Advances in the Diagnosis, Classification, Risk Stratification, and Management of Monoclonal Gammopathy of Undetermined Significance: Implications for Recategorizing Disease Entities in the Presence of Evolving Scientific Evidence

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Monoclonal gammopathy of undetermined significance (MGUS) is a premalignant plasma cell disorder present in more than 3% of the general population aged 50 years and older.^{1,2} MGUS is important to clinicians because it is associated with a 1% per year risk of progression to multiple myeloma (MM) or related malignancy.³ It is also important because it is found incidentally during the work-up of a variety of symptoms and disorders, and has confirmed and reported associations with numerous diseases commonly encountered in clinical practice, such as osteoporosis, peripheral neuropathy, and venous thrombosis.⁴ In addition, because MGUS is easily detected on blood tests and can be monitored noninvasively, it represents a readily accessible model to study the conversion of premalignancy to malignancy.⁵

In the past several years, new concepts and advances have emerged concerning the diagnosis, classification, risk stratification, and management of MGUS. This commentary highlights discoveries that provide valuable insight into the process of malignant transformation, as well as management strategies for dealing with a common premalignancy in which preventive strategies are thwarted by low rates of progression.⁶

CLINICAL DISEASE DEFINITIONS

Since MGUS was first described more than 30 years ago, the definition of the entity has evolved.7 Currently, 3 distinct clinical types of MGUS are identified: non-IgM (IgG or IgA) MGUS, IgM MGUS, and light chain MGUS (Table 1). Each clinical subtype is characterized by unique intermediate stages and progression events. For example, the more advanced premalignant stage of plasma cell proliferation in non-IgM MGUS is termed smoldering multiple myeloma (SMM) and is characterized by a much higher risk of progression to MM: 10% per year risk of progression for SMM vs 1% per year risk collectively for all forms of MGUS.⁸ The IgM type of MGUS is associated with a predisposition mainly to Waldenström macroglobulinemia and infrequently to IgM MM.9,10 Recently, a new disease entity termed light chain MGUS has been defined; it represents the premalignant precursor of a subtype of MM called light chain MM that accounts for nearly 20% of all new cases of MM.¹¹ The equivalent of SMM and smoldering Waldenström macroglobulinemia in the spectrum of light chain monoclonal gammopathies is called *idiopathic Bence Jones proteinuria* (Table 1).^{12,13}

What insight do these definitions offer in terms of the process to be used in establishing disease definitions? First, in each of these disease categories, one or more large cohorts of patients meeting a specified disease definition were assembled. Next, the natural history of each disease cohort was determined using sound epidemiological methods.^{3,8,9,11,12,14-16} As a result, we clearly know how to diagnose each of these entities accurately, and we also know the outcome of patients meeting the specific disease definition to assist with management and counseling. This approach to developing disease definitions is preferable to arbitrary criteria, in which the characteristics and outcome of patients meeting such criteria are unknown.

The specific criteria listed in Table 1 are of major importance in patient care and are based on the epidemiological and clinical studies that used clear criteria to define each entity.^{3,8,9,11,12,14-16} As a result of these large studies, we now know the prevalence, risk of progression, and natural history of non-IgM MGUS, IgM MGUS, and light chain MGUS. These studies illustrate that, although these disorders represent clonal proliferation of plasma cells, they do not behave like malignancies, and patients with these diagnoses should be reassured rather than being labeled as having a cancer. For example, patients with less than 10% infiltration of the marrow by lymphoplasmacytic cells have an overall survival that is as good as the general population at large, and should therefore not be labeled as having a lymphoma or Waldenström macroglobulinemia merely because the bone marrow pathology shows clonal proliferation of lymphoid cells.15 As our diagnostic methods improve and become increasingly sensitive, the line between

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Type of monoclonal	Premalignancy with low risk of	Premalignancy with high risk of		
gammopathy	progression (1%-2% per year)	progression (10% per year)	Malignancy	
IgG and IgA (non-IgM) ^b	 Non-IgM MGUS All 3 criteria must be met: Serum monoclonal protein 3 g/dL Clonal BM plasma cells <10% Absence of end-organ damage such as CRAB that can be attributed to the PCPD 	 Smoldering MM Both criteria must be met: Serum monoclonal protein (IgG or IgA) ≥3 g/dL and/or clonal BM plasma cells ≥10% Absence of end-organ damage such as CRAB that can be attributed to a PCPD 	 MM All 3 criteria must be met except as noted: Clonal BM plasma cells ≥10% Presence of serum and/or urinary monoclonal protein (except in patients with true nonsecretory MM) Evidence of end-organ damage that can be attributed to the underlying PCPD, specifically, Hypercalcemia: serum calcium ≥11.5 mg/dL or Renal insufficiency: serum creatinine >2 mg/dL or estimated creatinine clearance <40 mL/min Anemia: normochromic, normo- cytic with hemoglobin >2 g/dL below lower limit of normal or hemoglobin <10 g/dL Bone lesions: lytic lesions or severe osteopenia attributed to a PCPD or pathologic fractures 	
IgM	IgM MGUS ^c All 3 criteria must be met: • Serum monoclonal protein <3 g/dL • Clonal BM lympho- plasmacytic cells <10% • Absence of end-organ damage such as anemia, constitutional symptoms, hyperviscosity, lymphadenopathy, or hepatosplenomegaly that can be attributed to the underlying lympho- proliferative disorder	 Smoldering WM Both criteria must be met: Serum IgM monoclonal protein ≥3 g/dL and/or BM lymphoplasmacytic infiltration ≥10% No evidence of anemia, constitutional symptoms, hyperviscosity, lymphadenopathy, or hepatosplenomegaly that can be attributed to the underlying lymphoproliferative disorder 	 WM All criteria must be met: IgM monoclonal gammopathy (regardless of size of M protein) ≥10% BM lymphoplasmacytic infiltration (usual intertrabecular)) by small lymphocytes that exhibit plasmacytoid or plasma cell differentiation and a typical immunophenotype (eg, surface IgM*, CD5⁺, CD10⁻, CD19⁺, CD20⁺, CD23⁻) that satisfactorily excludes other lymphoproliferative disorders including chronic lymphocytic leukemia and mantle cell lymphoma Evidence of anemia, constitutional symptoms, hyperviscosity, lymphadenopathy, or hepato- splenomegaly that can be attributed to the underlying lymphopro- liferative disorder 	IgM myeloma All criteria must be met: • Symptomatic monoclonal PCPD charac- terized by a serum IgM monoclonal protein regardless of size • Presence of ≥10% plasma cells on BM biopsy • Presence of lytic bone lesions related to the underlying plasma cell disorder and/or translocation t(11;14) on FISH
Light chain	Light chain MGUS All criteria must be met: • Abnormal FLC ratio (<0.26 or >1.65) • Increased level of the appropriate involved light chain (increased K FLC in patients with ratio >1.65 and increased λ FLC in patients with ratio <0.26) • No immunoglobulin heavy chain expression on immunofixation • Clonal BM plasma cells <10% • Absence of end-organ damage such as CRAB that can be attributed to the PCPD	 Idiopathic Bence Jones proteinuria All criteria must be met: Urinary monoclonal protein on urine protein electrophoresis ≥500 mg/24 h and/or clonal BM plasma cells ≥10% No immunoglobulin heavy chain expression on immunofixation Absence of end-organ damage such as CRAB that can be attributed to the PCPD 	Light chain MM ^c • Same as MM except no evidence of immunoglobulin heavy chain expression on immunofixation	rish

TABLE 1. Disease Definitions for the Monoclonal Gammopathies: MGUS and Related Disorders^a

^a BM = bone marrow; CRAB = hypercalcemia, *r*enal insufficiency, *a*nemia, and *b*one lesions; FISH = fluorescent in situ hybridization; FLC = free light chain; MGUS = monoclonal gammopathy of undetermined significance; MM = multiple myeloma; PCPD = plasma cell proliferative disorder; WM = Waldenström macroglobulinemia.

^bOccasionally, patients with IgD and IgE monoclonal gammopathies have been described and will be considered part of this category as well.

^c Note that conventionally IgM MGUS is considered a subtype of MGUS, and similarly light chain MM is considered a subtype of MM. Unless specifically distinguished, when the terms MGUS and MM are used in general, they include IgM MGUS and light chain MM, respectively.

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malignancy and premalignancy will continue to blur. As our understanding of disease progression improves, it will become increasingly important to recognize that welldesigned epidemiological studies and clinicopathologic disease definitions will be required to separate patients who need treatment such as chemotherapy or stem cell transplant for cancer like myeloma¹⁷ from those who need no therapy and need reassurance.⁵

PATHOGENESIS AND CYTOGENETIC CLASSIFICATION

Race and ethnicity play a role in the pathogenesis of MGUS. African Americans, and blacks from Africa, have a 2- to 3-fold higher incidence of MGUS compared with whites.^{18,19} In contrast, the risk is lower in Asians from Japan²⁰ and in Mexicans.²¹ Advancing age,¹ male sex, family history,²² immunosuppression, and exposure to certain pesticides23 all increase the risk of MGUS. Understanding the mechanisms that underlie these risk factors will probably provide clues to the etiology of MGUS. The first step in the pathogenesis is likely an abnormal response to antigenic stimulation, mediated possibly by aberrant expression of toll-like receptors and overexpression of interleukin (IL) 6 receptors and IL-1 $\beta^{24,25}$ This then results in the development of primary cytogenetic abnormalities, either hyperdiploidy or immunoglobulin heavy chain (IgH) translocations (Table 2). The progression of MGUS to myeloma is likely secondary to a random second hit, the nature of which is unknown. Ras and p53 mutations, p16 methylation, myc abnormalities, and induction of angiogenesis are all associated with progression. In addition, there is increased osteoblast RANKL (receptor activator of nuclear factor kB ligand) expression and reduction in the level of its decoy receptor, osteoprotegerin, which results in osteoclast activation and increased bone resorption and turnover.²⁶ This is accompanied by increased levels of IL-3, IL-7, and dickkopf 1 that simultaneously inhibit osteoblast differentiation, leading to the characteristic pure lytic lesions typical of myeloma.27-29

In approximately 50% of MGUS cases, the primary pathogenetic event is likely hyperdiploidy, and in the remaining 50% the pathogenetic event is a translocation event at the IgH locus on chromosome 14q32.³⁰ These pathogenetic events result in at least 6 different cytogenetic subtypes of MGUS and myeloma (Table 2).^{31,32} The excess risk of MGUS in blacks is likely due to a higher predisposition to hyperdiploid MGUS on the basis of recent studies that suggest that the outcome of myeloma in blacks with lenalidomide-based therapy is better compared with that in whites.³³ These observations are important because they highlight the increasing complexity that has accompanied many of the advances

TABLE 2. Cytogenetic Classification of Monoclonal Gammopathy of Undetermined Significance (MGUS)

Cytogenetic abnormality	Affected genes
IgH translocated MGUS (50%)	
t(11;14)(q13;q32)	CCND1 (cyclin D1)
t(4;14)(p16;q32)	FGFR-3 and MMSET
t(14;16)(q32;q23)	C-MAF
t(6;14)(p21;q32)	CCND3 (cyclin D3)
t(14;20)(q32;q11)	MAFB
IgH non-translocated MGUS (50%)	
Hyperdiploid MGUS	Numerous

in cancer research. As more detailed genetic analysis is performed, it becomes apparent that what phenotypically was considered a single malignancy is actually several cytogenetically different diseases, each of which may have a different pathogenetic mechanism, and a different natural history and response to therapy.⁵

RISK STRATIFICATION

An abnormal serum free light chain ratio (ie, the ratio of free immunoglobulin κ to λ light chains in the serum), non-IgG MGUS, and a high serum M protein level (≥ 1.5 g/ dL) are 3 major risk factors for the progression of MGUS to myeloma (Table 3).³⁴ The risk-stratification model is helpful for patient counseling and management. The key concept applicable to other premalignancies is that, as the sensitivity of diagnostic tests increases, it is critical to develop risk-stratification models to distinguish patients with premalignancy of clinical relevance from those whose abnormalities do not result in a sufficiently higher risk compared with that in healthy people.

MANAGEMENT

We recently demonstrated the major problem with managing asymptomatic premalignant disorders like MGUS, which have a low but definite risk of progression to malignancy.³⁵ Preventive therapy cannot be justified without safe treatment options and evidence of benefit from phase 3 clinical trials. On the other hand, close follow-up without treatment can seldom identify progression before serious complications can occur. In these circumstances, risk stratification is needed to help guide optimal management and identification of biomarkers that signal malignant transformation before the onset of serious symptoms. Risk-stratification studies of MGUS indicate that follow-up is unnecessary for low-risk patients, whereas follow-up strategies, prevention trials, and continued research on biomarkers such as malignant immunophenotype and plasma cell proliferative rate for early detection of malignant transformation are needed for high-risk patients.35

Risk group	No. of patients	Relative risk	Absolute risk of progression at 20 y (%)	Absolute risk of progression at 20 y, accounting for death as a competing risk (%)			
Low-risk (serum M protein, <1.5 g/dL; IgG subtype, normal; free light chain ratio, 0.26-1.65	449	1	5	2			
Low-intermediate-risk Any 1 factor abnormal	420	5.4	21	10			
High-intermediate-risk Any 2 factors abnormal	226	10.1	37	18			
High-risk All 3 factors abnormal	53	20.8	58	27			

TABLE 3. Risk-Stratification Model to Predict Progression of Monoclonal Gammopathy of Undetermined Significance to Myeloma or Related Disorders

From Blood,³⁴ the American Society of Hematology.

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