Increased risk of melanoma in patients with chronic lymphocytic leukaemia: systematic review and meta-analysis of cohort studies

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An increased risk of melanoma has been variously reported in patients with chronic lymphocytic leukaemia (CLL), analogous with other immunosuppressed populations. To fully assess this association, we performed a systematic review and meta-analysis of available evidence from observational cohort studies. All such longitudinal studies of patients diagnosed with CLL that enabled quantitative assessment of the risk of melanoma compared with the general population were eligible. We identified seven studies from a search of all published literature to July 2014 in Medline, Embase and ISI science citation index databases. Data were pooled using a random-effects model. There was an almost four-fold increase in the risk of melanoma in patients with CLL compared with the general population (pooled standardized incidence ratio 3.88 [95% confidence interval (CI) 2.08-7.22]), although significant heterogeneity was evident among studies ($l^2 = 96.0\%$, $P_{\rm het}$ < 0.001). The risk of melanoma was higher for men with CLL (3.41; 95% CI 1.49-7.80) than women (2.61; 95% CI 1.13-6.01). CLL patients are at high risk of developing

Introduction

Melanoma is considered an immunogenic cancer, and populations with compromised immunity such as patients with solid organ transplants or those with HIV/AIDS are at increased risk [1,2]. Immunosuppression is also a characteristic of chronic lymphocytic leukaemia (CLL) [3], a malignancy belonging to the non-Hodgkin's lymphoma (NHL) group, and the most common leukaemia in adults [4]. Systemic therapy for CLL further impairs the immune response in these patients and although advances in treatment have improved response rates [5], this has been at the notable expense of prolonged, profound immunosuppression [6].

Increased risks of a range of second cancers among patients with CLL have been reported [4], and have been variously attributed to genetic susceptibility, treatment-related effects and shared risk factors, such as immuno-deficiency. Although an increased risk of melanoma has been reported in patients with CLL, the magnitude of the melanoma risk has yet to be reliably quantified as to date no systematic review of all relevant literature nor meta-analysis of the association has been carried out. A meta-analysis assessing the risk of melanoma in NHL patients reported a pooled relative risk of 1.85 [95% confidence interval (CI) 1.54–2.23] [7], but this analysis excluded CLL patients, who are more likely to present with

melanoma and the magnitude of the risk is higher than that found in other immunosuppressed populations. Our findings suggest that patients with CLL, as they are also at a higher risk of developing the more common skin cancers, would benefit from regular skin examinations. *Melanoma Res* 26:188–194 Copyright © 2016 Wolters Kluwer Health, Inc. All rights reserved.

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immune disturbances in comparison with other chronic lymphoproliferative disorders [8]. Given that CLL remains an incurable disease outside allogeneic stem cell transplantation [9] and survivors are living longer with advances in treatment, it is important to confirm the risk of melanoma in CLL patients so that they can benefit from regular screening of the skin, especially as these patients are also at high risk of developing keratinocyte skin cancers [10]. Understanding the association may also help to elucidate the role of shared aetiological factors. This systematic review and meta-analysis aimed to estimate the risk of melanoma in CLL patients compared with the risk of melanoma in the general population on the basis of all currently available evidence.

Methods

The systematic review and meta-analysis was performed in accordance with the Meta-analysis of Observational Studies in Epidemiology guidelines for reviews of observational Studies (MOOSE) [11] and we followed the PRISMA statement [12] to guide the reporting.

Literature search

Eligible studies up to July 2014 were identified by searching the Medline 1950 (US National Library of DOI: 10.1097/CMR.00000000000219

Medicine, Bethesda, Maryland, USA) database using PubMed software as the search interface; Embase 1966 database (Elsevier Science, Amsterdam, Holland) using the Embase search interface; and the ISI Science Citation Index using the ISI Web of Science search interface. We used the following medical subject headings terms or text words (both the US and UK spellings): melanoma, cancer, neoplasms, chronic lymphocytic leukaemia, CLL, leukaemia, non-Hodgkin's lymphoma, NHL, aetiology, cohort. The search was not limited to studies published in English. Relevant identified studies and review articles were included as citation search terms in the ISI Science Citation Index (1990 to the present) to identify subsequent studies that had referenced them. The abstracts of all identified studies were reviewed to exclude those that were clearly not relevant. The full texts of the remaining articles were read to determine whether they fulfilled the study inclusion criteria. Eligible studies were also identified by hand-searching the reference lists of retrieved articles.

Inclusion and exclusion criteria

We included all studies that were based on cohorts of patients diagnosed with CLL that enabled quantitative assessment of the risk of melanoma in CLL patients compared with the general population. Any discrepancies between investigators on inclusion of a study were resolved by joint evaluation of the manuscript. When multiple reports were published on the same population or subpopulation, we included in the meta-analysis the results from the longest follow-up or the most comprehensive data.

Data extraction and quality assessment

Two investigators independently abstracted data from identified studies using a standardized data abstraction form, with inconsistencies resolved by consensus. The following information was recorded for each study: study design, location, years of data collection, source and definition of cohort, number of cases, person-year duration of follow-up, age of study population, variables used for statistical adjustment, point estimates [relative risk, hazard ratio, or standardized incidence ratio (SIR)] and 95% CI. We evaluated the quality of primary studies using the Newcastle–Ottawa Scale [13], a validated technique for assessing the quality of observational studies.

Data synthesis and analysis

To pool individual study estimates for the risk of melanoma in CLL patients, we used the method of DerSimonian and Laird [14] using a random-effects model. Statistical heterogeneity among studies was assessed using the Q-statistic [15] (significance level at P < 0.10), and inconsistencies were quantified using the I^2 -statistic [16]. We carried out a sensitivity analysis by excluding one study at a time and calculated the pooled

relative risk for the remaining studies to evaluate whether the results could have been affected markedly by a single study. Subgroup analyses were carried out according to geographic region and sex. Finally, publication bias was evaluated through visual inspection of a funnel plot and using Begg's and Egger's tests [17,18]. All statistical analyses were carried out using Stata, version 10 (Stata Corporation, College Station, Texas, USA).

Results

Search

Details of the selection process for the eligible studies are shown in Fig. 1. A total of nine eligible studies were identified [19–27], three of which reported on overlapping populations [22,23,27] from Denmark, and one of three being restricted to a high-risk subgroup of CLL patients with a family history of cancer [23]. For these three studies, we therefore included only the most recent and comprehensive report [22], leaving a total of seven eligible studies.

Characteristics of included studies

The main characteristics of the studies included are listed in Table 1. The majority of the studies were of cutaneous melanoma. Studies were published between 1978 and 2011. Four of the studies were carried out in North America [20,21,25,26] and one each in Australia [19], Denmark [22] and Scotland [24]. With the exception of one study that reported on a clinical cohort from a single medical centre [21], all studies were national cancer registry linkage studies and melanoma diagnoses were ascertained through cancer registries for all seven studies. Two studies included patients diagnosed with small lymphocytic lymphoma (SLL) in their cohorts [20,21]. We included these studies as CLL and SLL are considered phenotypic variants on the same disease spectrum by many authors [28], including the WHO classification of tumours of haematopoietic and lymphoid tissues [29]. Moreover, there are few data describing the extent of immune dysfunction between patients with CLL versus SLL. The mean or the median follow-up time for the cohorts ranged from 3.4 to 6.3 years. All studies had accounted for age, sex and time period in their calculations. One study had additionally adjusted for ethnicity [20] (88% white), another reported that 100% of the study population was white [26] and no information on ethnicity was reported by the remaining five studies. With the exception of one study [24], all studies reported a statistically significant increased risk of melanoma.

Quality assessment

One study scored 8 out of a possible score of 9 using the Newcastle–Ottawa quality assessment scale for cohort studies [20]; the remaining studies scored 7 points.



Flow chart of literature search for studies on the association between CLL and melanoma. CLL, chronic lymphocytic leukaemia.

Outcome

The pooled SIR (pSIR) for studies reporting on risk of melanoma in CLL patients compared with the general population was 3.88 (95% CI 2.08–7.22), with evidence of significant heterogeneity ($I^2 = 96.0\%$, $P_{het} < 0.001$) (Fig. 2). All seven studies reported an increased risk that was statistically significant in all except one study [24]; the heterogeneity thus reflected differences in the size of the effect rather than the direction of the effect. In sensitivity analyses excluding each study in turn, the summary estimate ranged from 3.30 (95% CI 2.09–5.22) with the omission of Royle *et al.* [19] to 4.51 (95% CI 2.62–7.75) with the omission of Morton *et al.* [20]. There

was no evidence of publication bias (Begg P=0.764; Egger $P_{het}=0.358$).

The pSIR was higher for the four studies carried out in the USA (3.68; 95% CI 1.98–6.87; $I^2 = 90.1\%$, $P_{het} \le 0.001$) than for the two European studies (2.42; 95% CI 1.66–3.51; $I^2 = 0.0\%$, $P_{het} = 0.97$), whereas the single Australian study reported the highest SIR of all (7.74; 95% CI 6.86–8.73). On the basis of the four studies that presented SIRs stratified by sex [19,20,24,25], the pSIR was higher for men (3.41; 95% CI 1.49–7.80; $I^2 = 96.7\%$, $P_{het} < 0.001$) than women (2.61; 95% CI 1.13–6.01; $I^2 = 91.1\%$, $P_{het} < 0.001$). Several studies also

				Cohort (male)			Comparison population	Mean or median cohort
References	Location	Study period	Cohort description	[<i>v</i>]]	Cases (n) type	Study type	(measure of effect)	diagnosis age (follow-up)
Royle <i>et al.</i> [19]	Australia	1983-2005	CLL primary diagnosis	13580 (59)	272	Record linkage from	Australian population (SIR)	Median: 69 (4.3)
					100% cutaneous	state data		
Morton <i>et al.</i> [20]	USA, 11 cancer registries	1992-2006	CLL/SLL diagnosis	15 915 (58)	82	Record linkage	US population (SIR)	Mean: 67.2 (4.3)
					100% cutaneous			
Tsimberidou <i>et al.</i>	USA, University of Texas Anderson	1985-2005	CLL/SLL presentation	2028 (61)	19	Clinical cohort	US population (SIR)	Median: 58 (6.3)
[21]	Cancer Research Centre				NS			
Schöllkopf et al. [22]	Denmark	1943-2003	CLL diagnosis	12373 (61)	27	National record	Danish population (SIR)	Median: 70 (mean: 3.8)
					100% cutaneous	linkage		
VicKenna <i>et al.</i> [24]	Scotland	1975-1997	CLL index case	4016 (59)	9	National record	Scottish population (SIR)	Median: 72.2 (NR)
					100% cutaneous	linkage		
lravis <i>et al.</i> [25]	USA, 9 cancer registries	1973-1988	CLL first primary	9456 (58)	28	Record linkage	US population (O/E)	Mean: 70 (4.2)
			cancer		75% cutaneous			
					25% ocular			
Greene <i>et al.</i> [26]	USA, End Results Program NCI	1935-1971	CLL initial cancer	4869 (64)	6	Record linkage	US population (SIR)	Mean: NR (3.4)
	(100 hospitals)				NS			
CLL, chronic lymphoc	sytic leukaemia; NR, not reported; NS	, not specified;	O/E, observed/expected;	SIR, standardize	d incidence ratio; SLL,	small lymphocytic lymph	ioma.	

Table 1 Characteristics of the seven studies included in the meta-analysis of risk of melanoma in CLL patients

presented SIRs stratified by age at diagnosis [19,20,24] and time from diagnosis of CLL [19,20,24], although because there was heterogeneity in the categorization of time periods/age-groups, estimation of combined estimates of melanoma risk by age group was not possible. A sensitivity analysis excluding one study that included ocular melanoma [25] and two studies that did not specify melanoma type [21,26] resulted in a pSIR of 3.21 (95%) CI 1.20-8.60).

Discussion

We have examined all available evidence from observational studies reporting on the risk of melanoma in CLL patients and show that they have an almost four-fold increased risk compared with the general population. Cases were predominantly cutaneous and when we excluded studies where the site was mixed or unclear, risk was still increased by over three-fold. A similar but smaller increase in melanoma risk is also observed in other immunosuppressed populations including organ transplant recipients [1] (pSIR 2.3) and patients with HIV/AIDS (pSIR 1.5) [2] as well as in patients with NHL (pSIR 1.9) [7], but CLL (along with NHL more broadly [30–33]) is unusual because of a bidirectional relationship observed with melanoma [27,34]. Not only are CLL patients at increased risk of subsequent melanoma, but patients diagnosed with melanoma are at increased risk of subsequent CLL, suggesting the existence of shared risk factors. Risk factors for melanoma are well understood, namely sun-sensitive pigmentary phenotype and excessive exposure to ultraviolet radiation, but the aetiology of CLL remains to be fully elucidated [35,36]. Although several lines of evidence support a strong inherited genetic component [37], the role of potentially carcinogenic environmental factors such as ionizing radiation [38,39], pesticides and other chemicals [36,39,40] is less clear, with considerable heterogeneity in the published literature. Evidence on the association of CLL/NHL with exposure to UV radiation is contradictory [36,41]. There is some evidence of a genetic component in the common pathogenesis of CLL and melanoma. Genomewide association studies have identified germline mutations in genes in the TERT-CLPTM1L region, associated with telomerase activity, in both melanoma [42] and CLL [43]. Also associated with telomerase activity, loss-of-function mutations in POT1 have been found in familial melanoma [42] and have been found to be somatically mutated in CLL (3.5% of all CLL and 9% with an aggressive CLL subtype) [44]. Genome-wide association studies have also reported mutations in CASP8, involved in apoptosis, associated with melanoma [45] but less consistently with CLL [37]. Finally, longterm immunosuppression is a risk factor for both cancers [46], and is likely to be the strongest common underlying factor that explains the association.





Forest plot of the association between CLL and melanoma compared with the general population. Each line represents an individual study result with the width of the horizontal line indicating 95% CI, the position of the box representing the point estimate and the size of the box being proportional to the weight of the study. CI, confidence interval; CLL, chronic lymphocytic leukaemia; pSIR, pooled standardized incidence ratio.

The increased risk of melanoma in CLL patients is most likely explained by the state of immunosuppression inhibiting the antitumour response. The immune defects associated with the disease are complex and include decreased immunoglobulin synthesis as well as changes in the innate immune system including defects in complement activation, depressed cellular immunity and impaired phagocytosis [47]. Treatment of CLL can further impair the immune response, although currently, there is no evidence of an association between treatment with alkylating agents or nucleoside analogues and risk of melanoma [48,49], suggesting that treatment-related immunosuppression does not increase risk. The higher risk of melanoma found in CLL patients compared with other immunosuppressed populations [1,2] most likely reflects the more severe and prolonged degree of immunosuppression among the former.

All studies included in our meta-analysis received a highquality score using the Newcastle–Ottawa quality assessment scale for cohort studies [13]. There were some limitations in our meta-analysis. Our analyses may have been subject to publication bias because we did not search for unpublished studies, abstracts or the grey literature; however, inclusion of only results from peerreviewed studies provided greater assurance of the quality of those data we did include. The mean/median follow-up time for studies included in the meta-analysis ranged from 3.4 to 6.3 years, which may not have sufficiently covered the latent period for melanoma development, leading to a potential underestimation of the association. Limited control of confounding may also have influenced the results as the studies included did not have information on key melanoma risk factors including sun exposure, which has been inconsistently associated with the broader group of NHL [41] and also has an effect on immune function [50]. Similarly, five of the seven studies included did not specify the ethnic composition of the populations included. CLL is common in individuals of European descent, but is rare in Asians [51]; this is also true of melanoma, however, and thus lack of information on ethnicity is unlikely to have influenced the observed association.

Finally, although all studies included reported an increased risk, we observed significant heterogeneity in the magnitude of the effect. This heterogeneity was not explained by geographical variation, and we could not examine the effects of baseline differences in clinical parameters as these were mostly not reported. Studies included in our analysis were based on patients recruited over an extended time period (1935–2006), during which there have been considerable changes in treatment regimens [5].

Although CLL treatment advances have led to impressive benefits in terms of overall survival, this has been at the cost of substantial long-term immunosuppression [6]. The increased longevity of patients with CLL means that, on average, they are now at risk of melanoma for longer periods. It is unknown whether CLL patients diagnosed with melanoma are at greater risk of disease recurrence, but melanoma survival is worse in patients with CLL than in patients without CLL [52]. Given the heterogeneous clinical profile of CLL patients [53], collection of detailed clinical information on disease stage and treatment in future large prospective studies would aid better understanding of the strong association between CLL and melanoma that we have shown.

In summary, patients diagnosed with CLL are a high-risk population for the development of melanoma and as they are also at a higher risk of developing the more common types of skin cancer [10,34,54], these patients would benefit from routine skin examinations as well as counselling to avoid excessive sun exposure.

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Conflicts of interest

There are no conflicts of interest.

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