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Elevated serum free light chains are associated with inferior event free and overall survival in Hodgkin lymphoma

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Abstract

The serum free light chain (FLC) assay quantitates free immunoglobulin kappa and lambda light chains, which has prognostic value in plasma cell dyscrasias. However, there is limited data on serum FLC in lymphoid malignancies. We analyzed the association of pretreatment FLC with event-free survival (EFS) and overall survival (OS) in 100 patients with Hodgkin lymphoma (HL). Elevated polyclonal FLC were present in 30% of patients; these patients had an inferior EFS (HR = 4.84; 95% CI: 1.84–12.7) and OS (HR = 8.87; 95% CI: 2.35–33.52) compared to patients with normal FLC. Further studies of FLC in HL are warranted.

Introduction

The serum free light chain (FLC) assay measures the concentration of immunoglobulin (Ig) kappa (κ), and lambda (λ) light chains [1], which are produced by monoclonal and/ or polyclonal B-cell populations. The quantitation of serum FLC and the FLC κ/λ ratio has prognostic value in monoclonal gammopathy of undetermined significance, multiple myeloma, solitary plasmacytoma, and AL-amyloidosis [2–7].

Non-Hodgkin lymphoma (NHL) and Hodgkin Lymphoma (HL) are malignancies of B-cells that can be associated with a monoclonal serum Ig protein. In addition, these lymphoid malignancies often have a polyclonal B-cell infiltrate that can secrete Ig. To date, there have been limited studies that have examined the incidence and prognostic value of serum FLC in lymphoma. In 2005, in a study of 226 patients from our Lymphoma SPORE Serum Bank with NHL and chronic lymphocytic leukemia, we demonstrated that abnormal FLC ratios could be detected more frequently than immunofixation abnormalities; HL patients were not studied [8]. More recently, we showed that over 30% of diffuse large B-cell lymphoma (DLBCL) patients present with elevated FLC, and these patients are at increased risk of an event or death [9]. Others have reported that elevated FLC were associated with event free survival (EFS) in patients with early stage HL [10]. Here we report the association of

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elevated polyclonal FLC concentrations with EFS and overall survival (OS) in a prospectively enrolled cohort of patients with HL.

Methods

This study was reviewed and approved by the Human Subjects Institutional Review Board at the Mayo Clinic and University of Iowa, and written informed consent was obtained from all participants. Details on the patient cohort and FLC quantitation have been previously reported [9]. Briefly, lymphoma patients evaluated at the Mayo Clinic or University of Iowa within 9 months of diagnosis were prospectively enrolled in the University of Iowa/Mayo Clinic SPORE Molecular Epidemiology Resource from September 2002 to February 2008. Of the 219 HL patients enrolled, analyses were limited to the 100 patients with pretreatment research serum samples due to reduction in serum FLC observed in DLBCL patients during treatment [9]. Results for DLBCL are reported elsewhere [11]; here we report the results from HL.

The Freelite assay (The Binding Site, Birmingham, UK) was used to determine serum kappa and lambda FLC concentrations and the $\kappa:\lambda$ FLC ratio from the onstudy research serum samples [1,12]. All assays were performed at the Mayo Clinic Clinical Immunology Lab using kits provided courtesy of The Binding Site. Mayo Clinic reference ranges were used to define the normal range for kappa, lambda, and FLC ratio (0.26–1.65) [12]. Elevated FLC was defined as a kappa or lambda above the published reference range (κ >1.94 mg/dL, λ >2.63 mg/dL).

EFS was defined as the time from diagnosis to the first disease progression, retreatment, or death due to any cause. OS was defined as the time from diagnosis to death due to any cause. Patients without an event or death were censored at time of last known follow-up. Chi-squared tests and Wilcoxon rank-sum tests were used to examine the association of FLC with clinical, prognostic, and demographic variables. Cox proportional hazards models and Kaplan-Meier curves were used to assess associations of FLC abnormalities with time-to-event outcomes. Cox models were performed unadjusted as well as adjusted for the International Prognostic Score (IPS) (Figure 1) [13].

Results

100 HL cases with pretreatment research serum were analyzed. Patient characteristics are summarized in Table I. Thirty patients (30%) had an elevated serum FLC and in all 30 cases the elevation was polyclonal since the $\kappa:\lambda$ FLC ratios were within normal limits. The percentage of patients with elevated FLC was lower in early stage disease compared to advanced stage disease (17 vs. 45%, respectively, P = 0.003). GFR was slightly lower in patients with elevated FLC (median 105, range 78–200) compared to patients with normal FLC (median 125, range 46–322), though not statistically (P = 0.08) or clinically significant. Five additional patients had an abnormal FLC ratio without elevation of kappa or lambda.

At a median follow-up of 48 months (range, 27–87), 18 patients (18%) had an event and 12 patients (12%) had died. The cause of death was due to HL in 9 patients; pulmonary embolism, pneumonia, and aortic stenosis accounted for the other deaths. FLC elevation was significantly associated with age >45, advanced stage, poor ECOG PS, albumin <4.0 g/dl, hemoglobin <10.5 g/dl, leukocytosis (WBC > 15,000), B symptoms, elevated sedimentation rate, and unfavorable IPS. There was no association of FLC elevation with lymphopenia (WBC <600 g/dl), bulky disease, elevated creatinine, or histological subtype (lymphocyte-predominant HL (LPHL) vs. classical HL). Elevated FLCs were associated with poor EFS (HR = 4.58, 95% CI (1.80–11.69, P= 0.001) and OS (HR = 8.99, 95% CI (2.38–33.94, P=

0.001). The association remained significant for EFS after adjusting for the IPS (HR = 3.02, 1.02–8.89, P = 0.05); there were insufficient deaths to build a multivariable model for OS. After adjusting for GFR, elevated FLC remained associated with EFS: HR = 4.20, (95% CI: 1.57–11.23), P = 0.004. Elevated FLC were associated with poor EFS in both early (HR = 6.94, 1.32–36.4, P = 0.02) and advanced stage patients (HR = 3.07, 0.92–10.27, P = 0.07); FLC hazard ratio estimates were also similar across GHSG categories (data not shown). An abnormal FLC ratio without elevated FLC was not associated with outcome (data not shown).

Discussion

Classical HL is a unique tumor in that the majority of cells in the microenvironment are nonmalignant, consisting of reactive T- and B-cells, eosinophils, neutrophils, macrophages, and plasma cells [14]. The malignant cells—the Reed-Sternberg cells—have been demonstrated to be incapable of secreting functional Ig molecules and therefore would not be expected to produce FLC [15,16]. In our study, the FLC elevation is polyclonal in nature (elevated kappa and lambda light chains with a normal ratio) which supports the hypothesis that the FLC is being secreted by polyclonal B-cells, presumably in the HL microenvironment, although this needs to be established. It is known that polyclonal increase in FLC can be due to a number of host factors, such as increased age, impaired renal function, and general inflammation or immune stimulation [17]. In our study, we observed an association of polyclonal elevated FLC with more general host attributes (older age, B symptoms, and poor PS) and with sedimentation rate, an inflammatory marker. It is hypothesized that patients with elevated FLC will have elevations in inflammatory cytokines and chemokines. This should be evaluated in future studies to better understand the pathogenesis.

Serum FLC has the potential to be a useful clinical tool in the care of patients with HL. FLC quantitative analysis could be incorporated into a prognostic index used at diagnosis, which may be useful in predicting those patients who would benefit from a more aggressive first line of treatment. Future studies in HL should evaluate FLC at diagnosis and at response assessment time points to determine the impact on treatment on FLC values and if FLC correlates with clinical response. In summary, serum FLC is an intriguing biomarker in HL, and longitudinal studies are needed to describe the impact of treatment on FLC value, to understand the mechanisms of elevated FLC in HL, and validate these findings.

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Figure 1.

Kaplan Meier curves for event-free (A) and overall survival (B). Hazard ratios are unadjusted Cox models and Cox model adjusted for International Prognostic Score. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]

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TABLE I

Clinical Characteristics by FLC

	All Pa $N = N$	tients 100	Nori FLC A	mal / = 70	Elev FLC /	ated V = 30	
	N	%	N	%	N	%	<i>P</i> -value
Age, median (range)	40 (13	3-81)	36 (18	3-81)	54 (2	4-80)	
Male	54	54%	37	53%	17	57%	0.73
Age >45	39	39%	21	30%	18	60%	0.005
Stage							
Ι	10	10%	6	13%	1	3%	0.007
Π	43	43%	35	50%	8	27%	
III	26	26%	17	24%	6	30%	
IV	21	21%	6	13%	12	40%	
Albumin <4.0 g/dL	44	46%	22	33%	22	76%	0.001
Hemoglobin <10.5 g/dL	10	10%	4	6%	9	21%	0.02
ALC <600	52	53%	37	53%	15	54%	0.95
WBC >15,000	8	8%	33	4%	5	17%	0.03
B symptoms	40	40%	20	29%	20	67%	0.0004
Bulky disease	15	15%	11	16%	4	14%	0.79
ECOG PS >1	14	14%	9	%6	8	27%	0.02
ESR >ULN	36	51%	21	40%	15	83%	0.002
Creatinine >ULN	33	3%	-	1%	2	7%	0.15
SdI							
0-1	32	32%	27	39%	5	17%	0.0001
2	30	30%	27	39%	ю	10%	
3	18	18%	10	14%	8	27%	
4–7	20	20%	9	8%	14	47%	
Subtype							
Classical	93	93%	63	%06	30	100%	0.07
Lymphocyte predominant	٢	7%	7	10%	0	%0	
Initial treatment							
ABVD	85	85%	63	85%	22	85%	0.10

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	All Pat $N = 1$	ients 100	Nori FLC A	mal $7 = 70$	Elev: FLC A	ited '= 30		
	N	%	N	%	N	%	<i>P</i> -valı	lle
MOPP	2	2%	-	2%	-	2%		1
Stanford V	ŝ	5%	б	5%	2	5%		
Other/missing	8	8%	ю	8%	5	8%		
DSHSG								
Advanced	52	52%	28	54%	24	46%	0.00	07
Early favorable	30	30%	26	87%	4	11%		
Early unfavorable	18	18%	16	89%	2	13%		
TLC mg/dL, median (range)								
Kappa	1.44~(0.0	(0.6-9	1.14(0.0)	5-1.94)	2.55 (1.5	(0-6-0		
Lambda	1.50(0.2)	0-9.1)	1.26 (0.2))- 2.62)	2.58 (1.6	(3-9.1)		
Total FLC	2.90 (0.42	2-15.8)	2.56 (0.4.	2-4.56)	5.29 (3.7	t– 15.8)		

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