Risk of non-Hodgkin’s lymphoma following tuberculosis

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Summary To study the association between chronic infections and non-Hodgkin’s lymphoma (NHL), we assessed the risk of NHL in a Swedish cohort of 5050 individuals with tuberculosis 1939–1960. The overall relative risk was moderately increased, largely accounted for by high risks following severe tuberculosis diagnosed a long time ago. © 2001 Cancer Research Campaign http://www.bjcancer.com

Keywords: non-Hodgkin’s lymphoma; tuberculosis; cohort; Sweden

Over the past decades, the incidence of non-Hodgkin’s lymphoma (NHL) has increased in many western countries (Devesa and Fears, 1992) for reasons that are largely unknown. Chronic inflammation has been linked to increased occurrence of NHL (Holmes et al, 1989; Baekklund et al, 1998). Several moderate-sized case-control studies of patients with NHL have found associations with histories of chronic infections including tuberculosis (Bernard et al, 1984; Franceschi et al, 1989; Doody et al, 1992; La Vecchia et al, 1992; Tavani et al, 2000). During the first half of the 20th century, the incidence and case-fatality of tuberculosis decreased markedly (The Swedish Institute for Infectious Disease Control, 1989), in the same birth cohorts who have later experienced an increasing incidence of NHL. Based on this ecological correlation, and on the positive association between the two diseases reported in case-control studies, we hypothesized that individuals with a history of tuberculosis would be at increased risk of NHL.

At two Swedish tuberculosis dispensaries and one sanatorium with population-based catchment areas, we identified 5199 individuals consecutively diagnosed or admitted with tuberculosis in the years 1939–1960. For each individual, we collected information on date of birth, sex, date of entry (diagnosis or admission), location of tuberculosis (pulmonary vs extrapulmonary), result of sputum microscopy on admission (sanatorium patients up to 1953 only), admission to thoracic surgery (sanatorium patients up to 1953 only), and socioeconomic status (white collar, blue collar, unskilled worker, or missing, categorized on the basis of the occupation of the head of the family). The cohort was linked to the population-based Swedish cancer register for 1958–1996, whichever occurred first. 95% confidence intervals (95% CI) were calculated assuming a Poisson-distribution for the specific rates. Start of follow-up was set to last of: date of diagnosis of tuberculosis or 1 January 1952 (SMR) or 1958 (SIR). End of follow-up was set to date of death, emigration, or 31 December 1996, whichever occurred first. 95% confidence intervals (95% CI) were calculated assuming a Poisson-distribution for the observed cases (Breslow and Day, 1987). Poisson-regression was used to compare risk estimates (Breslow and Day, 1987).

2797 deaths were registered during 163 270 person-years of follow-up, the SMR being 1.40 (95% CI = 1.35–1.46). The SMR declined with time since tuberculosis, and was 1.17 (95% CI = 1.07–1.27) after 40 or more years since tuberculosis (P for linear trend < 0.0001). Individuals who entered the cohort 1939–1952 had a significantly higher SMR than those who entered 1953–1960 (adjusted for time since tuberculosis, data not shown). Admission (vs no admission) to thoracic surgery, positive (vs negative) direct microscopy, extrapulmonary (vs pulmonary) tuberculosis, and low (vs high) socioeconomic status were each associated with significantly higher SMRs (data not shown). 1073 cancers were registered and the SIR was 1.20 (95% CI = 1.13–1.28). The all-site SIRs were similar in all latency periods and did not vary significantly with calendar period of entry, admission to thoracic surgery, sputum smear status, location of tuberculosis, or socioeconomic status (data not shown). A total of 33 cases of NHL were registered and the SIR was 1.42 (95% CI = 0.98–1.99) (Table 1). The relative risk was highest after 40 or more years following entry (SIR = 2.25, 95% CI = 1.23–3.78, n = 14). The SIRs were close to unity in each 5-year interval up to 40 years since the tuberculosis, but above 2.0 in each subsequent interval (data not shown). Individuals who entered the cohort 1939–1952 had a significantly higher risk (SIR = 1.77, 95% CI = 1.16–2.57, n = 27) than those who entered 1953–1960, whose risk was not elevated (Table 1). When adjusted for each other, the elevated risks associated with more than 40 years of latency, and with entry during 1939–1952, remained largely similar yet no longer significant. Male sex, positive (vs negative) sputum smear, and admission (vs no admission) to thoracic surgery were associated with non-significantly higher SIRs, and the risks were similar for pulmonary and extrapulmonary tuberculosis (data not shown). However, 14 of the 33 cases of NHL occurred among individuals with either positive sputum smear or admission to
surgery (SIR = 2.77, 95% CI = 1.51–4.65), compared to five among individuals not admitted to surgery and with negative sputum microscopy (SIR = 0.70, 95% CI = 0.23–1.63), and 14 among individuals with no information on either characteristic (SIR = 1.26, 95% CI = 0.69–2.12, \( P \) for linear trend over estimates = 0.003). Among those smear-positive or admitted to surgery, the risk was significantly higher after 40 or more years since tuberculosis (SIR = 5.34, 95% CI = 2.44–10.1, \( n = 9 \)) compared to 0–39 years since tuberculosis (Table 1). Alternative categorizations of latency time resulted in a similar risk pattern. The SIR of Hodgkin’s disease was not significantly elevated (SIR = 1.65, 95% CI = 0.71–3.26, \( n = 8 \)) but displayed a similar pattern as NHL with the highest point estimate 40 or more years since the tuberculosis (SIR = 4.18, 95% CI = 0.51–15.1, \( n = 2 \)).

This study indicates that the risk of NHL following tuberculosis is increased, but only among individuals diagnosed with tuberculosis before 1953, and among them, in particular among individuals with severe tuberculosis. The increased overall risk of NHL corroborates the results of several case-control studies of patients with NHL (Bernard et al, 1984; Franceschi et al, 1989; Doody et al, 1992; La Vecchia et al, 1992; Tavani et al, 2000), although some have found no association (Tielsch et al, 1987; Cartwright et al, 1988). None of the case-control studies have, however, included characteristics of the tuberculosis. Two cohort-studies (performed for other purposes) of patients with tuberculosis analysed the risk for NHL (Howe et al, 1979; Davis et al, 1989) but neither found a positive association. However, in one study patients were followed for only 23 years, and proportionate mortality ratios were used which are inherently difficult to interpret in the presence of competing mortality (Howe et al, 1979). The other found an increased overall risk of dying from Hodgkin’s disease but no increased risk for the combined group of non-Hodgkin’s lymphoma, multiple myeloma and polycythemia vera. No separate analysis of non-Hodgkin’s lymphoma was performed (Davis et al, 1989). No analysis of latency was presented. An increased risk for NHL may thus well have been concealed in both these studies.

Patients with tuberculosis may differ from the general population with respect to several factors, such as socioeconomic status, use of tobacco, alcohol and coffee, exposure to ionizing radiation and isoniazid, and body mass index. Since none of these factors have been consistently identified as risk factors for NHL (Stott et al, 1976; Franceschi et al, 1989; Devesa and Fears, 1992; La Vecchia et al, 1992; Ron et al, 1994; Tavani et al, 1994; Nelson et al, 1997; Adami et al, 1998; Holly et al, 1999; Zhang et al, 1999), substantive confounding by these factors is unlikely. Surveillance bias, that is, if individuals with a history of tuberculosis received closer medical surveillance than healthy individuals, would more likely have resulted in generally increased risks rather than the observed calendar period-dependent risks. The classification of sputum smear status, location of tuberculosis, and surgical treatment does not take into consideration later relapses. Bias resulting from such misclassification would, however, attenuate rather than create differences between exposure-groups, and is thus an unlikely explanation for the observed variations in risk. The lack of information on severity during the 1953–1960 period of entry precludes a more detailed risk-analysis within this interval. However, it does not change the absence of overall risk increase during the interval, although the low number of expected cases in the long latency interval during this period makes it difficult to completely exclude a long-term effect. Validation-studies have found that one third of the registered cases of Hodgkin’s disease during the 1970s were misclassified NHL (Martinsson et al, 1992). We therefore also assessed the risk of Hodgkin’s disease. The similarity of the SIRs for NHL and for Hodgkin’s disease may be due to misdiagnosis of the former as the latter.

Severe tuberculosis (defined as individuals with positive sputum smear or admission to surgery, who had the highest overall mortality) was associated with a markedly increased risk of NHL. In contrast, individuals diagnosed after 1952, when curative chemotherapy (isoniazid) was introduced, were not at increased risk of NHL. Inflammatory intensity and duration are important risk determinants for malignant lymphomas complicating rheumatoid arthritis and coeliac disease (Holmes et al, 1989; Baecklund et al, 1998) and our results suggest that these factors also may influence the association between tuberculosis and NHL: limited infection did not increase NHL risk (tuberculosis diagnosed after 1952 or uncomplicated tuberculosis diagnosed 1939–1952), while the highest inflammatory load was associated with the highest risk increase (long-standing, severe tuberculosis diagnosed 1939–1952).

There is evidence of genetic variation in the susceptibility to tuberculosis, through genes encoding components of the inflammatory response (Bellamy et al, 1998). Since it has been proposed that low-secretor genotypes of TNF-alpha are associated with increased susceptibility to follicular NHL (Fitzgibbon et al, 1999), an alternative interpretation of our results is that NHL is a result of increased susceptibility rather than a consequence of tuberculosis itself.

### Table 1

| Period of entry | Subjects | 0–39 years since TB | 40+ years since TB | All latency intervals |
|----------------|---------|---------------------|-------------------|----------------------|
| 1939–1952      | All     | 1.38 (0.73–2.34) 13 | 2.42 (1.33–4.07) 14 | 1.77 (1.16–2.57) 27* |
|                | Smear positive or operated | 1.48 (0.48–3.46) 5 | 5.34 (2.44–10.1) 9* | 2.77 (1.51–4.65) 14* |
|                | No information | 2.22 (0.72–5.17) 5 | 3.70 (0.76–10.8) 3 | 2.61 (1.13–5.14) 8 |
|                | Smear negative, not operated | 0.77 (0.16–2.26) 3 | 0.61 (0.07–2.20) 2 | 0.70 (1.23–1.63) 5 |
| 1953–1960      | All     | 0.79 (0.29–1.73) 6 | 0.00 (0.00–5.52) 0 | 0.75 (0.27–1.63) 6 |
| Both periods   | All     | 1.11 (0.67–1.74) 19 | 2.25 (1.23–3.78) 14* | 1.42 (0.98–1.99) 33 |

*P for difference compared to entry in 1953–1960 = 0.04; \( P \) for difference compared to 0–39 years since tuberculosis = 0.02; \( P \) for linear trend from no, through missing, to yes = 0.001; \( P \) for difference compared to individuals with negative sputum smear and not operated on = 0.004; \( P \) for difference compared to 0–39 years since tuberculosis = 0.05.
In conclusion, this study suggests that the risk of NHL is increased in individuals with a history of tuberculosis, but only among individuals with tuberculosis, particularly severe, diagnosed a long time ago. Whether the observed risk increase is due to the tuberculosis itself, an underlying susceptibility, or associated exposures is not clear. Our results raise the question whether one reason behind the increasing incidence of NHL over the past decades is the increasing survival into older age groups of individuals with what used to be lethal infections of early adulthood, such as tuberculosis.

ACKNOWLEDGEMENTS

This study was supported by grants from the Swedish Heart Lung Foundation and from the National Heart and Lung Association.

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