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A prospective study of serum microbial translocation biomarkers and risk of AIDS-related non-Hodgkin lymphoma

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Abstract

Chronic immune activation is a harbinger of AIDS-associated non-Hodgkin lymphoma (AIDS-NHL), yet the underlying basis is unclear. Microbial translocation, the passage of microbial components from the gastrointestinal tract into the systemic circulation, is a source of systemic immune activation in HIV infection and may be an important contributor to the chronic B cell activation and subsequent AIDS-NHL development. We measured biomarkers of microbial translocation including bacterial receptors/antibodies, intestinal barrier proteins, and macrophage activation-associated cytokines/chemokines, in serum from 200 HIV-infected men from the Multicenter AIDS Cohort Study prior to their AIDS-NHL diagnosis (mean=3.9 years; SD=1.6 years) and 200 controls. Controls were HIV-infected men who did not develop AIDS-NHL, individually matched to cases on CD4+ T cell count, prior antiretroviral drug use, and recruitment year into the cohort. Biomarkers of bacterial translocation and intestinal permeability were significantly increased prior to AIDS-NHL. Lipopolysaccharide-binding protein (LPB), fatty acid binding protein 2 (FABP2) and soluble CD14 had 1.6-, 2.9-, and 3.7-fold increases in risk for each unit increase on the natural log scale, respectively. Haptoglobin had a 2.1-fold increase and endotoxin-core antibody a 2.0-fold decrease risk for AIDS-NHL (4th versus 1st quartile). Biomarkers of macrophage activation were significantly increased prior to AIDS-NHL: B-cell activation factor (BAFF), IL18, monocyte chemoattractant protein-1 (MCP1), Tumor Necrosis factor- α (TNF α), and CCL17 had 2.2-, 2.0-, 1.6-, 2.8-, and 1.7-fold increases in risk for each unit increase on the natural log scale, respectively. These data provide evidence for microbial translocation as a cause of the systemic immune activation in chronic HIV infection preceding AIDS-NHL development.

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Keywords

microbial translocation; and macrophage activation; AIDS-NHL; HIV

Introduction

Non-Hodgkin lymphoma (NHL) is an important cause of morbidity and mortality in people with HIV infection.^[1–3] While the incidence of AIDS-associated NHL (AIDS-NHL) has decreased with combination antiretroviral therapy (cART), it still results in 23–30% of AIDS-related deaths in countries with readily accessible cART.^[4–7] There are several AIDS-NHL subtypes; all are of B cell origin, yet subtypes are thought to differ in their pathogenesis.^[8–10] Primary central nervous system lymphomas (PCNSLs) are less common, but more aggressive, than Burkitt's lymphoma (BL), diffuse large B cell lymphoma (DLBCL), or other systemic tumors.^[11] Two major mechanisms are believed to contribute to the genesis of AIDS-NHL: i) the loss of immunoregulatory control of Epstein Barr virus (EBV) infected B cells, resulting from impaired T cell function; and ii) chronic B cell activation, which leads to DNA modifying events (oncogene mutations and translocations) that contribute to lymphomagenesis.^[12–14]

Epidemiologic evidence suggests a role for B cell stimulatory cytokines and molecules in the development of AIDS-NHL. Elevated levels of circulating cytokines (IL6, IL10, CXCL13, Tumor Necrosis factor- α [TNF α], Interferon- γ inducible protein [IP-10]/CXCL10), soluble receptors (sCD23, sCD27, sCD30), and markers of inflammation and immune activation (neopterin, κ and λ immunoglobulin free light chains) have been shown to precede the development of AIDS-NHL by up to 7 years.^[11, 15] Examining potential sources of chronic B cell activation in the setting of HIV infection may provide insights into the etiology of AIDS-NHL. Microbial components (e.g., lipopolysaccharide [LPS]) passing across the intestinal epithelial barrier into peripheral circulation (termed microbial translocation) has been shown to contribute to systemic immune activation in chronic HIV infection, and may be a contributor to the chronic B cell activation that predisposes to AIDS-NHL.^[16, 17] Microbial translocation and the consequential systemic immune activation is influenced by the loss of integrity and structural damage to the gastrointestinal barrier in chronic HIV infection, which can be measured by circulating barrier proteins including fatty acid binding protein (FABP) 2 and haptoglobin.^{[18][17]} The presence of LPS in the bloodstream leads to consumption of antibodies against LPS (anti-endotoxin-core protein IgM, otherwise known as EndoCAb), increased production of LPS binding protein (LBP) and receptor (sCD14), and increase in activation of macrophages and the secretion of various cytokines and chemokines.

In this study, we sought to determine the association between these biomarkers of microbial translocation, intestinal integrity, and macrophage activation and subsequent AIDS-NHL development, to support our hypothesis that microbial translocation contributes to chronic B cell activation and AIDS-NHL risk. We also examined several B cell activation markers to confirm prior reports of significant elevations preceding AIDS-NHL diagnosis.

Methods

Study population

This is a nested case-control study utilizing serum specimens from participants in the Multicenter AIDS Cohort Study (MACS), a prospective cohort study of the natural and treated history of HIV/AIDS in men who report sex with men. [19, 20] The participants were recruited from four U.S. metropolitan areas (Baltimore/Washington, Chicago, Los Angeles, and Pittsburgh), and study visits have been held at 6-month intervals beginning in 1983 (www.statepi.jhsph.edu/macs/mac.html). A total of 7,350 men have been enrolled in the MACS.

Participants were selected for study on the basis of pathologically confirmed AIDS-NHL after enrollment into the MACS and archival serum specimens available prior to cancer diagnosis (n=200). For each case, one control was selected randomly from all possible HIV-infected MACS participants who had not developed AIDS-NHL as of November 2014, matched on: (i) CD4+ T cell counts ($\pm 200/\text{ul}$); (ii) prior antiretroviral drug use (ever versus never); and (iii) recruitment year into the cohort (1984–85, 1987–91, or 2001+). Additionally, cases treated with a potent combination of antiretroviral drugs were matched to controls on time since first therapy, and cases who became HIV-infected after recruitment into the cohort were matched to controls by their seroconversion date.

Biomarker determination

Serum levels of haptoglobin and anti endotoxin-core protein IgM were determined by ELISA according to the manufacturer's directions (R&D Systems and Hycult Biotech, respectively). Serum levels of all other biomarkers were determined using the Luminex platform with custom-made panels produced by R&D systems. Briefly, Luminex microparticles pre-coated with analyte-specific antibodies were incubated with diluted serum samples, followed by a biotin-antibody and by a streptavidin-phycoerythrin conjugate. The fluorescence intensity of each analyte's microparticles was quantified using a Bioplex 200 (Luminex) System Analyzer (Bio-Rad), and the data analyzed using BioPlex Manager (v 4.1.1) software. The lower limit of detection (LLD) for each biomarker was set either as the lowest value that the BioPlex Manager software could calculate using the standard curve, or as the lowest value of the standard curve, whichever was smaller. For quality control, case and control samples were equally distributed across reaction plates, and replicates were included across the reaction plates to calculate coefficients of variation (CVs). All laboratory personnel were blinded to the case-control status of samples. Additional assay details including LLD, dilution factors, detectable fractions, and CVs, are presented in Supplemental Table 1.

Statistical analysis

We analyzed our biomarkers as natural log-transformed continuous variables. Additionally, in exploratory analyses, we examined biomarker categories by quartiles according to the distribution among the controls. We calculated odds ratios (ORs) and 95% confidence intervals (CIs) representing the risk of AIDS-NHL associated with one log-unit change in biomarker levels or risk of AIDS-NHL associated with each of the 2nd, 3rd, and 4th

quartiles compared to the 1st quartile, using conditional logistic regression. In addition to the matching factors, ORs were adjusted for covariates selected *a priori* for their association with AIDS-NHL in prior studies including age (continuous), race/ethnicity (categorical), and Hepatitis C virus (HCV) infection status. Each biomarker was tested for association with AIDS-NHL in separate regression models. Additionally, we tested for trends across quartiles using a continuous variable with values representing the medians of each category. We examined patterns of AIDS-NHL risk associated with biomarkers according to subgroups of systemic lymphomas or PCNSL. We also examined patterns of AIDS-NHL risk associated with biomarkers according to the time interval between serum sample collection and AIDS-NHL diagnosis. We classified this lag time into two categories: < 4 years or ≥ 4 years. These categories were selected according to the natural distribution of lag times and wanting to ensure approximately equal number of participants in each category. In addition to these stratified analyses, we also tested for statistical interactions between biomarkers and lagtime using interaction terms in the models. Lastly, we calculated pairwise correlations between all biomarkers using Pearson's correlation coefficient.

Results

Study population description

Cases and controls were similar in their distributions by recruitment year, CD4+ T cell count, and antiretroviral drug therapy, as expected based on the matched design (Table 1). The majority of controls and cases were non-Hispanic white (80.0% and 81.0% respectively). Cases tended to be older than controls, with 44.5% of cases ≥40 years, compared to 38.0% of controls. Cases were more likely to have acute or chronic HCV infection compared to controls (10.2% versus 6.5%). Both groups had relatively high levels of CD4+ T cells, with 51.5% of controls and 46.5% of cases having >400 CD4+ T cells/mm³, and the majority were antiretroviral drug naïve (94.0% of controls and 94.5% of cases). The mean time from serum date to NHL diagnosis was 3.9 years; standard deviation of 1.6 years and a range of 1 month to 12 years. The majority of cases (69.5%) were systemic lymphomas, and DLBCL was the major subtype (48.2%).

Microbial translocation biomarkers

Levels of sCD14, FABP2, and LBP were associated with AIDS-NHL risk, with 3.71-, 1.64-, and 2.97-fold increases in risk for each unit increase on the natural log scale, respectively (Table 2, Figure 1). Additionally, participants with marker levels in the highest versus lowest quartile for haptoglobin, and EndoCAb, were at 2.18-fold increased, and 2-fold decreased, risk of AIDS NHL, respectively. For some markers (haptoglobin, FABP2, and LBP), there appeared to be a threshold effect for the highest quartile: an increased AIDS-NHL risk was only observed for participants with marker levels in the highest versus lowest quartiles. In comparison, for sCD14, AIDS-NHL risk was elevated similarly across the 2nd, 3rd, and 4th quartiles. Also, AIDS-NHL risk was reduced similarly across the 2nd, 3rd, and 4th quartiles for EndoCAb.

In subgroup analyses by lag time between pre-diagnostic serum collection date and AIDS-NHL diagnosis date, EndoCAb, FABP2, and Haptoglobin demonstrated consistent

associations with AIDS-NHL risk across lag times, while other markers were more strongly associated with AIDS-NHL risk when measured close to diagnosis date (< 4 years, sCD14 and LBP, Table 2). There was an association between increased AIDS-NHL risk and higher levels of FABP2, haptoglobin, and LBP, particularly among those men with systemic lymphoma in comparison to those with PCNSL (Figure 1, Supplemental Table 2). Conversely, higher levels of sCD14 were more strongly associated with PCNSL versus systemic lymphoma. EndoCAb appeared to be similarly associated with decreased risk of systemic lymphoma and PCNSL.

Macrophage-associated cytokines and chemokines

BAFF, CCL17, ICAM, IL18, and MCP1 were associated with AIDS-NHL, with 2.21-, 1.68-, 1.66-, 1.93-, and 1.60-fold increases in risk for each unit increase on the natural log scale, respectively (Table 3, Figure 1). When modeled as quartiles, these biomarkers generally displayed dose-response relationships with AIDS-NHL risk (the higher the levels, the higher the risk). For sCD163, participants with levels in the highest versus lowest quartile were at 1.94-fold increased AIDS-NHL risk. CCL17 and MCP1 displayed similar associations with AIDS-NHL risk when measured < 4 years or ≥ 4 years prior to diagnosis date (Table 3). BAFF and IL18, were more strongly associated with AIDS-NHL risk when measured close to diagnosis date (< 4 years), while ICAM and sCD163 displayed stronger associations when measured ≥ 4 years prior to diagnosis. BAFF and IL18 also displayed stronger associations with systemic lymphoma than PCNSL, while CCL17, ICAM, MCP1 (Supplemental Table 2) were similarly associated with systemic and PCNSL risks. TRAIL was unassociated with AIDS-NHL overall, however we observed a statistically significant interaction ($p=0.04$) between TRAIL and lag time: higher TRAIL levels were associated with AIDS-NHL risk only when measured ≥ 4 years prior to AIDS-NHL diagnosis.

B cell activation markers

There was a significant association between B cell activation markers with AIDS-NHL risk, with ORs ranging from 1.23 to 4.91 for each unit increase on the natural log scale (Supplemental Table 3, Figure 1). These associations were observed when markers were measured close to diagnosis (< 4 years), and persisted when markers were measured ≥ 4 years prior to diagnosis. These biomarkers were significantly associated with both systemic lymphoma and PCNSL, although CXCL13 appeared to be more strongly associated with systemic lymphoma while sCD23 was more strongly associated with PCNSL (Supplemental Table 2, Figure 1).

We observed strong correlations between several markers of microbial translocation (sCD14), macrophage activation (sCD163, ICAM, BAFF, and IL18), and B cell activation (IP10 and sCD27), with Pearson's pairwise correlations ranging from 0.3 to 0.6.

Discussion

In this prospective study, we defined the association between serum levels of molecules associated with microbial translocation and immune activation/inflammation with subsequent AIDS-NHL diagnosis. Key novel observations were that markers of bacterial

exposure and host response to bacteria (sCD14, LBP, EndoCAb), intestinal barrier dysfunction (FABP2 and Haptoglobin), and macrophage activation (BAFF, CCL17, ICAM, IL18, MCP1, and sCD163,) were highly predictive of subsequent AIDS-NHL diagnosis in men with HIV infection. Importantly, these associations were observed after careful adjustment of HIV disease status, immune suppression, and antiretroviral drug therapy.

sCD14 displayed the strongest association with AIDS-NHL of all the microbial translocation markers examined. sCD14 is the soluble form of the LPS co-receptor and is produced by macrophages on exposure to LPS and other inflammatory activators.^[21, 22] Furthermore, sCD14 was more strongly associated with AIDS-NHL when measured close to the diagnosis date (< 4 years), and was more strongly associated with PCNSL than systemic lymphoma. PCNSL is an aggressive presentation of AIDS-NHL believed to arise principally due to the oncogenic properties of uncontrolled EBV reactivation in severely immunocompromised individuals. These observations are consistent with the hypothesis that microbial translocation is a consequence of compromised gut immunity (lower CD4+ T cell count), which occurs close to the time of AIDS-NHL diagnosis,^[23] and is more strongly associated with PCNSL in comparison to systemic lymphoma development.^[24] In a prior study, serum levels of sCD14 greater than or equal to 1.76×10^6 pg/ml (median value in controls) were associated with a 2.7-fold increased risk of AIDS-NHL, in samples collected a median of 1 year prior to AIDS-NHL diagnosis.^[25] Our results build on this prior observation in a larger study sample with a longer lag time from specimen collection to AIDS-NHL diagnosis.

LBP was another biomarker found to be strongly associated with subsequent AIDS-NHL diagnosis, particularly when it was measured close to NHL diagnosis date (< 4 years); however, in contrast to sCD14, LBP was more strongly associated with systemic lymphoma when compared to PCNSL. LBP is an LPS binding protein which promotes binding of LPS to its receptor. Additionally, we observed that levels of EndoCAb, a neutralizing antibody against LPS, were significantly reduced prior to AIDS-NHL, with consistent associations across lag times from blood collection to AIDS-NHL diagnosis, and between systemic lymphoma and PCNSL. In prior studies, LBP and EndoCAb have been described as indirect markers of LPS: LBP is positively associated while EndoCAb is inversely associated with LPS levels^[16]. Prior research suggests that HIV-infection impairs EndoCAb response.^[26] Thus, the observed inverse association between EndoCAb levels and AIDS-NHL risk is consistent with the notion that LPS in circulation among HIV-infected persons at risk for NHL depletes circulating EndoCAb, further supporting the hypothesized role of microbial translocation in AIDS-NHL.

Structural damage to the gastrointestinal tract in chronic HIV infection has been well-described,^[18] and the integrity of the intestinal barrier is an important factor in HIV-related systemic immune activation that may also influence lymphomagenesis.^[17] FABP2 (a lipid transport protein and marker of enterocyte damage) and haptoglobin (a physiological modulator of intercellular tight junctions) are markers of intestinal barrier integrity. For both of these markers, a significantly increased risk for AIDS-NHL risk was observed only among those men with the highest levels (4th quartile), particularly for systemic lymphoma. In prior studies, higher levels of FABP2 were seen to be associated with lower CD4+ T cell

counts, liver disease progression, and mortality in HIV infection.^[27–29] To our knowledge, no prior studies have examined serum haptoglobin in HIV infection or lymphoma.

We previously showed that pre-NHL diagnosis serum levels of markers related to B-cell activation (sCD23, sCD27, CXCL13, IL6, IL10, IP10, and TNF α) were associated with risk for AIDS-NHL.^[11, 13, 15, 30] In this study, we found that other cytokines and chemokines that are mainly secreted by activated macrophages (BAFF, CCL17, ICAM, IL18, MCP1, and sCD163) were associated with risk for AIDS-NHL.^[31–35] Macrophages have numerous functions related to inflammation, immunity, and tumor growth, and can be phenotypically polarized by the microenvironment to become M1 or M2. M1 typically secrete pro-inflammatory cytokines (TNF α , IL-6), while M2 mainly promote Th2 responses, immunosuppression, and tumor promotion.^[31, 36] It is interesting to note that many of the cytokines or chemokines associated with AIDS-NHL risk are typically produced by M1 polarized macrophages, including IP10, TNF α , IL18, IL6 and BAFF,^[31, 32, 36] which may be reflecting microbial translocation and activation of macrophages by LPS. Some notable M2 markers, including CCL17 and CD163,^[31, 36] were also associated with AIDS-NHL risk, possibly reflecting the presence of tumor-associated, tumor growth-promoting, macrophages. However additional studies are needed to elucidate these associations.

Interestingly, we found an association between pre-diagnosis serum levels of CCL17 (TARC) and the risk for AIDS-NHL. CCL17 was previously shown to be associated with the development of Hodgkin lymphoma (HL), suggesting that there may be common mechanisms contributing to the development of these lymphatic system cancers as they relate to CCL17.^[37] High levels of sCD163 (4th quartile) were associated with increased AIDS-NHL risk, particularly PCNSL (which tend to be EBV-associated). In prior work, sCD163 correlated with plasma EBV-DNA in patients with HL^[38]. Thus, we hypothesize that CCL17 and sCD163 may be associated with EBV reactivation in AIDS-NHL and HL.

We also found that pre-diagnosis levels of BAFF were associated with risk for AIDS-NHL. BAFF is a member of the TNF family of cytokines and plays an important role in normal B cell proliferation and activation. High levels of soluble BAFF may be contributing to prolonged B cell survival and increased susceptibility to mutation and transformation.^[39] Prior studies have found that BAFF was elevated in patients with NHL and correlated with aggressive disease.^[40]

The question of whether microbial translocation and macrophage activation markers are etiologically associated with AIDS-NHL development is impossible to determine in the context of an observational study. Interestingly, sCD14 was highly correlated with B cell activation markers (sCD27 and IP10), providing a link between microbial translocation and the B cell activation that precedes AIDS-NHL. However, pre-clinical disease in the cases may be responsible for the observed associations. Interestingly, we found that risk estimates for many of our markers were comparable when analyses were stratified by short (< 4 years) or long ($\geq$ years) lag time from specimen collection to AIDS-NHL diagnosis, suggesting that the associations are less likely to be due to undiagnosed disease. Some notable exceptions were sCD14 and LBP, both of which were more strongly associated with AIDS-NHL when

measured closer to NHL diagnosis, suggesting a role of undiagnosed AIDS-NHL in the observed associations.

Systemic immune activation occurs early in the course of HIV infection.^[16, 17] Therefore, it is reasonable to speculate that a failure to control immune activation may contribute to lymphomagenesis. In accordance with this, a recent study showed that plasma levels of LBP, FABP2 and IL6 were significantly elevated in patients from whom antiretroviral drug treatment was introduced during chronic HIV infection. In contrast, patients who started treatment with a potent combination of antiretroviral drugs at the time of HIV diagnosis were able to normalize their levels of LBP, FABP2 and IL6.^[41] This finding supports early introduction of therapy to limit systemic immune activation and thereby reduce the incidence of lymphomagenesis. Even though this is an interesting hypothesis, we were not able to test it in this study, as only a small percentage of cases and controls (6%) were treated with potent combinations of antiretroviral drugs.

The main strength of this study is the inclusion of a large sample of known AIDS-NHL cases with detailed longitudinal covariate data and specimen availability before AIDS-NHL. Compared to prior studies, we examined a larger number of biomarkers indicative of microbial translocation and related immune activation, with longer lag times from biomarker measurement to AIDS-NHL diagnosis. Also, our larger study population allowed us to examine differences in biomarker associations by lag time and site (CNS versus systemic). In addition to the markers presented in this paper, this study population has been characterized with respect to a number of important biomarkers related to AIDS-NHL, permitting us to evaluate our findings in this context. Although we controlled for CD4+ T-cell numbers and date of HIV infection among seroconverters, our observed associations may still be influenced by residual confounding by HIV progression. Also, our population consisted largely of white men who have sex with men, who developed lymphoma prior to initiation of HAART, potentially limiting the generalizability of the findings. We did not correct for type 1 error, thus we cannot exclude the possibility of chance findings. However, our goal of this study was to examine several biomarkers in a related pathway to demonstrate strength of evidence (not necessarily statistical significance for a single biomarker) for a role of microbial translocation in AIDS-NHL, thus type 1 error correction may not be appropriate.

Here we show that systemic immune activation in chronic HIV infection plays a prominent role in the development of AIDS-NHL. Additionally, the cytokine profile seen to precede AIDS-NHL diagnosis is consistent with an M1 macrophage activation, which in part may be driven by endotoxin exposure due to microbial translocation. Future research is needed to better understand their biologic basis. For example, studies are needed to better define how LPS drives chronic B-cell activation and subsequent AIDS-NHL development. If these results are replicated, attenuation of microbial translocation-induced immune activation could be an effective therapeutic strategy to reduce AIDS-NHL risk.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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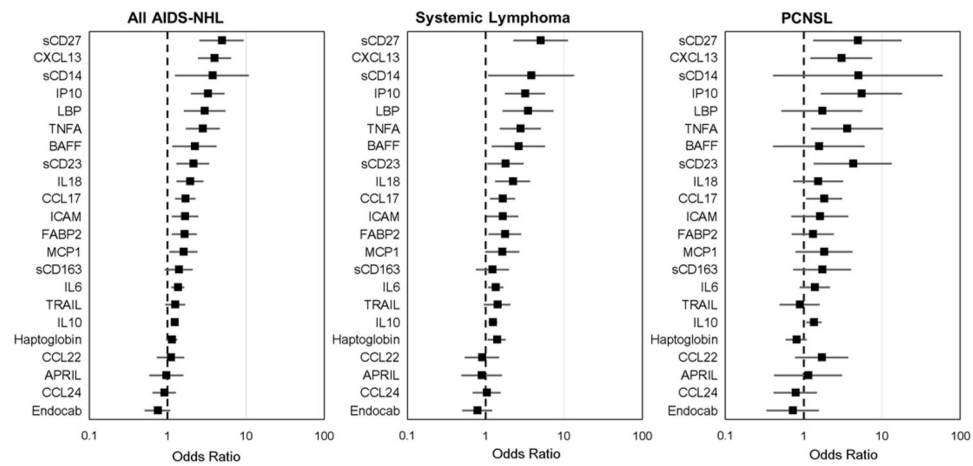


Figure 1. Overall AIDS-NHL, systemic lymphoma, and PCNSL risk associated with 1-unit increase in biomarkers on the natural log scale (OR=box, 95% CI=bar).

Table 1

Select characteristics of AIDS-NHL cases and controls

	HIV-infected controls (n = 200)		AIDS-NHL cases (n = 200)	
	Count	Percentage	Count	Percentage
Recruitment cohort				
1984–1985	169	84.5%	169	84.5%
1987–1991	24	12.0%	24	12.0%
2001+	7	3.5%	7	3.5%
Race				
White, non-Hispanic	160	80.0%	162	81.0%
Black, non-Hispanic	23	11.5%	17	8.5%
Hispanic	16	8.0%	21	10.5%
Asian or Pacific Islander	1	0.5%	0	
Age *				
< 30	31	15.5%	28	14.0%
30 – 39	93	46.5%	83	41.5%
40 – 49	61	30.5%	70	35.0%
≥ 50	15	7.5%	19	9.5%
Body mass index (mean ± SD) *	23.7 ± 2.8		23.9 ± 3.1	
HCV status *				
Negative	180	90.5%	171	86.8%
Acute or chronic infection	13	6.5%	20	10.2%
Cleared	6	3.0%	6	3.0%
Unknown	1		3	
CD4+ T-cells/mm ³ *				
< 200	40	20.0%	44	22.0%
200 – 399	57	28.5%	63	31.5%
≥400	103	51.5%	93	46.5%
Unknown				
Prior HAART exposure *				
No	188	94.0%	189	94.5%
Yes	12	6.0%	11	5.5%
Time from serum date to NHL diagnosis, years (mean ± SD)	N/A		3.9 ± 1.6	
NHL site				
Systemic			139	69.5%
Central Nervous System			61	30.5%
NHL subtype (systemic only)				
Diffuse large B-cell lymphoma			67	48.2%
Burkitt Lymphoma			23	16.5%
Lymphoplasmacytic lymphoma			2	1.4%
Peripheral T-cell lymphoma			2	1.4%

	HIV-infected controls (n = 200)		AIDS-NHL cases (n = 200)	
	Count	Percentage	Count	Percentage
Primary effusion lymphoma			1	0.7%
Follicular lymphoma			1	0.7%
NHL, NOS			43	30.9%
Tumor EBV status				
Negative			28	31.8%
Positive			60	68.2%
Unknown			88	

NHL, non-Hodgkin lymphoma; SD, standard deviation; HCV, hepatitis C virus; HAART, highly active antiretroviral therapy; EBV, Epstein-barr virus

* The reference date for these variables is the blood collection date which was used for testing serum biomarkers

Association between microbial translocation biomarkers and AIDS-NHL risk overall and stratified by time from blood collection to AIDS-NHL diagnosis

Table 2

	<i>All AIDS-NHL</i>									
	< 4 years time lag					≥ 4 years time lag				
	Controls, N	Cases, N	OR	95% CI		Controls, N	Cases, N	OR	95% CI	
sCD14 ng/mL	200	200	3.71	1.25 – 11.00		103	103	14.9	2.41 – 91.4	
< 1307	50	26	1			30	10	1		
1307–1529	50	56	2.03	1.06 – 3.87		26	25	3.51	1.23 – 10.0	
1529–1738	50	62	2.24	1.16 – 4.33		25	38	5.39	1.90 – 15.3	
> 1738	50	56	2.08	1.05 – 4.11		22	30	5.51	1.75 – 17.3	
EndoCAb MMU/mL	200	200	0.75	0.51 – 1.08		103	103	0.77	0.44 – 1.37	
< 39	50	76	1			21	36	1		
39–51	50	34	0.44	0.24 – 0.80		27	15	0.28	0.11 – 0.72	
52–73	50	49	0.55	0.31 – 0.10		32	29	0.50	0.21 – 1.15	
> 73	50	41	0.50	0.28 – 0.90		23	23	0.54	0.22 – 1.32	
FABP2 pg/mL	200	200	1.64	1.13 – 2.38		103	103	1.64	0.98 – 2.77	
< 694	50	37	1			27	20	1		
694–1009	50	49	1.87	0.94 – 3.71		27	29	2.37	0.90 – 6.24	
1010–1448	50	35	1.05	0.53 – 2.07		23	13	0.99	0.36 – 2.67	
> 1448	50	79	2.77	1.42 – 5.39		26	41	2.90	1.11 – 7.57	
Haptoglobin pg/mL	200	199	1.12	0.95 – 1.33		103	102	1.14	0.89 – 1.46	
< 462	50	38	1			21	14	1		
462–657	50	33	0.85	0.43 – 1.70		26	18	0.80	0.26 – 2.41	
657–932	50	55	1.38	0.76 – 2.48		24	27	1.90	0.63 – 5.74	
> 932	50	73	2.18	1.15 – 4.12		32	43	3.24	1.18 – 8.89	
LBP ng/mL	200	200	2.97	1.61 – 5.48		103	103	7.07	2.44 – 20.5	
< 8930	50	27	1			27	9	1		

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	All AIDS-NHL				< 4 years time lag				≥4 years time lag			
	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N	OR	95% CI
8931–11169	50	28	1.04	0.51 – 2.10	22	15	2.28	0.70 – 7.43	28	13	0.51	0.19 – 1.35
11169–13777	50	50	1.72	0.88 – 3.34	28	25	2.47	0.83 – 7.33	22	25	1.41	0.55 – 3.62
> 13777	50	95	3.40	1.79 – 6.45	26	54	7.32	2.43 – 22.1	24	41	2.04	0.84 – 4.94

N, number; OR, odds ratio; CI, confidence interval. ORs were adjusted for age at serum sample date (continuous), race/ethnicity (categorical), and HCV infection status at serum sample date.

Association between macrophage activation biomarkers and AIDS-NHL risk overall and stratified by time from blood collection to AIDS-NHL diagnosis

Table 3

		<i>All AIDS-NHL</i>						<i>< 4 years time lag</i>						<i>≥ 4 years time lag</i>					
		Controls, N		Cases, N	OR	95% CI		Controls, N		Cases, N	OR	95% CI		Controls, N		Cases, N	OR	95% CI	
APRIL pg/mL	200	200	200	200	0.96	0.58 – 1.58		103	103	103	1.15	0.56 – 2.34		97	97	97	0.78	0.37 – 1.63	
< 938	50	50	50	1	1			29	28	28	1			21	22	22	1		
938–1257	50	54	54	1.12	0.64 – 1.97			31	28	28	0.81	0.37 – 1.74		19	26	26	1.89	0.70 – 5.12	
1258–1703	50	50	50	1.02	0.56 – 1.84			24	26	26	1.54	0.66 – 3.58		26	24	24	0.75	0.29 – 1.94	
> 1703	50	46	46	0.91	0.48 – 1.72			19	21	21	1.34	0.52 – 3.44		31	25	25	0.64	0.24 – 1.74	
BAFF pg/mL	200	200	200	2.21	1.15 – 4.24			103	103	103	4.43	1.56 – 12.5		97	97	97	1.23	0.46 – 3.30	
< 1475	50	26	26	1	1			28	14	14	1			22	12	12	1		
1475–1816	50	38	38	1.33	0.69 – 2.58			27	18	18	1.16	0.42 – 3.25		23	20	20	1.51	0.57 – 3.97	
1817–2287	50	65	65	2.35	1.27 – 4.35			25	29	29	2.52	1.00 – 6.34		25	36	36	2.63	1.02 – 6.77	
> 2287	50	71	71	2.88	1.48 – 5.60			23	42	42	4.28	1.59 – 11.5		27	29	29	2.11	0.73 – 6.07	
CCL17 pg/mL	200	200	200	1.68	1.25 – 2.26			103	103	103	1.71	1.11 – 2.64		97	97	97	1.70	1.10 – 2.63	
< 311	50	32	32	1	1			20	18	18	1			30	14	14	1		
311–501	50	48	48	1.82	0.95 – 3.51			25	20	20	0.85	0.33 – 2.22		25	28	28	3.62	1.31 – 10.0	
502–759	50	38	38	1.26	0.66 – 2.38			35	15	15	0.60	0.23 – 1.55		15	23	23	3.86	1.31 – 11.4	
> 759	50	82	82	3.09	1.66 – 5.77			23	50	50	3.60	1.33 – 9.70		27	32	32	3.31	1.29 – 8.47	
CCL22 pg/mL	200	200	200	1.09	0.73 – 1.63			103	103	103	1.31	0.74 – 2.32		97	97	97	0.92	0.49 – 1.74	
< 609	50	49	49	1	1			24	24	24	1			26	25	25	1		
609–784	50	44	44	0.84	0.47 – 1.50			24	21	21	0.96	0.40 – 2.30		26	23	23	0.68	0.29 – 1.57	
785–1031	50	51	51	1.06	0.61 – 1.87			31	23	23	1.04	0.43 – 2.52		19	28	28	1.18	0.54 – 2.59	
> 1031	50	56	56	1.08	0.61 – 1.91			24	35	35	1.87	0.77 – 4.54		26	21	21	0.66	0.29 – 1.50	
CCL24 pg/mL	200	200	200	0.91	0.64 – 1.28			103	103	103	0.93	0.56 – 1.56		97	97	97	0.91	0.56 – 1.47	
< 595	50	43	43	1	1			27	26	26	1			23	17	17	1		

	<i>All AIDS-NHL</i>						< 4 years time lag						≥ 4 years time lag					
	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N	OR	95% CI	Controls, N	Cases, N
595–1010	50	74	1.99	1.11 – 3.59	27	33	1.52	0.66 – 3.49	23	41	3.87	1.47 – 10.2	23	41	3.87	1.47 – 10.2		
1011–1475	50	27	0.57	0.29 – 1.12	26	16	0.58	0.22 – 1.56	24	11	0.61	0.22 – 1.68	24	11	0.61	0.22 – 1.68		
> 1475	50	56	1.50	0.80 – 2.81	23	28	1.63	0.65 – 4.09	27	28	1.60	0.63 – 4.05	27	28	1.60	0.63 – 4.05		
ICAM ng/mL	200	200	1.66	1.12 – 2.48	103	103	1.55	0.84 – 2.87	97	97	2.10	1.07 – 4.12	97	97	2.10	1.07 – 4.12		
< 148	51	32	1		23	15	1		28	17	1		28	17	1			
148–196	49	32	0.90	0.43 – 1.88	25	17	0.96	0.34 – 2.66	24	15	0.94	0.29 – 2.99	24	15	0.94	0.29 – 2.99		
196–289	50	65	1.81	0.99 – 3.31	29	36	1.97	0.80 – 4.85	21	29	2.54	0.98 – 6.60	21	29	2.54	0.98 – 6.60		
> 289	50	71	2.00	1.09 – 3.67	26	35	1.84	0.77 – 4.38	24	36	3.41	1.23 – 9.45	24	36	3.41	1.23 – 9.45		
IL18 pg/mL	200	200	1.93	1.30 – 2.88	103	103	2.06	1.12 – 3.79	97	97	1.66	0.94 – 2.93	97	97	1.66	0.94 – 2.93		
< 978	50	27	1		25	10	1		25	17	1		25	17	1			
978–1418	50	47	1.83	0.94 – 3.56	23	24	2.84	0.85 – 9.42	27	23	1.22	0.50 – 2.96	27	23	1.22	0.50 – 2.96		
1419–1987	50	59	2.12	1.10 – 4.10	33	34	2.23	0.71 – 7.04	17	25	1.86	0.73 – 4.77	17	25	1.86	0.73 – 4.77		
> 1987	50	67	2.55	1.32 – 4.92	22	35	4.28	1.40 – 13.1	28	32	1.57	0.64 – 3.86	28	32	1.57	0.64 – 3.86		
MCP1 pg/mL	200	200	1.60	1.05 – 2.42	103	103	1.80	0.95 – 3.41	97	97	1.39	0.81 – 2.39	97	97	1.39	0.81 – 2.39		
< 356	50	39	1		24	18	1		26	21	1		26	21	1			
356–501	50	54	1.25	0.68 – 2.31	19	31	1.85	0.74 – 4.62	31	23	0.68	0.27 – 1.72	31	23	0.68	0.27 – 1.72		
502–601	50	36	0.96	0.50 – 1.84	33	13	0.58	0.21 – 1.55	17	23	2.34	0.83 – 6.62	17	23	2.34	0.83 – 6.62		
> 601	50	71	1.74	0.98 – 3.09	27	41	1.78	0.78 – 4.06	23	30	1.94	0.79 – 4.77	23	30	1.94	0.79 – 4.77		
sCD163 ng/mL	200	200	1.39	0.92 – 2.09	103	103	1.48	0.77 – 2.83	97	97	1.39	0.80 – 2.44	97	97	1.39	0.80 – 2.44		
< 715	50	37	1		21	19	1		29	18	1		29	18	1			
715–933	50	34	0.88	0.46 – 1.68	26	14	0.46	0.16 – 1.32	24	20	1.35	0.54 – 3.37	24	20	1.35	0.54 – 3.37		
933–1271	50	59	1.67	0.91 – 3.05	31	32	1.17	0.51 – 2.67	19	27	2.77	1.02 – 7.50	19	27	2.77	1.02 – 7.50		
> 1271	50	70	1.94	1.03 – 3.66	25	38	1.82	0.72 – 4.63	25	32	2.15	0.84 – 5.52	25	32	2.15	0.84 – 5.52		
TRAIL pg/mL	200	200	1.25	0.93 – 1.68	103	103	0.85	0.50 – 1.44	97	97	1.51	0.99 – 2.30	97	97	1.51	0.99 – 2.30		
< 46	51	41	1		21	23	1		30	18	1		30	18	1			

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	<i>All AIDS-NHL</i>					<i>< 4 years time lag</i>					<i>≥4 years time lag</i>				
	Controls, N	Cases, N	OR	95% CI		Controls, N	Cases, N	OR	95% CI		Controls, N	Cases, N	OR	95% CI	
46–63	49	45	1.14	0.61 – 2.13		22	20	0.95	0.38 – 2.35		27	25	1.38	0.54 – 3.56	
64–79	52	42	0.88	0.48 – 1.62		30	20	0.43	0.17 – 1.10		22	22	1.40	0.53 – 3.68	
> 79	48	72	1.71	1.00 – 2.91		30	40	1.08	0.50 – 2.32		18	32	2.34	1.03 – 5.32	

N, number; OR, odds ratio; CI, confidence interval. ORs were adjusted for age at serum sample date (continuous), race/ethnicity (categorical), and HCV infection status at serum sample date.