER	IRRSILPYG	DSM	1056.29	19.68	43.09 %	
/instgstrghastss <mark>yrrahspas</mark> tlpnspgstferkthvtrhayegsssgnsspe rkefassstrgrsgtrese <mark>lrvrlgsas</mark> pstrwtelddvkrllkgsrsasvsptrnss	2	1055	GET	FDYSELASH	VVS	1050.11	19.381	42.44 %	
kkgtvetkivta <mark>ssgsvsgty</mark> dat <mark>ildanlpsh</mark> vwsstlpagssmgtyhnnmtt	3	1283	RLL	STDASHSRG	SSS	898.89	16.697	36.56 %	
aysagsvfgvpnnmascsptlhpglstsssvfgmqnnlapslttlshgttt	4	211	DAT	ILDANLPSH	VWS	961.09	16.398	35.90 %	
spaavntgvstsaacttsvqsddllhkdckflilekdntpakkemel spasiaatsfsedtlkkekgaaynadsglkaeangdlktvstkgktt	5	841	PGP	AGPAGLPGH	QEV	757.85	16.234	35.55 %	
aatsisedtikkekqaaynadsgikaeangdiktvstkgktt ggvggagggpwgpapawcpcgsccswwkwllgl	6	220	PSH	VWSSTLPAG	SSM	876.01	15.634	34.23 %	
elerirrsilpygdsmdriekdrlqgmapaagadldkigl	7	1320	GAG	SLGAGGAFG	EAA	717.78	13.849	30.32 %	
engnlrgspgpkgdmgspgpkgdrgfpgtpgipgplghpg ggrgregpmgprge <mark>agppgsgek</mark> gergaagepgphgppgv	8	765	MPG	IRGPPGPSG	DPG	818.94	13.625	29.83 %	
glqglrgevglpgvkgdkgpmgppgpkgdggekgprgl	9	199	VTA	SSQSVSGTY	DAT	896.91	13.531	29.63 %	
pagpdghqgprgeqgltgmpgirgppgpsgdpgkpgl	10	247	SLL	NTNAYSAGS	VFG	865.85	13.391	29.32 %	
gkivtsegssmltvpgppgppgamgppgppgapgp	11	80	TSS	YRRAHSPAS	TLP	1026.14	12.928	28.31 %	
rgppgpsipgppgprgppgeglpgppgpgsflsns prghqgeqglpqfstsgsssfglnlggppgppgpg	12	1201	VED	LSSYLHTAG	LSF	930.03	12.403	27.16 %	
ggssstmyvsgppgppgppgppgsisssgqeiqqyise	13	423	TTA	DIHSYGSSG	GGG	903.91	12.396	27.14 %	
pgppgpvttitget <mark>fdyselash</mark> vvsylrtsgygvslfss rgylmgprgppgppgasgdgsllsldyaelssr <mark>ilsymss</mark>	14	837	PPG	APGPAGPAG	LPG	675.75	11.938	26.14 %	
1141 misiglpgppgppglpgtsyeellsllrgsefrgivgppgppgpgpgpgipgnvwssisved 1201 lssylhtadlsfipgppgpgpgpgpgpgpggpggalatyaaensdsfrselisyltspdv 1201 sfivgppgppgpgpgpgsdsrllstdashrgsssshssvrgssyssmstggggag 1321 lgaggafgeaagdrgpygtdigpgggygaaaeggmyagnggllgadfagdldynelavrv 1381 sesmqrqgllqgmaytvggppgqpgpgpgiskvfsaysnvtadlmdffqtygaiggp 1441 gqkgemgtpgpkgdrgpagppgppgppgpkgekgdkgdqvyagrrrrsiavkp	Bullous pemphigoid autoantigen BP180 binding to DRβ1*0402 Matrix: HLA (DR4) DRβ1*0402, Consensus sequence: ICFWHNHNM, Optimal Score: 44.604, Binding Threshold: 11.44 All rows highlighted in red represent predicted binders.								
	RANK	POS.	N	SEQUENCE	С	MW (Da)	SCORE	% OPT.	
	1	1134	SSR	ILSYMSSSG	ISI	926.06	16.967	38.04 %	
					110	1000.00	And in case of the local division of the loc		
	2	472	LGL	LLTWLLLLG	LLF	1000.32	15.227	34.14 %	
	2 3	472 142	LGL ESE	IRVRLQSAS	PST	1000.32	15.227 14.347	34.14 % 32.17 %	

Figure 3. Predicted CD4 T cell epitopes in Bullous Pemphigoid Antigen 2 (BP180). Predicted BP180-specific CD4 T cell epitopes restricted by HLA-DQ7(DQB1*0301) and HLA-DR β 1*0402 are shown in yellow and green, respectively. T cell epitopes that are predicted to bind both HLA-DR β 1*0402 and HLA-DQ7(DQB1*0301), are show in red. All shown peptides exceed the Binding Threshold (see Methods for details). POS. = Position; N = N-terminal; C = C-terminal; MW = Molecular Weight (Daltons); %OPT. = Optimal Score.

These T cells, through CD40-CD40L interaction, then stimulate two different B cells to produce two different antibodies. The second scenario may be that an Antigen Presenting Cell (APC) may internalize two separate antigens. The APC then stimulates 2 different T cells, each specific to either of the antigens [107–109]. The third scenario may be that a macromolecule complex of 2 or more antigens is internalized by an APC that stimulates two different T cells, making each specific for one antigen within the macromolecule [107,110–112].

Through CD40-CD40L interaction, B cells are enlisted resulting in the production of two different autoantibodies. Finally, the fourth scenario may be that the surrounding tissue gets damaged as a result of the presence of one autoimmune disease [107,113,114]. The damage exposes the immune system to a previously sequestered epitope, which is then internalized by an APC that stimulates a T cell that, in turn, recruits a B cell resulting in the production of a second antibody.

In patients with Stevens-Johnson syndrome, it has been demonstrated that if extensive inflammation and damage to surrounding tissue ensues, the patients often develop OCP [115]. Moreover, it has been found that the amount and type of antibodies produced are influenced by the MHC II genes [116]. Furthermore, studies indicate that binding groove flanking residues of core nonameric peptides play a role in peptide selection because the perfect fitting of the peptide residues within the HLA II allele binding groove cannot ensure that the whole peptide will fit completely [66].

Based on the studies by Reche et al. [54-56] using RANKPEP program, we have shown that MMP, PV, and MCTD antigens bear CD4 T cell epitopes restricted by DRβ1*0402 and DQβ1*0301 alleles that can stimulate an immune response (Figures 1, 2, and 3). There are patients who simultaneously have PV and MCTD and MMP and MCTD but lack the HLA II genes associated with MCTD. Nonetheless, the sera of these patients contain antibodies to snRNP70 and have the clinical features of MCTD. Therefore, within the four scenarios, it is theoretically possible for the patients who lack the HLA II genes associated with MCTD to benefit from the presence of DRβ1*0301 to produce autoantibodies to snRNP70 based on epitope spreading. This explanation provides a possible molecular basis to the presence of these two autoimmune diseases occurring in the group of patients described in this manuscript. Also, Since $DQ\beta1*0302$ and *0603 alleles have peptide binding specificities similar to $DQ\beta1*0301$, it is also hypothetically possible for those alleles to produce a similar effect as *0301.

Although the model presented in this manuscript is hypothetical, it provides a basis in which it demonstrates that the molecular structure and the binding properties of the HLA II genes and the possible T cell epitopes within the relevant antigens may play a critical role in the simultaneous presence of two autoimmune diseases in the patients described in the study. Also, it follows that the T and B cells stimulated are specific leading to the production of specific autoantibodies. It is important to highlight that the authors are aware that this is not the only explanation to the observed phenomena, but are utilizing the available data in order to propose a possible explanation. It is also possible that other unknown genetic or non-genetic factors may have resulted in the production of two autoimmune diseases simultaneously in the same patient.

The genetic factors were emphasized because the patients are all unrelated. The authors acknowledge that the major limitation of this study is the fact that it lacks in-vitro studies in which T cells from these patients were simultaneously stimulated with the relevant T cell epitopes described in the manuscript, and proliferative responses were demonstrated. However, the authors felt that providing the information from this computer model to the readership would be of significant benefit to investigators in this field, because they have provided a rational basis for designing and conducting future experiments.

These observations are of clinical and biological importance. These studies demonstrate that, in some individuals with an inherited genetic predisposition towards autoimmune disease, the immune system may be aversely stimulated to produce two autoimmune diseases. Conventional therapy may then be less effective, and the course of the illness may become more chronic. This combination of two autoimmune diseases occurring simultaneously has the potential to influence the patient's morbidity and possible mortality, and poses a challenge to the treating physicians.

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