A Novel Class of Anti-DNA Antibodies Identified in BALB/c Mice

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Summary

We have characterized four IgG monoclonal antibodies (mAbs) derived from BALB/c mice that bind double-stranded DNA (dsDNA) with high affinity. The hydridomas were selected for expression of a member of the V_H S107 family. Three of the four cell lines use the V_H 11 gene and one uses the V_H 1 gene. These antibodies exhibit many characteristics of pathogenic anti-DNA antibodies. They are high affinity and not broadly crossreactive. Unlike the anti-DNA antibodies in autoimmune mice, they exhibit no somatic mutation in their V_H genes. These results demonstrate that somatic mutation of V_H S107 genes is not necessary for generating high affinity dsDNA binding. The fact that such antibodies have not previously been reported suggests that they are rare and that their expression may be downregulated in both nonautoimmune and autoimmune individuals.

Recent analyses of autoantibodies have established that there are two qualitatively different classes of autoantibodies. One class is the "natural" autoantibodies, made by autoantibody-producing B cell lines derived from both normal individuals and individuals with autoimmune disease; the other class can be derived only from autoimmune individuals. The physiologic function of the ubiquitous "natural" autoantibodies is not known and it is assumed that they have no pathogenic potential (1-5). "Natural" autoantibodies have been induced in normal and autoimmune strains both in vitro and in vivo by stimulation with bacterial LPS, a polyclonal B cell activator (6-9). Monoclonal autoantibodies derived from LPSstimulated B cells are generally of the IgM isotype, are widely crossreactive with a variety of autoantigens and exogenous antigens, and are encoded by unmutated germline genes. Most of these antibodies are low affinity, but some may exhibit moderate binding to a particular autoantigen (10). However, because they are pentameric IgM, it is difficult to attribute this binding to true affinity. It has been suggested that these autoantibodies are the product of a particular B cell lineage that displays the CD5 (Ly1) marker on the membrane (11, 12).

In contrast, $(NZB \times NZW)F_1$ and MRL/lpr autoimmune strains spontaneously produce various autoantibodies that differ in a number of parameters from natural autoantibodies. Autoimmune animals produce high affinity autoantibodies of the IgG isotype that are not widely crossreactive and are encoded by somatically mutated genes (13-16). Autoantibodies with these characteristics are a hallmark of both the human disease SLE and of the (NZB \times NZW)F₁ and MRL/lpr mouse models of the human disease. Furthermore, these autoantibodies are believed to have pathogenic potential, since autoantibodies eluted from diseased kidneys of autoimmune mice exhibit the same characteristics: they are IgG, high affinity, and not broadly crossreactive (1, 17–21). Autoantibodies with these characteristics have not previously been found in nonautoimmune strains. Some investigators have speculated that these disease-associated autoantibodies may be the somatically mutated progeny of natural autoantibodies (22).

In this paper we describe four anti-DNA antibodies secreted by hybridomas derived from BALB/c mice. Three of the four hybridoma antibodies reported here were derived from animals immunized with an anti-I-J antibody (23), and a fourth was produced in an animal that received a single immunization of phosphorylcholine (PC)¹ coupled to a protein carrier. These anti-dsDNA hybridomas were selected for expression of a member of the \$107 V_H family. The \$107 gene family has four members, (V_H1, V_H3, V_H11, and V_H13) and is known to be important in the protective immune responses to pneumococcal bacteria $(V_H 1)$ and to influenza $(V_H 11)$ in BALB/c mice. Previous studies showed that a single amino acid substitution in the PC-binding, V_H1-encoded myeloma cell line S107, resulted in production of the U4 antibody, which no longer bound PC but had acquired reactivity with dsDNA (24). The analysis of the U4 mutant suggested that the S107 gene family might contribute to the in vivo produc-

¹ Abbreviations used in this paper: ARS, p-azo-phenylarsonate; HGG, human gammaglobulin; PC, phosphorylcholine.

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tion of anti-DNA antibodies and studies of serum and hybridoma antibodies have shown that the S107 family is indeed used in the anti-DNA response in both the (NZB \times NZW)F₁ and MRL/lpr strains of mice (13–15, 25). The antibodies reported here have features of autoantibodies found in autoimmune disease; they are IgG, have high affinity for dsDNA and are not broadly crossreactive. They differ from the autoantibodies found in autoimmune disease because they are encoded by germline V_H genes containing no somatic mutations. Two are encoded by germline V_k genes also; the germline V_k genes for the other two antibodies are not known so it is not possible to determine whether the expressed V_k genes are unmutated, although this is likely to be the case. Autoantibodies with these characteristics in autoimmune disease display extensive somatic diversification.

The detailed analysis of these four antibodies leads us to posit a new third class autoantibody, a class of highly specific IgG anti-DNA antibodies that are encoded by nonmutated germline genes and that may not generally be expressed either in normals or in individuals with lupus-like autoimmune disease. The existence of these antibodies raises new questions about the nature of germline gene-encoded antibodies and the regulatory defect that may be present in autoimmunity.

Materials and Methods

Cell Lines. 8-wk-old female BALB/c mice were immunized intraperitoneally with either anti-I-J^d 50 μ g in CFA or PC conjugated to human gammaglobulin (HGG), 100 μ g in CFA. Mice were boosted intravenously with either 100 μ g PC coupled to KLH, 100 μ g p-azo-phenylarsonate (ARS)-KLH, or 100 μ l of an antiserum to the T15 idiotype (Table 1). The antiidiotype was prepared as follows: rabbits were immunized with S107 protein and high titered antiserum was then absorbed on an irrelevant mouse IgA antibody (W1329)-Sepharose to render it variable region specific. Spleen cells from the immunized mice were fused to the nonproducing HAT-sensitive myeloma cell line X63Ag8.653 (26). Hybridomas were plated in 96-well tissue culture plates in HATcontaining medium and screened 2 wk later by RNA dot blot for expression of a T15 V_H gene.

RNA Dot Blot Analysis. Cells were lysed in 96-well plates and filtered through nitrocellulose with a 96-well manifold (Schleicher & Schuell, Keene, NH) and hybridized to a 210-bp nick-translated probe that detects all members of the S107 V_H gene family (27, 28).

mRNA Preparation and Sequencing. Total RNA was isolated from hybridoma cells by extraction with guanidinium thiocyanate. Samples were dissolved in 10 mM Tris-Cl, 1 mM EDTA, 400 mM NaCl heated to 68°C and loaded three times onto oligo(dT)cellulose columns (Collaborative Research Inc., Bedford, MA). mRNA was eluted with TE (10 mM Tris-Cl, 1 mM EDTA) and precipitated with ethanol. The precipitate was dissolved in mRNA annealing buffer (250 mM KCl, 10 mM Tris-Cl, 1 mM EDTA) for sequencing. mRNA was annealed to radiolabeled Ig constant region oligonucleotide primers and dideoxy sequencing was performed according to Geliebter et al. (29).

Normalization of Cell Supernatants for Ig Concentration. Cells were grown in serum-free medium containing 5.0 μ g/ml bovine insulin, 3.5 μ g/ml human transferrin, 0.1% sodium selenite, and 0.1% ethanolamine, in a 1:1 mixture of DME (1,000 mg glucose per liter) and Ham's F12 nutrient mixture (Hazelton, St. Lenexa, KS). The Ig content of each culture supernatant was determined by ELISA. 96-well polystyrene E.I.A. plates (Costar Corp., Cambridge, MA) were coated for 2 h at 37°C with an anti-IgG isotypic reagent (Fisher Biotech, Orangeburg, NY) appropriate for each cell line. Plates were blocked with 1% BSA in PBS overnight at 4°C, and cell supernatants or dilutions of commercially purified Igs (Litton Bionetics, Charleston, SC) were added and the plates were incubated for 90 min at 37°C. Plates were washed three times with PBS, 0.05% Tween, and then incubated with alkaline phosphatase-conjugated anti-mouse IgG isotypic reagent for 90 min at 37°C. Plates were washed again and developed with alkaline phosphatase substrate (Sigma Chemical Co., St. Louis, MO). Plates were read in a Titertek ELISA reader (Flow Laboratories, McLean, VA) at 405 nm.

Millipore Filter Assay. Nick-translated DNA was made doublestranded by passage through a $0.45-\mu$ m nitrocellulose Millex syringe filter (Millipore Products Division, Bedford, MA). Cell supernatants normalized for Ig concentration were incubated with 10,000 cpm of ³²P-labeled, dsDNA in 96-well nitrocellulosebottom Millititer-HA plates (Millipore Products Division) for 90 min at 37°C. The dsDNA/cell supernatant mixtures were then filtered through the nitrocellulose and washed three times with 1× SSC (0.15 M NaCl, 0.015 M citrate, pH 7.2) on a 96-well vacuum filtration apparatus (Millipore Products Division). The wells were punched out and counted on an LKB-RackBeta (Pharmacia LKB Biotechnology, Inc., Piscataway, NJ) scintillation counter.

DNA ELISA. Round-bottomed, polyvinyl plates (Falcon Labware, Oxnard, CA) were coated with 100 μ l per well of poly-Llysine (50 μ g/ml), and incubated for 2 h at 37°C. Plates were shaken dry and rinsed with distilled water. 100 μ l of calf thymus (CT) DNA either filtered (as above for dsDNA) or boiled for 10 min

Cell line	Immunization protocol	Isotype	V _H	D _H	J _H	Vĸ	Jĸ
2B11.1	αI-J ^d , 5 wk Ars-KLH, 4 d	IgG2b	1	DFL16.2	3	8	1
C8.5	αI-J ^d , 7 wk αT15, 4 d	IgG1	11	DFL16.1	1	V _k 20	5
7D2.G12	αI-Jª, 4 wk PC-KLH, 3 d	IgG3	11	DFL16.1	1	12, 13	1
R4A.12	PC-HGG, 1 wk	IgG3	11	?	4	1	1

Table 1. Anti-dsDNA mAbs from BALB/c Mice

and quickly cooled on ice (ssDNA) was added at 4°C overnight. The plates were blocked with 1% BSA in PBS for 90 min, washed three times with PBS, 0.05% Tween 20, and then 50 μ l of normalized serum-free supernatant or 50 μ l of serum of 1:500 dilution in 1% BSA/PBS was applied. After a 90-min incubation and washing, a goat anti-mouse κ chain reagent or a sheep anti-mouse IgG reagent conjugated to horseradish peroxidase (Fisher Biotech, Orangeburg, NY) was applied and incubated for 90 min at 37°C. Plates were then washed again, ABTS peroxidase substrate (Kirkegaard and Perry Laboratories, Gaithersburg, MD) was applied, and color development was monitored at 405 nm.

Inhibition of dsDNA Binding. 100 μ l per well of CT DNA (ds or ss) at various dilutions in PBS was preincubated with 15 μ l of serum-free cell supernatant normalized to 40 μ g/ml (final concentration 6.3 μ g/ml) at 37°C for 1 h in a 96-well, nitrocellulosebottomed plate. 10 μ l containing 15,000 cpm of ³²P-labeled dsDNA was added per well and the incubation was continued for an additional hour at 37°C. The mixtures were filtered through the nitrocellulose on a 96-well vaccuum filtration apparatus and the wells were washed and counted as above.

ELISAs for Other Antigenic Specificities. Binding to thyroglobulin, mycobacteria 65-kD protein, PC-BSA, p-aminophosphorylcholine, and L-o-glycerophosphorylcholine was measured as follows. Antigen was diluted to 10 μ g/ml PBS and 50 μ l was added to each well of a 96-well polystyrene EIA plate (Costar Corp.) for 2 h at 37°C. The plates were blocked with 100 μ l per well of 1% BSA in PBS at 4°C overnight. Plates were washed three times with BSA 0.05% Tween 20. Samples were added at concentrations ranging between 1 and 20 μ g/ml (50 μ l per well) at 37°C for 90 min, washed three times with PBS, 0.05% Tween 20 and then incubated with a peroxidase-linked anti-mouse κ chain reagent (Fisher Biotech) for 90 min at 37°C followed by three washes with PBS 0.05% Tween 20 and a final rinse with dH₂O. Plates were developed with hydrogen peroxide and ABTS (Kirkegaard and Perry Laboratories) and read at 405 nm.

For analysis of binding to phosphatidylethanolamine, phosphatidylinositol, and phosphatidylserine, antigens were diluted to 10 μ g/ml in methanol and 50 μ l per well of antigen was added to polystyrene plates. The plates were allowed to dry overnight at room temperature, then were blocked for 2 h at 37°C with 1% BSA in PBS. The remainder of the assay was performed as above.

For analysis of binding to myosin, mouse cardiac myosin was diluted to 2.5 μ g/ml in a buffer containing 36.6 mM NaHCO₃, 12.8 mM Na₂CO₃, and 50 mM tetrasodium pyrophosphate. Polystyrene plates were coated with 100 μ l of myosin solution per well for 1 h at 37°C. The buffer was then discarded and the plate was allowed to dry overnight (30). The plate was blocked with 1% BSA/PBS for 2 h at 37°C and the assay performed as above.

For analysis of binding to influenza, polystyrene plates were coated with 50 μ l of 2.5 \times 10⁵ HAU/ml of influenza virus strain PR8. The plates was dried overnight at room temperature and given a 15-min wash with methanol followed by two PBS washes. The plates were then blocked as usual with 1% BSA/PBS and the assay was performed as above.

Results

BALB/c mice were immunized using two protocols intended to elicit anti-DNA antibodies encoded by members of the S107 heavy chain variable region gene family (V_H family). One strategy of immunization was designed to permit expression of anti-dsDNA antibodies by attempting to in-

terfere with normal suppression of autoimmunity. This protocol involves the use of an anti-I-J^d antibody, WF18.2B15 (23). The I-J molecule, present on suppressor T cells, is believed to play a role in mediating suppressor cell activity since antibody to I-J can block induction of suppressor activity. Our laboratory has previously shown that polyclonal antiidiotype to anti-I-I can abrogate suppression in an in vitro assay system (23). In preparing monoclonal antiidiotypic reagents to anti-I-J^d, we observed high-titered anti-dsDNA antibodies as well as antiidiotype in BALB/c (H-2^d) mice immunized with anti-I-J^d. The anti-DNA response of several groups of mice immunized with anti-I-J^d was monitored by ELISA. Results from one group are illustrated in Fig. 1. The repetitive cycling of the anti-DNA response over a period of several months was reproduced in large numbers of mice. The high-titered anti-DNA activity in the anti-I-J^d immunized mice led us to use this model to generate anti-dsDNA mAbs in BALB/c mice. Subsequent analysis showed that many of these animals also had elevated S107 idiotype in their serum. In an effort to bias towards the production of S107 V_{H} encoded anti-dsDNA antibodies, we boosted the mice with PC-KLH or anti-S107 idiotype. As a control, we boosted with the hapten ARS that has also been implicated in an anti-DNA response but elicits antibodies encoded by a different V_H gene family (5). The second strategy for the production of S107-encoded anti-DNA antibodies was to immunize BALB/c mice with PC in an attempt to isolate U4-like antidsDNA mAbs that might arise by in vivo somatic mutation of anti-PC antibodies encoded by an S107 V_H gene.

Spleen cells from the immunized mice were fused to the nonproducing myeloma cell line X63Ag8.653 by conventional methods. The hybridomas were screened first by RNA dot blot using a heavy chain probe that detects all members of the S107 V_H gene family. Supernatants from wells expressing an S107 heavy chain were analyzed for isotype. We were in-



Figure 1. Mouse sera were diluted 1:500 and analyzed by dsDNA ELISA. The ELISA was developed with a peroxidase-conjugated anti-mouse IgG reagent. The mean dsDNA binding plus 2 SD of a panel of eight unimmunized age-matched BALB/c has been subtracted from the values that are shown.

Table 2.	Antigen	Binding
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Cell line	dsDNA binding
	cpm
2B11.1	1,950
C8.5	1,211
7D2.G12	1,069
R4A.12	1,218
S107	304
U4	1,008

Hybridoma cell supernatants were analyzed by millipore filter assay at 2 μ g/ml. The S107 parental (IgA) non-DNA binding cell line and the U4 mutant (IgA) DNA binding cell line are described elsewhere (24).

terested primarily in T15 antibodies of the IgG isotype. All T15 IgG antibodies were, therefore, analyzed for their ability to bind dsDNA by millipore filter assay. Four IgG-secreting hybridoma cell lines were isolated and are described in Table 1. The anti-DNA activity of the antibodies secreted by these cell lines was compared with that of the S107 (PC-binding) IgA antibody and the U4 (dsDNA binding) mutant IgA antibody (Table 2). Each antibody bind DNA as well as or better than the dimeric IgA U4 antibody and equivalently to a 1:1,000 dilution of serum from an autoimmune mouse (data not shown).

The heavy and light chain variable regions of these four antibodies were analyzed by mRNA dideoxy sequencing and the results are shown in Figs. 2 and 3. Fig. 2 a shows the heavy chain variable region genes compared to their corresponding germline genes. All of the antibodies use unmu-

A	IgG T15 ANTI-dsDNA ANTIBODIES		C VH1
VH1 2B11.1 VH11 C8.5 7D2.G12 R4A.12	GAGGTGAAGCTGGTGGAATCTGGAGGAGGCTTGGTACAGCCTGGGGGTTCTCTG	AGACTCTCCTGTGCAA 70	2811. VH11 C8.5 702.0 R4A.1
VH1 2B11.1 VH11 C8.5 7D2.G12 R4A.12	CTTCTGGGTTCACCTTCAGTGĂŢ <mark>TTCTACATGGĂ</mark> ĞTGGGTCCGCCAGCCTCCAGC	GGAAGAGACTGGAGTG 140 -AGCT	VH1 2B11. VH11 C8.5 7D2.0 R4A.1
VH1 2B11.1 VH11 C8.5 7D2.G12 R4A.12	GATTGCTGCAAGTAGAAACAAAGCTAATGATTATACAACAGAGTACAGGTACAGTGCATCT -T-G-G-TTT-T	GTGAAGGGTCGGTTC 210	VH1 2B11. VH11 C8.5 7D2.0
VH1 2B11.1 VH11 C8.5 702.G12 R4A.12	ATCGTCTCCAGAGACACTTCCCAAAGCATCCTCTACCTTCAGATGAATGCCCTG -C-AT-ACATCACA	AGAGC TGAGGACACTG 280	R4A.1 VH1 2811. VH11
VH1 2B11.1 VH11 C8.5 7D2.G12 RA4.12 B	CCATTTATTACTGTGCAAGAGAT C	30.3	CB.5 7D2.0 R4A.1 VH1 2B11. VH11 CB.5 7D2.0 R4A.1
2811 .1	N DLF16.2 N JH3 ATTACTACGGCTAC TTTGCTTACTGGGGCCAAGGGACTGTGGTCAC -A GGTT	D in Frame TGTCTCCGCAG 	2811. C8.5 7D2.0 R4A.1
C8.5 7D2.G12	N DFL16.1 N JH1 TATTACTACGGTAGTAGCTAC CTGGTACTTCGATGTCTGGGGCGCAGGGACCA 	CGGTCACCGTCTCCTCAG +	
R4A	N D? N JH4 CTATGGACTACTGGGGTCAAGGAACCTCAGTC AGGATCC	ACCGTCTCCTCAG	Figu mAl prop

C																						
VH1 2811.1	E -	v -	к -	L -	V -	E -	s -	G -	G -	10 G -	L -	v -	Q -	Р -	G -	G -	s -	L -	R -	20 L -	s -	с -
VH11 C8.5 7D2.G12 R4A.12							-															
VH1 2811.1	A _	T -	s -	G -	F -	т -	F	30 S -	D	F -	Y -	M -	E -	W	v -	R -	Q -	40 P -	Р -	G -	к -	R -
VH11 C8.5 7D2.G12 R4A.12								T - - -		Y - -	s - -	-										A - - -
VH1 2B11.1	L -	E	W -	1 -	A -	50 A -	s -	52 R -	A N -	В К -	C A -	N -	D	Y	T -	T -	E -	Y -	60 S -	A -	s -	v -
VH11 C8.5 7D2.G12 R4A.12				L - -	G - -	F - - -	I - - -	-					G - -			-			-			
VH1	ĸ	G	R	F	I	v	70 S	R	D	т	s	Q	s	I	ι	Y	80 L	Q	82 M	A N	B A	C L
VH11 C8.5 7D2.G12 R4A.12	-			-	T - -	I - - -	-			N - - -					-						- - -	
VH1 2811.1	R -	A	E -	D -	т -	A -	I -	90 Y -	Y -	с -	A -	R -	95 D		т	т	A	100 т	A V	B F	C	D
VH11 C8.5 7D2.G12 R4A.12					\$ - -		T - - -			-		-	Ā	Y I R	Y N I	Y Y P	G G M	s s	s s	S ₩	L Y	W F
2B11.1 C8.5 7D2.G12	E Y	F	A D D	Y V V		9999	QAA	0000	Ť T T		vvv	110 T T T T	VVV	S S S S	5555							

Figure 2. Heavy chain variable region sequences of BALB/c anti-dsDNA mAbs obtained by mRNA dideoxy sequencing compared with their appropriate V_H , D_H , and J_H genes. Overlining indicates the CDRs.

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A	١							281	11 V _K									c								7D2.	G12 V	ĸ							
	CGC Arg	GCT Ala	GTG Val	FR1 1 GAC Asp	ATT Ile	GTG Val	ATG Met	TCA Ser	CAG G1n	TCT Ser	CCA Pro	TCC Ser	LO TCC Ser	CTA Leu	GCT Ala	GTG Val	TCA Ser		GTT Vai	GTC Va1	AGA Arg	TGT Cys	FRI 1 GAC Asp	ATC Ile	CAG Gln	ATG Met	ACT Thr	CAG Gln	1C1 Ser	CCA Pro	GCT A1a	10 TCA Se r	CTG Leu	fCT Ser	GCA Ala
GTT Val	GGA G1y	GAG Glu	AAG Lys	GTT Val	20 ACT Thr	ATG Met	AGC Ser	TGC Cys	CDR1 AAG Lys	TCC Ser	AGT Ser	27 CAG G1n	27A AGC Ser	27B CTT Ser	27C TTA Leu	270 TA T Tyr	27E AGT Ser	TCT Ser	GTG Val	GGA Gly	GAA G lu	ACT Thr	GTC Val	20 ACC Thr	ATC I le	ACA Thr	TGT Cys	CDR1 GGA Gly	GCA Ala	AGT Ser	GAG G1u	AAT Asn	ATT J le	30 TAC Tyr	GGT Gly
27F AGC Ser	AAT Asn	CAA G1n	30 AAG Lys	AAC Asn	TAC Tyr	TTG Leu	GCC Ala	FR2 TGG Trp	TAC Tyr	CAG Gîn	CAG G1n	AAA Lys	40 CCA Pro	GGG Gly	CAG Gln	TCT Ser	CCT Pro	GCT Ala	TTA Leu	AAT Asn	FR2 TGG Trp	tat Tyr	CAG Gln	CGG Arg	AAA Lys	40 CAG G1n	GGA G1y	AAA Lys	TCT Ser	CCT Pro	CAG Gln	CTC Leu	CTG Leu	ATC Ile	TAT Tyr
AAA Lys	CTG Leu	CTG Leu	ATT Ile	TAC Tyr	CDR2 50 TGG Trp	GCA Ala	TCC Ser	ACT Thr	AGG Arg	GAA G1u	TCT Ser	FR3 GGG G1y	GTC Val	CCT Pro	60 GAT Asp	CGC Arg	TTC Phe	CDR: 50 GGT Gly	GCA Ala	ACC Thr	AAC Asn	TTG Leu	GCA Ala	GAT Asp	FR3 GGC Gly	ATG Met	TCA Ser	60 TCG Ser	AGG Arg	TTC Phe	AGT Ser	GGC Gly	AGT Ser	GGA Gly	TCT Ser
ACA Thr	GGC Gly	AGT Ser	GGA Gly	TCT Ser	GGG G1y	ACA Thr	70 GAT Asp	TTC Phe	ACT Thr	CTC Leu	ACC Thr	ATC Ile	AGC Ser	AGT Ser	GTG Val	AAG Lys	80 GC⊺ Ala	GGT G1y	AGA Arg	70 CAG G1n	TAT Tyr	TCT Ser	CTC Leu	AAG Lys	ATC I 1e	AGT Ser	AGC Ser	CTG Leu	CAT His	80 CCT Pro	GAC Asp	GA⊺ Asp	GTT Val	GCA A1a	ACG Thr
GAA Glu	GAC ASP	CTG Leu	GCA Ala	GTT Val	TAT Tyr	TAC Tyr	TGT Cys	CDR3 CAG G1n	90 CAA G1n	TAT Tyr	TAT Tyr	AGC Ser	TAT Tyr	95 CCT Pro	J _k CGG Arg	ACG Thr	FR4 TTC Phe	TAT Tyr	TAC Tyr	CDR3 TGT Cys	CAA G1n	90 AAT Asn	GTG Val	tta Leu	AGT Ser	GCT Ala	95 CCG Pro	J _k TGG Trp	ACG Thr	FR4 TTC Phe	GGT Gly	GGA Gly	GGC G1y	ACC Thr	AAG Lys
GGA G1y	GGC G1y	ACC Thr	AAG Lys	CTG Leu	GAA G 1u	ATC Ile	AAA Lys	C _k CGG Arg	GCT Ala	GAT Asp								CTG Leu	GAA G1u	ATC Ile	AAA Lys	CGG Arg	C _k GCT Ala	GAT Asp											
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B GTC Val	TCT Ser	GAT Asp	TCT Ser	AGG Arg	GCA Ala	FR1 1 GAA Glu	ACA Thr	C8 ACT Thr	5 V _k GTG Val	ACC Thr	CAG Gîn	TCT Ser	CCA Pro	GCA Ala	10 TCC Ser	CTG Leu	TCC Ser		d TCC Ser	AGC Ser	AGT Ser	FR1 1 GAT Asp	GTT Val	GTG Val	ATG Met	R4A ACC Thr	.12 V CAA Gln	ACT Thr	CCA Pro	CTC Leu	10 TCC Ser	CTG Leu	CCT Pro	GTC Val	AGT Ser
B GTC Val ATG Met	TCT Ser GCT Ala	GAT Asp ATA Ile	TCT Ser GGA Gly	AGG Arg GAA Glu	GCA Ala AAA Lys	FR1 1 GAA G1u GTC Va1	ACA Thr 20 ACC Thr	C8 ACT Thr ATC Ile	GTG GTG Val AGA Arg	ACC Thr TGC Cys	CAG Gin CDR1 ATA Ile	TCT Ser ACC Thr	CCA Pro AGC Ser	GCA Ala ACT Thr	10 TCC Ser GAT Asp	CTG Leu ATT Ile	TCC Ser 30 GAT Asp	CTT Leu	d TCC Ser GGA G1y	AGC Ser GAT Asp	AGT Ser CAA G1n	FR1 GAT Asp GCC Ala	GTT Val 20 TCC Ser	GTG Val ATC Ile	ATG Met TCT Ser	R4A ACC Thr TGC Cys	CAA Gln CDR: AGA Arg	ACT Thr 1 TCT Ser	CCA Pro AGT Ser	CTC Leu 27 CAG G1n	10 TCC Ser 27A AGC Ser	CTG Leu 278 CTT Leu	CCT Pro 27C GTA Val	GTC Val 270 CAC His	AGT Ser 27E AGT Ser
B GTC Val ATG Met GAT Asp	TCT Ser GCT Ala GAT Asp	GAT Asp ATA Ile ATG Met	TCT Ser GGA Gly AAC Asn	AGG Arg GAA Glu FR2 TGG Trp	GCA Ala AAA Lys TAC Tyr	FR1 GAA Glu GTC Val CAG Gln	ACA Thr 20 ACC Thr CAG Gln	C8 ACT Thr ATC Ile	GTG Val AGA Arg 40 CCA Pro	ACC Thr TGC Cys GGG Gly	CAG Gln CDR1 ATA Ile GAA GIu	TCT Ser ACC Thr	CCA Pro AGC Ser CCT Pro	GCA Ala ACT Thr AAG Lys	10 TCC Ser GAT Asp CTC Leu	CTG Leu ATT Ile CTT Leu	TCC Ser 30 GAT Asp ATT Ile	CTT Leu AAT Asn	d TCC Ser GGA G1y GGA	AGC Ser GAT Asp 30 AAC Asn	AGT Ser CAA G1n ACC Thr	FR1 1 GAT Asp GCC Ala TAT Tyr	GTT Val 20 TCC Ser TTA Leu	GTG Val ATC Ile CAT His	ATG Met TCT Ser FR2 TGG Trp	R4A ACC Thr TGC Cys TAC Tyr	CAA GIN CDR AGA Arg CTG Leu	k ACT Thr 1 TCT Ser CAG Gln	CCA Pro AGT Ser AAG	CTC Leu 27 CAG G1n 40 CCA Pro	10 TCC Ser 27A AGC Ser GGC	CTG Leu 27B CTT Leu CAG Gln	CCT Pro 27C GTA Val TCT Ser	GTC Val 270 CAC H1s CCA Pro	AGT Ser 27E AGT Ser AAG Lys
B GTC Val ATG Met GAT Asp TCA Ser	TCT Ser GCT Ala GAT Asp CDR2 50 GAA Glu	GAT Asp ATA Ile ATG Met 22 GGC Gly	GGA GGA Asn AAT Asn	AGG Arg GAA Glu FR2 TGG Trp ACT Thr	GCA Ala AAA Lys TAC Tyr CTT Leu	FR1 1 GAA Glu GTC Val CAG Gln CGT Arg	ACA Thr 20 ACC Thr CAG Gln CCT Pro	C8 ACT Thr ATC Ile AAGG Lys FR3 GGA Gly	GTG GTG Val AGA Arg 40 CCA Pro	ACC Thr TGC Cys GGG Gly CCA Pro	CAG Gln CDR1 ATA Ile GAA Glu 60 TCC Ser	TCT Ser ACC Thr CCT Pro CGA	CCA Pro AGC Ser CCT Pro TTC Phe	GCA Ala ACT Thr AAG Lys TCC Ser	10 TCC Ser GAT Asp CTC Leu AGC Ser	CTG Leu ATT Ile CTT Leu AGT Ser	TCC Ser 30 GAT ASp Ile GGC Gly	CTT Leu AAT Asn CTC Leu	d TCC Ser GGA Gly CTG Leu	AGC Ser GAT ASP 30 AAC ASD 11e	AGT Ser CAA Gîn ACC Thr TAC	FR1 1 GAT Asp GCC Ala TAT Tyr CDR: 50 AAA Lys	GTT Val TCC Ser TTA Leu GTT Val	GTG Val ATC Ile CAT His TCC Ser	ATG Met TCT Ser FR2 TGG Trp AAC Asn	R4A ACC Thr TGC Cys TAC Tyr CGA	.12 V CAA Gln CDR: AGA Arg CTG Leu TTT Phe	k ACT Thr 1 TCT Ser CAG Gln TCT Ser	CCA Pro AGT Ser AAG Lys FR3 GGG Gly	CTC Leu 27 CAG G1n 40 CCA Pro GTC Val	10 TCC Ser 27A AGC Ser GGC G1y CCA Pro	CTG Leu 278 CTT Leu CAG Gln GAC Asp	CCT Pro 27C GTA Val TCT Ser	GTC Val 270 CAC H1s CCA Pro TTC Phe	AGT Ser 27E AGT Ser AAG Lys AGT Ser
B GTC Val ATG Met GAT Asp TCA Ser TAT Tyr	GCT Ala GAT Asp CDR2 GAA Glu GGT GIy	GAT Asp ATA Ile ATG Met 2 GGC Gly ACA Thr	GGA GIy AAC Asn AAT Asn Asp	AGG Arg GAA Glu FR2 TGG Trp ACT Thr Thr Phe	GCA Ala AAA Lys TAC Tyr CTT Leu GTT Val	FR1 1 GAA Glu GTC Val CAG Gln CGT Arg	ACA Thr 20 ACC Thr CAG GIn CCT Pro	C8 ACT Thr ATC Lys FR3 GGA G1y ATT Ile	GTG Val AGA Arg CCA Pro GTC Val GAA Glu	ACC Thr TGC Cys GGG Gly CCA Pro AAC	CAG Gln CDR1 ATA Ile GAA Glu 60 TCC Ser ATG Met	TCT Ser ACC Thr CCT Pro CGA Arg CTC Leu	CCA Pro AGC Ser Pro TTC Phe 80 TCA Ser	GCA Ala ACT Thr AAG Lys TCC Ser GAA Glu	10 TCC Ser GAT Asp CTC Leu AGC Ser GAT	CTG Leu ATT Ile CTT Leu AGT Ser GTT Val	TCC Ser 30 GAT ASp ATT Ile GGC Gly GCA Ala	CTT Leu AAT Asn CTC Leu GGC Gly	d TCC Ser GGA Gly CTG Leu AGT Ser	AGC Ser GAT ASP 30 AAC Asn Ile GGA GIy	AGT Ser CAA GIn ACC Thr TAC Try TCA Ser	FR1 1 GAT ASP GCC Ala TAT Tyr CDR: 50 AAA Lys GGG Gly	GTT Val 20 TCC Ser TTA Leu GTT Val ACA Thr	GTG Val ATC Ile CAT His TCC Ser 70 GAT Asp	ATG Met TCT Ser FR2 TGG Trp AAC Asn TTC Phe	R4A ACC Thr TGC Cys TAC Tyr CGA Arg ACA Thr	.12 V CAA Gln CDR: AGA Arg CTG Leu TTT Phe CTC Leu	k ACT Thr TCT Ser CAG Gln TCT Ser AAG Lys	CCA Pro AGT Ser AAG Lys FR3 GGG Gly ATC Ile	CTC Leu 27 CAG G1n 40 CCA Pro GTC Val	10 TCC Ser 27A AGC Ser GGC Gly CCA Pro	CTG Leu 27B CTT Leu CAG Gìn 60 GAC Asp	CCT Pro 27C GTA Val TCT Ser AGG Arg GAG Glu	GTC Val 270 CAC His CCA Pro TTC Phe 80 GCT Ala	AGT Ser 27E AGT Ser AAG Lys AGT Ser GAG Glu
B GTC Val ATG Met GAT Asp TCA Ser TAT Tyr GAT Asp	GCT Ala GAT Asp CDR22 500 GAA Glu GGT GIy TAC	GAT Asp ATA Ile ATG Met GGC Gly ACA Thr TAC	GGA Gly AAC Asn AAT Asn 70 GAT Asp TGT Cys	AGG Arg GAA Glu FR2 TGG Trp ACT Tr Phe CDR CDR	GCA Ala AAA Lys TAC Tyr CTT Leu GTT Val 3 90 CAA GIn	FR1 1 GAA Glu GTC Val CAG Gln CGT Arg TTT Phe AGT	ACA Thr 20 ACC Thr CAG GIn CCT Pro ACA Thr GAT Asp	C8 ACT Thr ATC Ile AAG Gly ATT Ile AACC Asn	GTG Val AGA Arg 40 CCA Pro GTC Val GTC Val GTC Val	ACC Thr TGC Cys GGG Gly CCA Pro AAC Asn 95 CCT Pro	CAG Gln CDR1 ATA Ile GAA Glu GAA Glu CCC Ser ATG Met J _k CTC Leu	ACC Thr CCT Pro CGA Arg CTC Leu ACG Thr	CCA Pro AGC Ser CCT Pro TTC Phe 80 TCA Ser FR4 TTC Phe	GCA Ala ACT Thr AAG Lys TCC Ser GAA Glu GGT	10 TCC Ser GAT Asp CTC Leu AGC Ser GAT Asp GCT Ala	CTTG Leu ATTT Ile CTTT Leu AGTT Ser GTT Val GGGG Gly	TCC Ser 30 GAT Asp ATT Ile GGC Gly GCA Ala	CTT Leu AAT Asn CTC Leu GGC Gly GAT	d TCC Ser GGA Gly CTG Leu CTG Leu	AGC Ser GAT Asp 30 AAC Asp 30 AAC Ile GGA Gly Gly	AGT Ser CAA Gin ACC Thr TAC Try TCA Ser GTT Cai	FR1 1 GAT ASP GCC Ala TAT Tyr CDR: COR: COR: COR: COR: COR: COR: COR: CO	GTT Val 20 TCC Ser TTA Leu GTT Val ACA Thr TTC Phe	GTG Val ATC Ile CAT His TCC Ser 70 GAT Asp	ATG Met TCT Ser FR2 TGG Trp AAC Asn TTC Phe CDR: TCT Ser	R4A ACC Thr TGC Cys TAC Tyr CGA Arg ACA Thr 3 90 CAA G1n	.12 V CAA GIN CDR: AGA Arg CTG Leu TTT Phe CTC Leu AGT Ser	k ACT Thr 1 TCT Ser CAG Gln TCT Ser AAG Lys ACA Thr	CCA Pro AGT Ser AAG Lys FR3 GGG Gly ATC Ile CAT	CTC Leu 27 CAG Bln 40 CCA Pro GTC Val AGC Ser GTT Val	10 TCC Ser 27A AGC Ser GGC G1y CCA Pro AGA Arg 95 CCT Pro	CTG Leu 27B CTT Leu CAG Gìn GAC Asp GTG Val J _k TGG Trp	CCT Pro 27C GTA Val TCT Ser AGG Arg GAG Glu	GTC Val 270 CAC H1s TTC Phe 80 GCT Ala FR4 TTC Phe	AGT Ser AGT Ser AAG Lys AGT Ser GAG G1u

Figure 3. ~ Light chain sequences of BALB/c anti-dsDNA mAbs obtained by mRNA dideoxy sequencing.

tated germline V_H genes except C8.5, which has a GAT to GCC substitution in V_H 11, thereby generating an alanine residue instead of an aspartic acid at position 101 in the protein sequence. Because these substitutions in C8.5 occur at the D_HJ_H junction, it is probable that these changes arose by addition of N sequence and not by somatic point muta-

tion. Fig. 2 *b* shows the $D_H J_H$ regions of all four antibodies as compared with the germline genes from which they are derived. Only the 2B11.1 antibody has a nucleotide substitution that is unquestionably due to somatic mutation. A substitution in the J_H3 region of CTG for GTG in amino acid number 108 as seen in Fig. 2 *c* causes a conservative amino acid replacement of a leucine for a valine in the fourth framework region. The sequence of the 5' end of the DFL16.2 region in the 2B11.1 antibody is AAT instead of ATT at amino acid position 96 which leads to an isoleucine to asparagine change. This may also represent N sequence rather than mutation. In this antibody the D region is out of frame relative to the most commonly used translation frame. In contrast, the C8.5 and 7D2.G12 DFL16.1 genes are read in the most commonly used frame. Both the C8.5 antibody and the 7D2.G12 antibody have changes at the 3' end of their D regions. The substitution of a TCC for TAC at amino acid position 100B in C8.5 causes a tyrosine to serine change, and the AGC to AGT change at amino acid position 100A in 7D2.G12 is silent. These latter three changes are likely to have resulted from N sequence addition because of their proximity to the VD or DJ junctions. Finally, the R4A.12 antibody uses an extremely short D region that is not related to any of those previously described, therefore its usual reading frame can not be determined, nor can the degree of somatic mutation it exhibits be determined.

The light chain genes expressed by these cell lines are all different from each other, as shown in Fig. 3. The 2B11.1 antibody uses a V_k gene that is a member of the V_k 8 family (Fig. 3 a). Its sequence is completely identical to a productively rearranged but unexpressed light chain gene of the plasmacytoma PC3609 from an NZB mouse whose antigen binding specificity is unknown (31, 32), and it is also 98% homologous to an antilysozyme antibody, Gloop 5 (33). While the 2B11.1 V_K gene is highly homologous to two rearranged V_k genes, it is only 88.3% homologous to a germline V_k gene, V-ser, cloned from liver DNA (34). Therefore, the 2B11.1 V_k is probably derived from an as yet unidentified germline gene and may itself, like the productively rearranged unexpressed but PC3609 V_k , constitute the unmutated sequence. The 7D2.G12 antibody uses a light chain V region that belongs to the V_k 12,13 family (Fig. 3 c). It exhibits greatest

dsDNA INHIBITION



Figure 4. Cold CT-dsDNA (15 μ /ml) was preincubated with serumfree supernatant from hybridoma cell lines for 1 h at 37°C followed by addition of 15,000 cpm nick-translated dsDNA and incubation for 1 h at 37°C. Final concentration of mAb was 6.3 μ g/ml. Serum from an MRL/lpr mouse was diluted 1:1,000 in serum-free supernatant as the DNA binding of this dilution of serum corresponds to the DNA binding of mAb at 6.3 μ g/ml.

homology (84.2%) to a germline gene K2 cloned from the MOPC-149 myeloma and from BALB/c embryo DNA (35). It is likely again that the appropriate V_k germline gene has not yet been cloned and sequenced. The C8.5 cell line uses a V_k gene that is a member of the V_k20 gene family (Fig. 3 b) (36) and the R4A.12 V_k gene is completely identical to a V_k1 germline gene, K5.1, cloned from BALB/c embryo liver DNA (Fig. 3 d) (37).

The relative binding of these S107 V_H-encoded antibodies for dsDNA was analyzed by inhibition assays. The monoclonal antibody concentrations used in these assays were 0.63 μ g per well (Fig. 4). All four antibodies were inhibited to 50% of maximum dsDNA binding by ds-calf thymus DNA concentrations between 0.2 and 1.0 micrograms per well. From these data, one can derive an apparent binding constant of 10⁹ to 10¹⁰ for these antibodies using the equation of Nieto et al. (38).

The four anti-DNA antibodies were also analyzed by ELISAs to determine their antigenic cross-reactivity. Because the antibodies are encoded by the V_H1 and V_H11 gene members of the S107 V_H family, they were analyzed for binding to PC-KLH and influenza, and no binding to either antigen was seen. They were also analyzed for binding to mycobacterium 65kd protein, phosphatidylethanolamine (PE), phosphatidylinositol (PI), phosphatidylserine (PS), L- α -Glycerophosphorylcholine, and *p*-aminophosphorycholine. No binding was seen to any of these exogenous antigens. Binding was analyzed to the autoantigens thyroglobulin and murine cardiac myosin and no binding was demonstrated (Table 3).

Discussion

We have isolated four anti-dsDNA mAbs from BALB/c mice that resemble the pathogenic type autoantibodies

Table 3. Antigen Reactivity on ELISA

	Reactivity on ELISA											
Antigens	2B11.1	C8.5	7D2,G12	R4A.12								
ssDNA	_	-	+/-*	+/								
Thyroglobulin		_	-									
Myosin	_	-	-	_								
Cardiolipin	-	-	÷+/	+/ -								
Influenza	-	_	-									
Mycobacterium	-	_	-	-								
Phosphatidylethanolamine	-	_	-	-								
Phosphatidylinositol	-	-	-	-								
Phosphatidylserine		-	-	-								
L-α-Glycerophosphorylcholine	-	_	-	-								
p-Aminophosphorylcholine	-	-	-	-								

All ELISAs were performed using antibody at a concentration of $10 \,\mu$ g/ml. (+/-) Binding above the mean of irrelevant antibodies, but <2 SD above the mean.

described in autoimmune mouse strains, such as (NZB \times NZW)F₁ and MRL/*lpr*, far more than "natural" autoantibodies found in normal mouse strains. Two strategies were used to induce the production of anti-DNA antibodies in normal mice. One strategy was intended to induce the production of these antibodies through somatic mutation of a response to the bacterial antigen PC. After immunization with PC-KLH we observed that the serological response to dsDNA undergoes a transient rise and fail simultaneously with the response to PC-KLH (data not shown). This coordinate response should occur if the antigens are potentially cross-reactive so that certain epitopes on PC-KLH can also stimulate secretion of anti-DNA antibodies. Alternatively, as suggested by the U4 mutant, somatic mutation of anti-PC antibodies could give rise to anti-DNA antibodies in a PC-immunized host.

The second strategy we used to induce anti-DNA antibodies in BALB/c mice was to immunize with anti-I-J^d antibody in an attempt to interfere with the normal pathways for regulating an immune response (39). In a similar system, Gibson et al, demonstrated that autoantibodies could be induced in the offspring of B10.A(3R)(H-2^b) mothers and B10.A(5R)(H- 2^k) fathers when the pregnant females were immunized with paternal lymphoid cells (40). This immunization induced the production of anti-I-J^k antibodies in the gestating mothers, which crossed the placenta to the offspring and presumably led to the depletion or inactivation of suppressor T cells. When the offspring were immunized with rat red blood cells, they developed a crossreactive response to mouse red blood cells that is not normally seen in naive syngeneic animals. We believe that the long-term cycling (over 30 wk; data not shown) of the level of anti-DNA antibodies that we see in anti-I-J^d immunized BALB/c mice might be due to a similar perturbation in the ability of these animals to suppress an autoimmune response. It is not possible, however, to determine whether a given antibody arose as a result of a particular immunization protocol. More extensive serological and molecular analysis must be undertaken in order to ascertain the role that these immunizations might play in eliciting these antibodies, or whether they might occur in unimmunized animals as well.

Although anti-DNA activity is most commonly measured by ELISA, this method detects low affinity crossreactive antibodies as well as the high affinity, monospecific antibodies that are seen in autoimmune disease (41). We, therefore, chose to measure the binding of our antibodies to dsDNA by a liquid phase assay (millipore filter assay). The millipore filter assay is used as a clinical assay because it is believed to measure the high affinity anti-dsDNA antibodies that bind in solution and titers of anti-DNA antibody measured with this assay correlate with disease activity. All four antibodies display high binding of dsDNA in solution. A low concentration $(0.2-1.0 \mu g)$ of unlabeled dsDNA inhibitor causes a 50% inhibition of binding. These antibodies bind dsDNA with similar avidity to the IgG mAbs that have been isolated from autoimmune strains. Most autoantibodies from nonautoimmune animals bind antigen only at very high antibody concentrations. While it is not possible to obtain an affinity constant for anti-dsDNA antibodies, these antibodies display apparent antigen binding constants of 10⁹ to 10¹⁰, which is, in fact, greater then the binding constants of "natural" germline, gene-encoded pentameric IgM antibodies for DNA. In addition to their high affinity, these antibodies are highly specific for nucleic acid antigens, and unlike IgM "natural" autoantibodies exhibit little or no crossreactivity to a panel of autoantigens and to exogenous antigens.

We analyzed only IgG antibodies as we expected to study the somatic mutations of germline genes of the S107 gene family that can lead to anti-DNA activity. The four autoantibodies described here have V_H sequences that are germlineencoded with limited junctional diversity. Only one antibody, 2B11.1, has a clearly identifiable somatic point mutation in the middle of the J_{H3} gene in framework 4. Three of the four antibodies express the V_H11 gene that encodes most of the S107 V_H-expressing anti-DNA antibodies identified in autoimmune mice. Moreover, 6G6, a V_H11-encoded antibody from a CBA/J mouse (42), was shown to have specificity for dsDNA (43). This antibody is IgM and has only one nucleotide difference from the CBA/J germline V_H11 gene at the 3' end of the antibody, resulting in an aspartic acid to alanine substitution. Therefore, it is possible that the V_H11 germline gene may predispose an antibody to bind DNA, even in an unmutated state.

Three of the four antibodies use a member of the DFL16 family, the significance of which is unclear. No such predominance is seen in the VDJ sequences of anti-DNA antibodies that have been reported from autoimmune mice or from normal mice. However, since these antibodies describe a novel class of autoantibodies that are high affinity for DNA despite being germline gene encoded, and it has been shown that CDR3 is important for DNA binding, it is possible that the usage of the DFL16 family may be significant. It will be necessary to examine more antibodies of this class to understand the importance of specific D region families for DNA binding activity. However, it has been suggested that anti-DNA antibodies often contain in their D regions positively charged amino acids such as lysine, arginine, and histidine or the polar but uncharged amino acids glutamine, glycine, and asparagine which can form covalent bonds to nucleic acid bases (15, 44, 45). An absolute requirement of a given number of these residues for DNA binding to occur has not been established, however, and the presence of one such amino acid in the CDR3 of each of the antibodies may be sufficient for dsDNA binding. Also it has been proposed that the D regions in autoantibodies are often read in an unusual frame (46). Two D regions in these antibodies are unremarkable in this respect; however, a third is read in an uncommon frame and the fourth derives from an unidentified D segment. The D region from R4A.12 antibody is very short and cannot be assigned to any of the previously described D regions. Extremely short D regions have been found in antidextran antibodies, and have also been described in anti-RNA antibodies (15). Because these D's are so short it is often difficult to assign them to a family, or to distinguish D segment from N sequence.

It is more difficult to ascertain whether somatic mutation has occurred in the V_k genes. The R4A.12 V_k gene clearly

expresses an unmutated sequence, while the other three antibodies probably use germline genes that have not yet been cloned and sequenced. Because there is a rearranged V_k sequence that is 100% identical to the 2B11.1 V_k , we believe that the 2B11.1 Vk gene represents a germline gene sequence that has not as yet been cloned. The sequence that is identical to the 2B11.1 V_k is an unexpressed double recombination product from an NZB plasmacytoma, PC3609. The expressed light chain from this IgG2b secreting cell line is a member of the V_K14 family, while the unexpressed light chain is a member of the V_K8 family. Because this rearrangement is not expressed, and has not undergone selection by antigen, the likelihood that it represents an unmutated sequence is increased. The C8.5 and the 7D2.G12 cell lines both use V_k genes that are members of families whose germline sequences have not been completely analyzed, so we cannot know whether these hybridoma Vk sequences represent germline sequences or not. Previous studies, however, have shown that in general somatic mutations in the heavy and light chain variable regions tend to accumulate concurrently. Thus, when the heavy chain variable regions do not exhibit mutations, the light chain variable regions are also unmutated.

We have described a set of anti-DNA antibodies exhibiting unusual properties that do not permit their classification either as "natural" autoantibodies or as autoantibodies associated with autoimmune disease. These antibodies are IgG, bind DNA with high avidity, do not crossreact with non-nucleic acid antigens, and are the products of unmutated Ig genes. These binding and isotypic properties are, in general, associated with the somatically mutated antibodies that are part of autoantibody responses seen in autoimmune disease; however, the apparent absence of somatic mutation that is seen in these antibodies is a property characteristic of "natural" autoantibodies. We are currently investigating the pathogenic potential of these antibodies in BALB/c mice.

These observations generate several questions regarding the regulation of autoantibodies in nonautoimmune and autoim-

mune animals. We have previously suggested that anti-dsDNA antibodies with high avidity for antigen arise in autoimmune individuals and autoimmune mice by somatic mutation of germline Ig genes. Data from many laboratories have confirmed this hypothesis; to date, all anti-dsDNA antibodies from autoimmune animals or patients with SLE reflect the accumulation of somatic mutations. The present data indicate that antibodies that have some of the features associated with pathogenic autoantibodies need not only arise through the accumulation of somatic mutations. Contrary to the observations made previously about germline, gene-encoded autoantibodies, we have shown that such autoantibodies may be neither low affinity nor broadly crossreactive. While it had been suggested that germline gene-encoded anti-dsDNA antibodies are not found because germline genes do not encode such high avidity IgG antibodies, our data suggest that the more likely explanation is that in both normal and autoimmune individuals B cells expressing such antibodies are deleted or rendered anergic when they arise as IgM antibodies in an immature B cell. In autoimmune mice high affinity IgG anti-dsDNA antibodies are routinely found; however, they are always the products of somatic mutation. The nonmutated high binding antibodies described here have not been found, presumably because their regulation is intact.

This might imply two pathways of regulation; one for less mature B cells without somatic mutation and one for more mature B cells that have accumulated somatic mutations. Matzinger and Guerder have formulated a similar hypothesis regarding the differential regulation of immature and mature T cells (47). We hypothesize that in autoimmune disease the defect may exist solely in the regulation of mature B cells whose autospecificity is acquired by somatic mutation. For future studies it will be important to determine the frequency of these monospecific high avidity anti-dsDNA autoantibodies in the preimmune repertoire and to understand their cellular origin and regulation in both nonautoimmune and autoimmune mice.

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