Associations between B-cell non-Hodgkin lymphoma and exposure, persistence and immune response to hepatitis B

Although 90-95% of adults recover completely from Hepatitis B (HBV) infection, a minority are unable to clear the virus.¹ Epidemiological studies have demonstrated an increased risk of B-NHL among those with persistent HBV and B-NHL.²⁻⁵ However, the roles of exposure per se, occult infection, antibody response and viral clearance remain unclear.

Occult HBV infection (OBI) signifies persistence of the viral genome in blood and liver tissue in patients without hepatitis B surface antigen (HBsAg), with or without antibodies to hepatitis B core (anti-HBc) or hepatitis B surface (anti-HBs). Most subjects with OBI are in fact positive for anti-HBc and negative for HBsAg, while some are seronegative for all HBV markers.⁶ The association between OBI and B-NHL is clinically important due to the risk of reactivation of HBV following immunotherapy or chemotherapy. Few studies, however, have addressed OBI as a risk factor for NHL.⁷⁻⁹

Immune response to HBV (defined as anti-HBs) is elicited either by natural response to viral exposure (anti-HBc⁺ with anti-HBs⁺) or by vaccination (anti-HBc- with anti-HBs⁺). Limited data have suggested a negative association between the presence of anti-HBs and B-NHL.^{7,8}

Heredity plays a role in NHL etiology, as evidenced by the doubling of risk in first-degree relatives of NHL patients.¹⁰ It is unknown whether host genetic factors related to viral clearance are also related to NHL susceptibility.

The role of HBV infection in B-NHL has not previously been evaluated in Israel or the West Bank, where HBsAg seroprevalence has been estimated at 0.22% and 1.8%, respectively.^{11,12} In a case-control study (*Online*)

Supplementary Methods), we explored associations between exposure, persistence, and immune response to HBV with overall B-NHL as well as with two major sub-types: diffuse large B-cell lymphoma (DLBCL) and follicular lymphoma (FL). Additionally, we assessed whether a family history of hematopoietic malignancies were associated with HBV persistence.

We recruited 823 (516-Israelis/307-Palestinians) incident cases of B-NHL (median time from diagnosis- three months). DLBCL was the most common histology, comprising 427 (52%) cases, while FL was diagnosed in 186 (23%). We recruited 808 healthy controls (414-Israelis/394-Palestinians) from individuals accompanying patients to hospital or health centers. Cases and controls (*Online Supplementary Table S1*) differed in distributions of sex, age, marital status and family history of hematopoietic malignancies (*P*<0.01 for all); we adjusted for these variables in our models accordingly.

HBV serology was performed in 96.8% of participants (see results of individual markers, Table 1a). Most seromarkers (with the exception of eAg), including anti-HBc, HBsAg, anti-HBs and e-antibodies (anti-HBe), were more prevalent among Palestinians than Israelis. In contrast, antibody response to vaccine was more prevalent among Israelis. In addition, the combination of anti-HBc⁺ without anti-HBs, representing lack of immune response to HBV, was more common in Palestinians (9.0% vs. 3.0% in Israelis, P<0.0001) (Table 1b). Case-control association patterns were similar in the two groups.

Comparing cases and controls we found that persistence of HBV, as evidenced by HBsAg⁺, was associated with DLBCL (odds ratio (OR)=2.39, 95% confidenceinterval (CI):1.13-5.06). Moreover, the prevalence of HBsAg⁺ among those exposed to HBV (anti-HBc⁺) was higher in cases than in controls (15% vs. 7.8%, P=0.03), indicating lower viral clearance rates among cases. However, no case-control differences were found regarding exposure to HBV (anti-HBc⁺) (OR=0.94, CI:0.72-1.23)

а	Controls			Overall			B-NHL	DLBCL		FL		
	lsr	Pal	P *	lsr	Pal	P *	lsr	Pal	P *	lsr	Pal	P *
Biomarker	%	%		%	%		%	%		%	%	
HBsAg+	0.5	3.1	< 0.01	2.0	5.4	0.01	2.9	7.1	0.05	1.5	2.3	0.57
Anti-HBc⁺	14.6	34.0	< 0.0001	14.9	33.7	< 0.0001	16.2	35.7	< 0.0001	14.9	31.0	0.02
Anti-HBs+	28.5	38.1	< 0.01	21.3	29.2	0.01	18.5	30.5	< 0.01	24.3	30.2	0.43
Anti-HBe+	11.1	21.7	< 0.0001	10.0	17.7	0.002	9.9	17.1	0.03	13.3	23.3	0.12
HBeAg+	0.5	0.0	0.50	0.4	0.7	0.64	0.5	1.0	1.0	0.7	0.0	1.0
b Biomarker combinations												
OBI	14.1	30.5	< 0.0001	12.5	27.9	< 0.0001	12.7	28.1	< 0.0001	13.4	28.6	0.02
OBI with Anti-HBe+	8.3	17.4	< 0.0001	6.7	11.4	0.02	6.4	10.0	0.18	9.0	16.3	0.18
HBsAg+ with Anti-HBe+	0.5	3.1	< 0.01	1.8	4.7	0.02	2.5	6.2	0.06	1.5	2.3	0.57
Naturally immune	11.6	24.8	< 0.0001	9.4	20.8	< 0.0001	8.9	21.0	< 0.01	10.4	23.8	0.04
Immune via vaccine	16.9	13.6		11.2	8.1		8.9	9.5		13.4	4.8	
Lack of immune response	3.0	9.0	< 0.0001	5.5	12.8	< 0.0001	7.4	14.8	0.02	4.5	7.1	0.45
Total No.	414	394		516	307		210	217		143	43	

Table 1. Seropositivity for HBV and HCV biomarkers in Israelis versus Palestinians for Overall B-NHL, DLBCL, FL patients and controls.

B-NHL: B-cell non-Hodgkin lymphoma; DLBCL: diffuse large B-cell lymphoma; FL: follicular lymphoma; Isr: Israelis; Pal: Palestinians; HBsAg: hepatitis B surface antiber; Anti-HBc: hepatitis B core antibodies; Anti-HBs: hepatitis B surface antibodies; Anti-HBe: hepatitis B e antibodies; HBeAg: hepatitis B e antigen; Combinations: OBI (anti-HBc', HBsAg'); Naturally immune (anti-HBc', anti-HBs'); Immune via vaccine (anti-HBc', anti-HBs'); Lack of immune response (anti-HBc', anti-HBs'); *Fisher's exact and 2 tests comparing positive biomarker or combination versus negative.

or OBI (OR=0.84, CI:0.64-1.12) (Figures 1-2), anti-HBe or HBeAg prevalence (*data not shown*).

The presence of anti-HBs antibodies was inversely associated with B-NHL (OR=0.73, CI:0.57-0.93) and DLBCL (OR=0.68, CI:0.51-0.92). This pattern held for antibody response to vaccine (OR=0.67, CI:0.48-0.93 for overall B-NHL; OR=0.55, CI:0.36-0.85 for DLBCL), but was not statistically significant for natural antibody response (OR=0.76, CI:0.56-1.04). Conversely, a lack of antibody response was positively associated with DLBCL (OR=1.62, CI:0.1-2.61).

As expected, having a first-degree relative with hematopoietic cancer was associated with overall B-NHL (OR=1.69, CI:1.16-2.48) and DLBCL (OR=1.83, CI:1.17-2.87). Interestingly, this variable was also strongly associated with persistent HBV infection among controls (OR=6.80, CI:1.14-23.8).

In summary, we confirm that persistent HBsAg carriers had a significantly increased risk of DLBCL and a non-significant increased risk for overall B-NHL. The European Epilymph study reported a non-significant increased risk of B-NHL in HBsAg carriers (OR=1.58, CI:0.69-3.64) and DLBCL (OR=1.50, CI:0.47-4.82),² while a Turkish study reported OR=1.26, CI:0.65-2.46 and OR= 2.68, CI:1.19-6.01 for B-cell lymphoid tumors and DLBCL, respectively, and a strong relation for FL (OR=5.48, CI:1.02-29.5). Korean investigators reported a significant positive association between HBsAg and NHL (hazard ratio=1.74, CI:1.45-2.09) and a twofold risk of DLBCL in a cohort study⁴ and adjusted OR (2.09, CI:1.11-3.92) in a casecontrol study,⁵ respectively. In the latter study, investigators did not detect HBV-DNA in lymphoma tissues, but found HBV S, X, and C genes in DNA extracted from peripheral blood mononuclear cells, suggesting an indirect effect on lymphomagenesis.⁵

In the current study, the presence of anti-HBc was not associated with B-NHL or its subtypes, implying that exposure to HBV per se is not a risk factor for B-NHL. This finding is supported by a single Japanese cohort study of 20,360 subjects.¹³

Chen *et al.*, defining OBI as HBsAg⁻ and HBV-DNA⁺, reported a non-significant positive association with B-NHL.⁹ DNA extracted from tumor samples also failed to detect the HBV genome.

The inverse association between the presence of anti-HBs with B-NHL and DLBCL is consistent with Marcucci *et al.*'s report of an OR=0.61 (CI:0.44-0.85), both in indolent (OR=0.57, CI:0.38-0.88) and aggressive (OR=0.63, CI:0.44-0.93) B-NHL.⁷ Similarly, Wang *et al.* reported an OR of 0.60 (CI:0.40-0.70).⁸ In contrast, Kim *et al.* found no significant difference in anti-HBs⁺ prevalence comparing NHL patients with subjects with nonhematological malignancies and non-malignant conditions.¹⁴

Distinguishing between naturally acquired and vaccine-associated immunity is not clear cut as HBV DNA was not available to confirm actual exposure status. Nevertheless, we found a significant inverse association of B-NHL and DLBCL with anti-HBs overall. Similarly, Wang *et al.* reported a lower proportion of immune individuals among B-NHL cases.⁶ In contrast, the Epilymph study found no association between vaccine immunity and B-NHL.² However, the OR for DLBCL (OR=0.56, CI:0.29-1.06) was similar to the current study's point estimate.

Lack of immune response showed a significant positive association with DLBCL in our study. Likewise, Wang *et al.* concluded that patients with B-NHL may show lower clearance of the virus.⁸ Marcucci *et al.* also

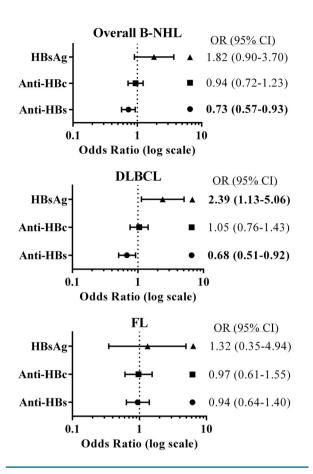


Figure 1. Forest plots showing the odds ratio (OR) and 95% confidence interval (Cl) for combinations of hepatitis B (HBV) biomarkers and their associations with overall B-cell non-Hodgkin lymphoma (B-NHL), diffuse large B-cell lymphoma (DLBCL) and follicular lymphoma (FL) for the pooled populations (Israelis and Palestinians). ORs were stratified by population (Israelis, Palestinians), sex and age categories (four-year grouping); adjusted for marital status, education (yrs), family history of hematopoietic malignancies in firstdegree relatives. Hepatitis biomarkers: hepatitis B surface antigen (HBsAg), hepatitis B core antibodies (anti-HBc) and hepatitis B surface antibodies (anti-HBs). Bold font indicates *P*<0.05.

demonstrated that lack of an antibody response was positively associated with B-NHL (OR=2.05, CI:1.24-3.37). These findings suggest that immune competence may be protective against B-NHL. Alternative explanations include diminished immune response to HBV in B-NHL patients due to the lymphoma itself or reduction in antibody titres due to treatment, especially rituximab,15 or waning immunity with age. Most of the sera among cases (75%) were collected post-treatment. However, in a subgroup with available paired samples, pre- and posttreatment sera showed agreement [(Kappa=1.0 for HBsAg and anti-HBs biomarkers, and Kappa=0.81 (CI:0.55-1.00) for anti-HBc)] due to two patients with positive pre-treatment anti-HBc⁺ becoming seronegative post-treatment. Wang et al.'s study, in which 85% of cases were recruited before treatment, also demonstrated a significant negative association between anti-HBs⁺ and B-NHL.

Intriguingly, we found a higher proportion of individuals with positive family history among HBsAg carriers than non-carriers in the control group. This association has not previously been reported and indicates the possi-

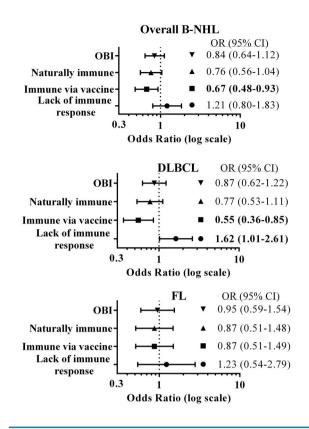


Figure 2. Forest plots showing the odds ratio (OR) and 95% confidence interval (Cl) for combinations of hepatitis B (HBV) biomarkers and their associations with overall B-cell non-Hodgkin lymphoma (B-NHL), diffuse large B-cell lymphoma (DLBCL) and follicular lymphoma (FL) for the pooled populations (Israelis and Palestinians). ORs were stratified by population (Israelis, Palestinians), sex and age categories (four-year grouping); adjusted for marital status, education (yrs), family history of hematopoietic malignancies in firstdegree relatives. HBV biomarkers: hepatitis B surface antigen (HBsAg), hepatitis B core antibodies (anti-HBC) and hepatitis B surface antibodies (anti-HBs). Combinations: Occult hepatitis B infection (OBI: anti-HBc', HBsAg⁻; Naturally immune (anti- HBc', anti-HBs'); Immune via vaccine (anti-HBc-, anti-HBs'); Lack of immune response (anti-HBc', anti-HBs'); Bold font indicates P<0.05.

bility of a joint inherited susceptibility to both diminished viral clearance and hematologic malignancy. Alternatively, it could reflect increased NHL risk in families where vertical transmission has occurred.

Limitations of this study include the inclusion of spouses as controls for 12% of enrolled cases. Use of spouse controls could, theoretically, alter case-control comparisons and bias the study toward the null. However, no couples were concordant for HBsAg⁺; in one couple, both members were anti-HBc⁺. Sensitivity analysis excluding spouse controls did not alter any of the reported associations (*not shown*). Other drawbacks are the lack of information on specific HBV genotypes, which may influence type and degree of immune responses.

The study's strengths include its unique population and the examination of several hepatitis biomarkers, enabling assessment of combinations and their association with B-NHL. Although a number of studies have examined associations between HBV seromarkers and B-NHL, few have clearly distinguished the roles of exposure, persistence and viral clearance in disease etiology.

Higher seroprevalence observed among Palestinians

implies that this population is at increased risk for HBV complications, including hepatocellular carcinoma and, potentially, B-NHL. It is particularly important to screen this population and all groups with high levels of exposure prior to immunosuppressive treatments in order to prevent potentially fatal flare-ups of pre-existing hepatitis infection.

In conclusion, our findings support an association between persistent HBV infection and B-NHL. We raise the possibility that hereditary factors may be related both to susceptibility to lymphoma and viral persistence, and that exposure per se does not explain the observed association between HBV and B-NHL. Prospective studies may clarify both the role of HBV vaccination in the prevention of B-NHL in endemic populations, and long-term B-NHL risk in individuals with poor antibody response to vaccine. The results of this study prompt a rethinking of the mechanism of virus-associated B-NHL beyond effects of chronic antigenic stimulation or viral integration, to include host response to infection as a marker of disease susceptibility.

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