Immunocytochemical Detection of Kappa and Lambda Light Chain V Region Subgroups in Human B-cell Malignancies

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We have used a sensitive immunoperoxidase method and highly specific anti-light chain antisera to determine the light chain variable region (V_L) subgroup nature of cytoplasmic (c) and cell surface (s) Ig expressed by human monoclonal plasma cells and B lymphocytes. The immunocytochemical characterization of cIg and sIg used antisera specific for the established κ light chain V_{κ} subgroups $(V_{\kappa l}, V_{\kappa l l}, V_{\kappa l l l}, and V_{\kappa l V})$ and the λ light chain V_{λ} subgroups $(V_{\lambda l}, V_{\lambda ll/V}, V_{\lambda ll}, V_{\lambda lV}, and V_{\lambda Vl})$. Studies were performed using cytospin preparations of bone marrow-, peripheral blood-, and lymph nodederived cells from patients with multiple myeloma, amyloidosis AL, and Waldenström's macroglobulinemia and with low-, mid-, and high-grade B-cell malignancies. The V_{λ} or V_{λ} subgroup of the cIg or slg also could be identified after deparaffinization and enzyme treatment of formalin-fixed, paraffinembedded specimens. For those patients who had monoclonal serum or urinary Igs, there was complete concordance between the V_L subgroup of the secreted Ig and that of the cIg or sIg. The percentage distribution of V_{κ} or V_{λ} subgroups on the sIg of cells from patients with chronic lymphocytic leukemia (CLL) and other cytomorphologic types of B-cell malignancies differed from that found for κ - or λ type Bence Jones proteins obtained from patients with multiple myeloma, amyloidosis AL, and Waldenström's macroglobulinemia. In contrast to the plasma cell and lymphocytoid plasma cell diseases. a relative predominance of certain V_L subgroups, ie, $V_{\kappa IV}$, $V_{\lambda III}$, and $V_{\lambda IV}$, and the absence of the amyloid-associated $V_{\lambda VI}$ subgroup were found in CLL and related diseases. The immunocytochemical techniques used make possible a rapid means to demonstrate B-cell monoclonality and provide further evidence for the selective expression of certain

V_L genes in buman B-cell neoplasia. (Am J Pathol 1990, 137:855-862)

B-cell-related neoplasms represent monoclonal proliferative diseases that are distinguished by distinctive immunoglobulin (lg) phenotypic properties. The clonal nature of these disorders is evidenced by the expression of lg genes and their monoclonal protein products. One such product, common to all classes of lq, is the kappa (κ) or lambda (λ) light polypeptide chain. Human κ and λ light chains can be classified into multiple variable region (V_1) subgroups on the basis of characteristic amino acid and nucleic acid sequence homologies.¹⁻³ Such subgroups have also been differentiated serologically.^{4,5} Antisera, prepared against monoclonal light chains, can recognize each of the chemically defined V_x and V_{λ} subgroups, designated $V_{\kappa l}$, $V_{\kappa ll}$, $V_{\kappa ll}$, $V_{\kappa lv}$ and $V_{\lambda l}$, $V_{\lambda l l / V}$, $V_{\lambda l l l}$, $V_{\lambda l V}$, $V_{\lambda v l}$ (proteins classified chemically as V_{AV} react with specific anti- $V_{\lambda II}$ antisera). In addition, multiple V_L sub-subgroups have been demonstrated serologically,5,6 and these presumably (like the V_L subgroups) represent the products of single or closely related V₁ genes. A functional role for V₁ subgroups (and sub-subgroups) is shown by the association of particular V₁ genes with certain types of autoantibodies,⁷⁻¹³ amyloidosis AL,¹⁴ and B-cell neoplasms.¹⁵⁻¹⁸

To study further the distribution and importance of V_L subgroups as biologic markers of B-cell lymphoproliferative disorders, we have used highly specific anti-light chain antisera prepared in our laboratory to analyze immunocytochemically the cellular Ig of monoclonal B-cell-related populations found in patients with multiple myeloma, Waldenström's macroglobulinemia, amyloidosis AL, chronic lymphocytic leukemia (CLL), and other forms of malignant lymphoma. These studies have demonstrated the capability of such antisera to identify the V_L

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subgroup nature of cytoplasmic (c) and cell-surface (s) Ig in bone marrow-, peripheral blood-, and lymph nodederived cytospin preparations and in tissue sections. The ability to demonstrate immunocytochemically the V_x or V_λ subgroup provides a unique diagnostic tool for the establishment of B-cell monoclonality, as well as a rapid and effective means of determining V_L gene usage in human B-cell neoplasia.

Methods and Materials

Anti-light Chain Antisera

The methods used in the preparation and characterization of our polyclonal anti-human κ or λ light chain and V_L subgroup-specific antisera have been described previously.⁴ Briefly, albino New Zealand rabbits were immunized with κ or λ Bence Jones proteins or V_L-related fragments and the reactivity of the antisera was determined through comparative immunodiffusion analyses using completely sequenced light chains representative of the human V_x and V_{λ} subgroups. The antisera were rendered specific for κ or λ light chain or V_L subgroup-associated determinants, as established by ELISA and immunoblotting analyses, through extensive absorption with appropriate heterologous light chains and non-lg proteins.

Immunophenotyping

Specimens containing mononuclear cells obtained from bone marrow, peripheral blood, or teased from lymph nodes were enriched by ficoll density gradient (Histopaque-1077, Sigma Diagnostics, St. Louis, MO) and prepared by centrifugation onto clean glass slides using a Shandon cytocentrifuge (Shandon, Inc., Pittsburgh, PA). The slides were fixed for 30 minutes in acetone, air-dried overnight, and washed extensively in a 0.25-mol/l (molar) sodium phosphate, 0.15 mol/l sodium chloride (PBS) buffer, pH 7.2. For intact fresh tissue specimens, $4-\mu m$ sections were cut in a cryostat, placed on 0.5% gelatincoated slides, and dried overnight. They were fixed and washed as previously stated. Immunostaining was performed using the avidin-biotin complex (ABC) method (Vectastain Peroxidase ABC kit, Vector Laboratories, Inc., Burlingame, CA) and serial dilutions of our primary rabbit anti-human light chain antisera, as well as a nonimmune rabbit serum. All were diluted in 1% bovine serum albumin. Slides were flooded with these reagents for 1 hour and washed in PBS. Bound antibody was visualized by exposure of the slides to a peroxidase substrate solution containing the chromagen 3,3' diaminobenzidine (DAB). Lerner-3 hematoxylin (Lerner Labs, New Haven, CT) was

used as the counterstaining reagent, and slides were mounted using a nonaqueous mounting medium (ACCU·Mount 60^{TM} , Scientific Products, McGaw Park, IL).

Formalin-fixed, paraffin-embedded bone marrow was cut on a microtome to a thickness of 6 μ m and mounted on 0.01% poly-L lysine-coated slides. The sections were deparaffinized using a xylene-related solvent (AmeriClear[™], Scientific Products) and brought through a series of graded alcohols to PBS. Sections were washed extensively and to block endogenous peroxidases, slides were submerged in 0.15% hydrogen peroxide/anhydrous methanol for 30 minutes. Next the bone marrow was pretreated with ficin (GammaZyme-F[™], Gamma Biologicals, Inc., Houston, TX) for 30 minutes at 37° C. The sections were overlaid with appropriate dilutions of rabbit anti-human V₁ antisera and incubated overnight at 4° C. Detection and visualization of bound antibody was accomplished using the ABC method, with DAB used as the substrate chromagen.

Clinical analyses were done using a direct and an indirect immunoperoxidase method. Cells or tissue from patients were routinely examined with commercially available primary reagents, which included peroxidase-conjugated sheep polyclonal anti- κ and anti- λ light chain antisera and anti- μ , - δ , - γ , and - α heavy chain antisera (Tago, Inc., Burlingame, CA) and with unconjugated mouse monoclonal anti-LEU 1 (CD5), -LEU 14 (CD22), and -LEU 4 (CD3) antisera (Becton-Dickinson, Mountain View, CA). For the monoclonal antibodies, a biotin-labeled, horse anti-mouse IgG antiserum (Vector) was used as the secondary reagent.

Proteins

An agarose gel immunofixation procedure (Paragon Electrophoresis System, Beckman, Brea, CA) was used routinely to examine serum and urine specimens for the presence of monoclonal lg components. Because of the generally low amounts of urinary Bence Jones proteins in patients with CLL or malignant lymphoma (in contrast to those with multiple myeloma), 100-ml volumes of randomly collected or 24-hour urine specimens were dialyzed exhaustively against deionized, distilled water and then lyophilized. Before electrophoresis and immunofixation, the urine specimens were reconstituted in water to a final protein concentration of 50 mg/ml. The V_L subgroup of monoclonal urinary κ or λ chains was determined by comparative immunodiffusion analyses using anti-V₁ subgroup-specific antisera.⁴ For these analyses, urine specimens were appropriately diluted to yield a final monoclonal light chain concentration of 0.5 to 2 mg/ml.

Proteins	Anti-κ (R 360)†	Anti-V _{×I} (R 374)	Anti-V _{⊀ll} (R 538)	Anti-V _{⊀⊪} (R 294)	Anti-V _{⊀Ⅳ} (R 178)	Proteins	Anti-λ (R 358)	Anti-V _{al} (R 505)	Anti-V _{או/v} (R 583)	Anti-V _{all} (R 269)	Anti-V _{alv} (R 510)	Anti-V _{avi} (R 394)
κl	+	+	0	0	0	λΙ	+	+	0	0	0	0
ĸII	+	Ó	+	Ō	Ō	λII/V	+	0	+	0	0	0
ĸIII	+	Ó	0	+	0	λlĺ	+	0	0	+	0	0
κIV	+	0	0	0	+	λΙν	+	0	0	0	+	0
						λνι	+	0	0	0	0	+

Table 1. Reactivity by ELISA of Polyclonal Anti-human V_{\star} and V_{λ} Subgroup-specific AntiseraUsed for Immunophenotyping Analyses*

* Reactivity of proteins designated as + or 0 based on at least a >2.5-fold difference in optical density at 405 nm. † Antiserum designation.

Patients

The diagnosis of CLL was made on the basis of established hematologic criteria and that of low-, mid-, or highgrade malignant lymphoma on the basis of morphologic criteria according to the Working Formulation. With rare exception, specimens for analyses were obtained before chemotherapy or radiotherapy was begun.

The B-cell nature of the cell populations obtained from 40 patients with CLL and other low-grade malignant lymphomas and 13 patients with mid- or high-grade lymphomas was evidenced by their CD22⁺, CD3⁻ reactivity and by the presence of slgM or, less commonly, slgM and slgD. CD5 expression was limited to the CLL/low-grade populations, in which 37 of the 40 cases were found to be positive.

Results

Specificity of Anti-light Chain Antisera

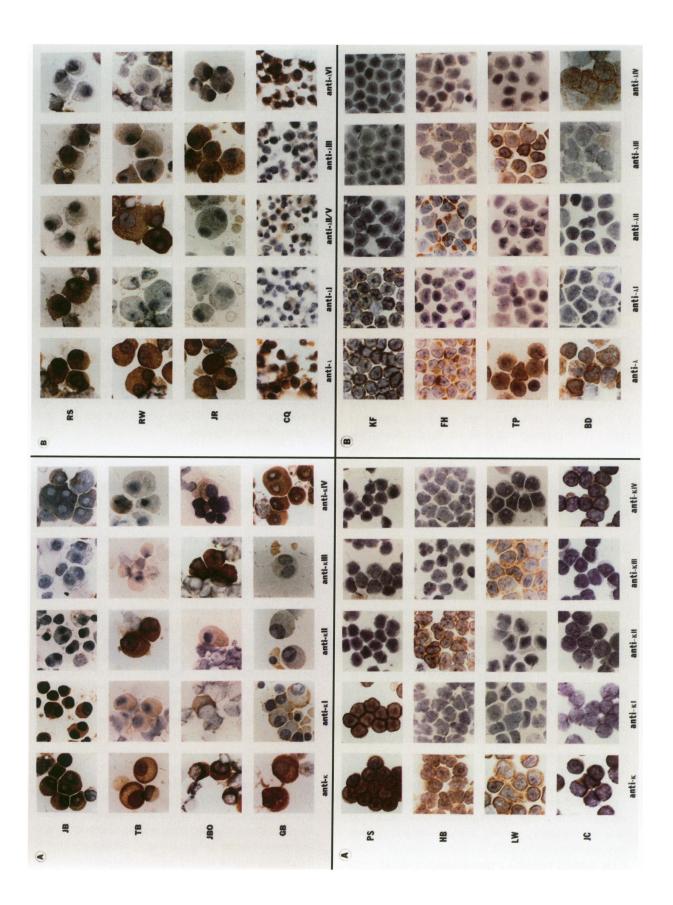
The anti-light chain antisera used as primary antibodies for the immunocytochemical analyses of cellular Ig included two sets of reagents. One, designated 'anti- κ ' or 'anti- λ ,' recognized, respectively, antigenic determinants common to all κ or λ light chains. The second, designated 'anti- V_{κ} ' or 'anti- V_{λ} ,' recognized V region determinants unique to each of the known human V_{L} subgroups, ie, anti- V_{L} subgroup-specific antisera.

The specificity of these anti- κ /anti- λ and anti-V_L subgroup antisera was unambiguously demonstrated by ELISA. For these analyses, the anti-light chain antisera were tested against a panel of monoclonal κ - and λ -type light chains as well as intact IgG, IgA, and IgM proteins of known V_L subgroups, as established by amino acid sequence.^{1,4} As indicated in Table 1, the anti- κ antiserum recognized all κ -type Igs, regardless of V_{κ} subgroup, whereas the anti-V_{κ II}, anti-V_{κ III}, anti-V_{κ III}, and anti-V_{κ IV} antisera were specific for κ I, κ III, κ III, and κ IV type Igs, respectively. Similarly the anti- λ antiserum reacted with all λ -type Igs, whereas the anti-V_{λ II}, anti-V_{λ III}, anti-V_{λ III}, anti-V_{λ IV}, and anti-V_{XVI} antisera recognized only λ I, λ II/V, λ III, λ IV, and λ VI type Igs, respectively. Further confirmation of the specificity of these reagents resulted from immunoblotting experiments in which monoclonal Igs of known V_x or V_x subgroup were subjected to SDS polyacrylamide gel electrophoresis before subtyping analyses. Immunocytochemically the specificity of the anti-V_x and anti-V_x antisera also was apparent in blocking experiments in which cytospin preparations were overlaid before antisera application with monoclonal light chains representative of each V_L subgroup. The reactivity of a particular anti-V subgroup-specific antiserum was abrogated by subgroupconcordant monoclonal light chains but not by light chains of other subgroups.

Immunocytochemical Determination of V_L Subgroups

Multiple Myeloma, Amyloidosis AL, and Waldenström's Macroglobulinemia

Our initial efforts were directed toward the determination of the V, or V_{λ} subgroup of Ig molecules present in the cytoplasm of monoclonal plasma cell populations. For this purpose, we used plasma cell-enriched cytospin specimens isolated from the bone marrow or peripheral blood of patients with multiple myeloma who had serum or urinary monoclonal lgs. Sufficient numbers of slides were prepared to permit the application of normal (nonimmune) rabbit serum and, depending on the κ - or λ -type of the cytoplasmic lg (as determined with the general anti- κ and anti- λ chain antisera), of the 4 anti-V, or 5 anti-V, subgroup-specific antisera. For comparative purposes and to eliminate nonspecific background staining, the slides were incubated with a minimum of three serial dilutions of each antiserum. Similar methods were used to examine, after deparaffinization and enzyme treatment, formalinfixed, paraffin-embedded bone marrow specimens. As illustrated in Figures 1A and B, the V₁ subgroup nature of the clg was evident among the κ - and λ -type monoclonal plasma cells examined (no cases of the rare V_{AV} sub-



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Figure 1. Immunocytochemical characterization of the V_x and V_λ subgroups expressed by the light chains of cytoplasmic Ig (cIg) in monoclonal plasma cell populations. Bone marrow-derived plasma cells from patients with multiple myeloma (cytospin preparations) or amyloidosis AL (deparaffinized, enzyme-treated sections) examined using the ABC immunoperoxidase technique and the anti-light chain antisera as indicated; hematoxylin counterstain; × 1000. A (Top left): Kappa. Reactivity with anti-k and anti- $V_{x,u}$, $V_{x,uv}$, and $V_{x,v}$ antisera indicated the k1, k11, k111, or k1V nature of the cIg in cells from multiple myeloma patients JB, TB, JBO, and GB, respectively. B (Top right): Lambda. Reactivity with anti- λ and anti- $V_{x,v}$, $V_{\lambda uv}$, $v_{\lambda uv}$, antisera indicated the k1, k1, k111, or k1V nature of the cIg in cells from multiple myeloma patients RS, RW, and JR and amyloidosis AL patient CQ, respectively (all patients' cells were tested with the anti- $V_{x,v}$ and found to be negative). Figure 2. Immunocytochemical characterization of the V_x and V_x subgroup expressed by the light chains of surface Ig (sIg) in chronic lymphocytic leukemia. Cytospin preparations of peripheral blood-derived lymphocytes examined using the ABC immunoperoxidase technique and the anti-light chain antisera as indicated the k1, k11, k110. A (BOTOM left): Kappa. Reactivity with anti-x and anti- $V_{x,v}$, $V_{x,u}$, and $V_{x,v}$ antisera indicated the k1, k11, $V_{x,v}$, $V_{x,v}$,

group⁵ were available in this population for study). In studies of cells obtained from 20 patients with multiple myeloma, plasma cell leukemia, or amyloidosis AL, there was complete concordance between the immunocytochemical V_x or V_λ subgroup classification and that obtained through serologic classification of the patient's monoclonal serum lg (lgG or lgA) or urinary Bence Jones protein.

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We also examined bone marrow- and pleural fluid-derived lymphocytoid plasma cells obtained from four patients with Waldenström's macroglobulinemia. In such cases, it was also possible to identify the V_x or V_λ subgroup nature of the clgM. The clg V_L subgroup determined immunocytochemically correlated with that found for the light chains of the secreted IgM protein.

Chronic Lymphocytic Leukemia and Other B-cell-Related Neoplasms

The immunocytochemical methodology used for determination of the V_L subgroup of clg also proved suitable for analyses of slg in other types of monoclonal B-cell populations. Lymphocyte-enriched (more than 90%) preparations were obtained by cytocentrifugation of heparinized peripheral blood or bone marrow specimens from patients with CLL and from cell suspensions prepared from the lymph nodes, tumor masses, or pleural fluid of patients with malignant lymphoma. Using the general anti-k and anti- λ chain antisera, the virtually exclusive slg^{κ +} or $slg\lambda^+$ nature of these cells was shown. Through the application of the anti- V_{\star} or anti- V_{λ} subgroup-specific antisera, it was possible to determine immunocytochemically the particular V_L subgroup expressed by these homogeneous populations of cells (as is illustrated for CLL cells in Figures 2A and B). In contrast, an admixture of $slg\kappa^+$ and slg λ^+ cells was found when samples from two patients with B-cell lymphoproliferative disorders were analyzed with the general anti- κ and anti- λ reagents. In each case the demonstration that the slg κ or slg λ cells expressed, respectively, a single V_{κ} or V_{λ} subgroup permitted the establishment of the monoclonal nature of the disease.

The anti-V_L subgroup-specific antisera also were used successfully to analyze intact sections of lymph node biopsies from patients with B-cell malignancies. In comparative studies involving five patients, the V_L phenotype was as apparent for cells present in fresh-frozen specimens as it was for the isolated cells in cytospin preparations.

Although we have found considerable variation in the intensity of slg staining using the immunoperoxidase technique, we have been able to classify the V_L subgroup expressed by the slg in 53 of 60 patients studied to date. For those patients having a detectable urinary monoclonal Ig, there was agreement between the V_x or V_x subgroup of the slg and that of the excreted lg. For the 40 patients with CLL and other low-grade lymphomas, 28 of these cases were slg κ^+ . Of these, 17 were classified as V_{sl}, 4 as $V_{\kappa ll}$, 4 as $V_{\kappa ll}$, and 3 as $V_{\kappa lV}$. The 12 slg λ^+ cases included 1 classified as $V_{\lambda l}$, 1 as $V_{\lambda ll/V}$, 7 as $V_{\lambda lll}$, and 3 as $V_{\lambda lV}$; no $V_{\lambda VI}^{+}$ cells were found among the slg λ^{+} patients. Among the 13 patients with mid- or high-grade lymphomas, 8 slg κ^+ cases included 3 V_{k1}, 1 V_{k11}, 2 V_{k111}, and 2 V_{k1v}; for the 5 slg λ^+ patients, 3 were classified as V_{$\lambda l}, and 2 as V_{<math>\lambda ll}$ </sub></sub> (Table 2).

The percentage distribution of the V_x and V_λ subgroups for all 53 patients (36 lgx⁺ and 17 lgλ⁺) is given in Table 3 and is compared to the distribution of V_x and V_λ subgroups found on serum lg (based on analyses of more than 200 κ- and λ-type Bence Jones proteins^{2,4}). Among the slgx⁺ cases, the percentage of V_{xIV} was higher and that of V_{xIII} was lower than expected. For the slgλ⁺ patients, a predominance of subgroups V_{λIII} and V_{λIV}, a decrease in V_{λII/V}, and an absence of V_{λVI} were found.

Discussion

Heretofore, the V_L subgroup nature of the lg expressed by monoclonal B-cell populations found in patients with multiple myeloma and other types of malignant lymphoproliferative disorders has been based primarily on analyses of the biologic products of these cells; namely, myeloma proteins, Waldenström macroglobulins, and Bence Jones proteins. The extent of this information has been

	Number	V _* subgroup				V _λ subgroup					
Pathologic Classification*		ĸl	ĸII	ĸIII	кIV	λΙ	λII/V	λIII	λIV	λVI	
Low grade											
slgκ	28	17	4	4	3						
slgλ	12					1	1	7	3	0	
Mid/high grade											
slgĸ	8	3	1	2	2						
slgλ	5					3	0	2	0	0	
Total	53	20	5	6	5	4	1	9	3	0	

Table 2. V_{κ} and V_{λ} Immunophenotypes in Malignant Lymphoma

* International Working Formulation.

limited, however, due to the unavailability of serum or urine specimens or, as in the case of 'nonsecretory' forms of B-cell–related neoplasms, a paucity of circulating or excreted monoclonal lgs. Using a sensitive immunoperoxidase method and our highly specific anti-light chain antisera, we have demonstrated the feasibility of determining immunocytochemically the V_L subgroup expressed by slg as well as by clg. The methodology used has been effective for analyses of cytospin preparations of bone marrow, peripheral blood, or lymph node-derived plasma cells and lymphocytes, as well as of fresh-frozen or formalin-fixed, paraffin-embedded tissue. We also found this technique useful for discriminating between monoclonal and polyclonal B-cell–related disorders.

For patients with multiple myeloma, amyloidosis AL, or Waldenström's macroglobulinemia, the V_L subgroup nature of clg expressed by monoclonal plasma cells or lymphoplasmacytoid cells was readily determined, even in those patients with amyloidosis who typically have minimal bone marrow plasmacytosis.¹⁹ In all cases studied, there was complete concordance between the clg V_L subgroup and that of the serum or urinary monoclonal Ig. Other than the demonstrated association of the V_{XVI} subgroup with amyloidosis AL^{5.14} and the predominance of the V_{XII/V} subgroup among λ -type Waldenström macroglobulins,⁵ an association between V_L subgroups and particular clinicopathologic features of disease remains to be established.

The V_L subgroup of slg expressed by monoclonal Bcell populations in patients with CLL and other types of Bcell malignancies could also be determined immunocytochemically. The distribution of slg V_r or slg V_λ subgroups in this patient population, however, differed from that found from analyses of the monoclonal serum and urinary Ig products of patients with multiple myeloma, amyloidosis AL, and Waldenström's macroglobulinemia.2,4,5 Our data showed that among 40 patients with CLL and other low-grade lymphomas and 13 with mid- or highgrade disease, certain V_L subgroups predominated while others were decreased or absent. For $slg\kappa^+$ cells, the percentage of V_{xIV} was higher and that of V_{xIII} was lower than expected based on the percentage distribution of these V_s subgroups among monoclonal lgs. Even more striking was the unusual V₁ subgroup distribution among slg λ^+ cells in which a predominance of the $V_{\lambda III}$ and (rare) $V_{\lambda IV}$ subgroups was found. We attribute the absence of V_{AVI} cells to the apparent exclusive association of this subgroup with amyloidosis AL.5,14

Our immunocytochemical data thus provide evidence for the predominance of certain V_L subgroups in B-cell neoplasms. One such subgroup, V_{klV}, is of interest because of the preferential association of klV light chains with certain types of cold agglutinins, anti-intermediate filaments, and other types of autoantibodies.² The significance of the V_{λlH} and V_{λlV} subgroup dominance among the slg λ^+ population studied is presently unknown, although it should be noted that a recently characterized monoclonal lgM λ rheumatoid factor also contained λ IIItype light chains.²⁰ Additional studies of monoclonal slg κ^+ and slg λ^+ populations will be required to substantiate the apparent over-representation and clinical relevance of particular slg V₁ subgroups in B-cell disorders.

Cells from virtually all our patients with CLL and lowgrade lymphomas expressed, as expected,²¹ the CD5

Table 3. Comparison of the V_L Subgroup Distribution (%) Found on Serum Igk and Ig λ with that Found on the Cells of 36 Patients with $sIgk^+$ and 17 Patients with $sIg\lambda^+$ B-cell–Related Malignancies

	V _* subgroup					V _λ subgroup					
Source	κl	ĸll	ĸIII	κIV	Source	λΙ	λII/V	λIII	λIV	λΙV λVΙ	
Serum (lg _k) %*	56	10	30	4	Serum (Igλ) %*	26	38	22	3	11	
Lymphoma (lgr) %	55	14	17	14	Lymphoma (Igλ) %	23	6	53	18	0	

* Based on serologic analyses of >200 κ and λ Bence Jones proteins.^{2,4}

marker. This pan T-cell antigen is infrequently found on a relatively small population of normal tonsillar or spleenderived B cells²² but is commonly detected on low-grade neoplastic B-cell populations^{23,24} as well as on auto-reactive lymphocytes.^{25–28} Through studies of CD5⁺, slg_K CLL cells and IgMk autoantibodies, conservation in light (and heavy) chain V gene usage has been demonstrated for certain forms of malignant and reactive B-cell populations.^{13,15–18,29,30} Remarkably the κ chains found on the slg of malignant B cells in \sim 25% of patients with CD5⁺, slg κ CLL (and other low-grade, small cell lymphomas) and the light chains of IgMr rheumatoid factors are members of the V_{rillb} sub-subgroup, express a cross-reactive idiotype (CRI) designated 17.109, and are products of the Humkv 325 germ-line gene.¹¹⁻¹³ Furthermore the resistance of this gene to somatic mutation has been evidenced by the virtual sequence identity of cDNA from cases of CD5⁺, 17.109⁺ B-cell malignancies and also by the strikingly homologous killb protein sequences of Humkv 325-encoded 17.109⁺ IgM_K autoantibodies.¹⁵⁻¹⁸ In contrast, other morphologic types of B-cell lymphomas (eg, follicular cell) are usually CD5⁻ and rarely express the 17.109 CRI; in such cases, restriction in Ig V gene usage and conservation has not been apparent.¹⁷ Other examples of conserved V_L and V_H genes have been reported in patients with monoclonal B-cell neoplastic and autoimmune diseases.31,32

The demonstration that the V_L (or V_H) genes of monoclonal CD5⁺ B cells are relatively resistant to somatic mutation (in contrast to light and heavy chain genes in malignant plasma cell and other cytomorphologic types of B-cell neoplasms) has given new insight into the pathogenesis of CLL and related lymphocytic malignancies. Our finding that the V_L subgroup distribution differs in multiple myeloma versus lymphocytic malignancies provides further evidence of fundamental biologic, as well as clinical, differences between these diseases. The ability to identify immunocytochemically the VL nature of the Ig expressed by monoclonal B-cell populations offers a rapid and sensitive means to determine V_L gene usage in neoplastic and reactive disorders. The demonstration that particular V_L genes are preferentially associated with specific B-cell-related diseases has pathophysiologic importance, as well as diagnostic and therapeutic implications.

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