REVIEW ARTICLE

Lung cancer risk at low cumulative asbestos exposure: meta-regression of the exposure–response relationship

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Abstract

Purpose Existing estimated lung cancer risks per unit of asbestos exposure are mainly based on, and applicable to, high exposure levels. To assess the risk at low cumulative asbestos exposure, we provide new evidence by fitting flexible meta-regression models, a notably new and more robust method.

Methods Studies were selected if lung cancer risk per cumulative asbestos exposure in at least two exposure categories was reported. From these studies $(n = 19)$, we extracted 104 risk estimates over a cumulative exposure range of 0.11–4,710 f-y/ml. We fitted linear and natural spline meta-regression models to these risk estimates. A natural spline allows risks to vary nonlinearly with exposure, such that estimates at low exposure are less affected by estimates in the upper exposure categories. Associated

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relative risks (RRs) were calculated for several low cumulative asbestos exposures.

Results A natural spline model fitted our data best. With this model, the relative lung cancer risk for cumulative exposure levels of 4 and 40 f-y/ml was estimated between 1.013 and 1.027, and 1.13 and 1.30, respectively. After stratification by fiber type, a non-significant three- to fourfold difference in RRs between chrysotile and amphibole fibers was found for exposures below 40 f-y/ml. Fibertype-specific risk estimates were strongly influenced by a few studies.

Conclusions The natural spline regression model indicates that at lower asbestos exposure levels, the increase in RR of lung cancer due to asbestos exposure may be larger than expected from previous meta-analyses. Observed potency differences between different fiber types are lower than the generally held consensus. Low-exposed industrial or population-based cohorts with quantitative estimates of asbestos exposure a required to substantiate the risk estimates at low exposure levels from our new, flexible metaregression.

Keywords Amphiboles - Asbestos - Chrysotile - Exposure - Lung cancer - Meta-analysis

Introduction

It is widely accepted that exposure to asbestos is related to an excess risk of lung cancer [1]. However, studies exploring the exposure–response relationship have shown a large variability in excess risk per unit of exposure. Berman and Crump [2] showed in a meta-analysis that such variations might be related to different fiber size distributions and fiber type. Within fiber type, relatively longer

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fibers were associated with a higher increased lung cancer risk compared to shorter fibers. Moreover, chrysotile was estimated to be less potent than amphiboles by a factor ranging between 6 and 60. The meta-analysis of Hodgson and Darnton [3] had comparable findings with respect to potency differences between chrysotile and amphiboles. In a recent meta-analysis by Lenters et al. [4], it was shown that variations in risk estimates of lung cancer might additionally be explained by differences in quality aspects of the applied exposure assessments methodology besides fiber type.

In all previous meta-analyses $[2, 4, 5]$ except for the one of Hodgson and Darnton [3], a fixed, average excess risk per fiber year (expressed as the potency, that is, the socalled K_L value, of asbestos for causing lung cancer) was estimated by combining the K_L values obtained for each study. However, most of the lung cancer studies included in the meta-analyses were not very recent and notably involved heavily exposed individuals. Currently, certainly in the Western world, it is unlikely that individuals are exposed to levels previously generally studied, because handling of asbestos declined gradually after the 1970s and dropped severely in the 1990s due to directives on protecting workers exposed to asbestos [6]. As a consequence, current interest lies in estimating excess risk accurately at relatively low exposures.

Previously, linear extrapolation has been applied to estimate risks at low exposure levels. However, such extrapolation is heavily dependent on estimates at high exposures, rendering extrapolated risk estimates at low exposure uncertain. For example, the population-based study of Gustavsson et al. [7] found a significant excess risk of lung cancer at low levels of cumulative asbestos exposure, which was much higher than could be expected by simple linear extrapolation from cohorts with higher exposures.

To accurately derive acceptable exposure limits and underpin compensation claims, better evidence is needed about the asbestos-related risk of lung cancer at low exposures. We provide new evidence by fitting nonlinear meta-regression models to existing data, which is notably new in meta-analyses. The flexibility of these models ensures that the exposure–response relationship can vary with exposure levels and is less affected by estimates in the upper exposure categories [8]. Moreover, the advantage of our method is that we combine all existing risk estimates at low exposures and obviate the need to extrapolate below the study-specific exposure range. Hence, our method provides a more robust estimate of exposure-specific lung cancer risks than previous meta-analyses. In addition, we stratified our results by fiber type to explore a potential potency difference between chrysotile and amphibole fibers at relatively low cumulative exposure levels.

Methods

Identification of included studies

The same selection criteria were applied as in the metaanalysis by Lenters et al. [4]. Briefly, occupational studies from MEDLINE and EMBASE were selected if lung cancer risk per cumulative exposure in at least two exposure categories was reported. Furthermore, the cumulative exposure needed to be reducible to units of total number of fiber years (f-y/ml), which is defined as the product of the concentration of asbestos fibers per milliliter of air measured by phase-contrast microscopy (PCM), and the duration of exposure in years. PCM measures fibers that are longer than 5 μ m, thicker than approximately 0.25 μ m, and with an aspect (length-to-width) ratio >3 . Studies with only one exposure category were excluded because no studyspecific exposure–response relationship could be derived. The selection criteria resulted in 18 industry-based cohort studies, including one nested case–control, and one general population-based case–control study (see Table 1 for details). In all studies, exposure to asbestos was based on data from stationary or personal monitoring.

Extraction of data from the incorporated studies

Information about the study design, study characteristics, and exposure categories were extracted from each study. To obtain risk estimates for the 15 studies with standardized mortality ratios (SMRs), observed and expected lung cancer cases were extracted for each exposure category. The relative risks (RRs) and their confidence intervals (CIs), size of the study population, and number of lung cancer cases were extracted for each exposure category with lung cancer occurrence among the two cohort studies with an assigned reference group. For the two case–control studies, the odds ratios (ORs) and their confidence intervals, and the number of lung cancer cases and controls were included. The adjusted ORs and corresponding CIs for the study of Gustavsson et al. [7] were obtained via direct communication with the authors. For the purpose of this meta-analysis, all measures of association, that is, ORs, RRs, and SMRs, were considered estimates of the RR of asbestos exposure and lung cancer occurrence.

To assign a specific point estimate of cumulative exposure to the extracted risk estimates, we used the mean of the exposure category, when described in the original publication. If not described, the midpoint of the range of the exposure categories was used. For open-ended, uppermost exposure categories, the midpoint was calculated as 5/3 times the lower bound of those categories (as proposed by the asbestos advisory committee of the Unites States Environmental Protection Agency in 2008). For example,

the midpoint estimate for an open-ended category of >100 fiber years was calculated as $5/3 * 100 = 167$. For additional details on data extraction, we refer to Lenters et al. [4].

Modeling the exposure–response relationship

We hereby expanded on the study of Lenters et al. [4] in which they investigated the role of quality of the asbestos exposure assessment to potentially explain heterogeneity in linear exposure–response slope estimates. They showed that the linear exposure–response slope estimates can be influenced by measurement error. Moreover, linear extrapolations to lower exposures based on these estimates likely yield a large uncertainty as they did not focus on the actual shape of the exposure–response curve. To improve estimates in the low exposure range, we assessed the shape of the exposure–response curve by fitting nonlinear metaregression models to all available data estimates.

From the 19 studies, we extracted 104 risk estimates (i.e., study points of the RR for lung cancer at a given exposure level) over a cumulative exposure range of 0.11–4,710 f-y/ml. To accurately estimate associations in the lower exposure range based on all available data points, we used a previously developed macro for applying linear and nonlinear regression models to the reported risk estimates $[8]$. In this macro, the natural logarithm (LN) of the reported risk estimates was inversely weighted by their variance [9]. As risk estimates (ORs and RRs) within a single study are correlated, the variance of the risks was corrected by estimating the covariance between different risk estimates using the method of Greenland [10]. For studies reporting SMRs, no covariance was estimated as it can be assumed that the independence assumption does hold for SMRs since the total population is used as the reference group instead of a subsample.

The regression models applied consisted of full linear models (model type 1), and natural splines with prespecified knots at the 20th, 50th, and 80th percentiles (model type 2). The natural spline is a flexible model and allows risks to vary nonlinearly with exposure, such that estimates at low exposure are less affected by estimates in the upper exposure categories [11]. The two model types were fitted with (option A) and without (option B) an intercept where model A assumes a difference in background rate of lung cancer between exposed and non-exposed individuals and model B assumes no difference. A model with intercept has been used in previous studies to account for potential differences in background risk [2, 4, 5, 12]. However, if an intercept above $RR = 1$ is due to measurement error instead of differences in background risk, it is more appropriate to model the exposure–response relationship without intercept [13]. To accommodate potential betweenstudy heterogeneity, the regression models allowed for random study-specific intercepts and exposure effects [9]: LN RR = $\beta_0 + \beta_1^*$ exposure $+\sigma_{u0}^2 + \sigma_{u1}^2 + \sigma_{e0}^2$ (model option A); LN RR = β_1^* exposure $+\sigma_{u1}^2 + \sigma_{e0}^2$ (model option B);

where β_0 is the common intercept across studies, β_1 is the common slope associated with exposure across studies, σ_{u0}^2 is the estimated variance of the intercept between studies, σ_{ul}^2 is the estimated variance of the slope between studies, and $\sigma_{\rm e0}^2$ is the variance of the individual risk estimates. (For the spline models an additional spline variable was estimated by using third-order polynomials to fit a nonlinear slope [11]).

As a result, an additional component of the variance explaining the between-study heterogeneity was considered in weighting each observation [14]. Models were fitted using maximum likelihood (ML) estimation, and goodness of fit was assessed with the deviance (-2 log likelihood) criterion. For accurate estimation of the parameters, models were refitted using restricted maximum likelihood (REML). A variance components structure was used to compute the between-study variances for option A.

The results on the LN scale were retransformed to the 'normal' scale to identify the variation in RR as a function of exposure. We calculated the RRs and their CIs for low cumulative exposure levels of 4 and 40 f-y/ml. These levels were selected because occupational exposure standards have been endorsed from levels of 2 to 0.1 f/ml over an 8-h time weighted average in the past decades [15, 16]. Over a working life exposure of 40 years, we expect the cumulative exposure levels of workers over the last decades to be somewhere between 4 and 40 f-y/ml. For models with an intercept (option A), the predicted RR at zero exposure may not be equal to 1. To relate the estimated risk at a specific exposure level to an RR of 1 at zero exposure, models were refitted to the data points from which the common predicted intercept was subtracted. Results were stratified by fiber type (i.e., chrysotile, amphibole, or mixed). For comparison, RRs were also calculated based on estimates from previous published meta-analyses [2–5].

Sensitivity analyses

The sensitivity of the predicted risk to the inclusion of specific studies was assessed with a 'jackknife' analysis, in which studies are excluded one by one [17]. The sensitivity of the predicted risk at low exposure to the inclusion of risk estimates corresponding to high exposures $(>100 \text{ f-y/ml})$ was assessed by fitting models excluding these data. In addition, results were stratified to studies that included a latency in their estimates between exposure and lung cancer and studies that included no latency.

⁶ The highest exposure category (corresponding to 2,400 f-y/ml) for which no risk estimate could be calculated was excluded from the analyses because of lack of observed cases The highest exposure category (corresponding to 2,400 f-y/ml) for which no risk estimate could be calculated was excluded from the analyses because of lack of observed cases

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Software

All analyses were performed with SAS version 9.2 (SAS Institute Inc., Cary, NC, USA). Significance tests between fiber-type-specific estimates were assessed with use of simulating the fiber-type-specific risk distribution in R version 2.10.1 (The R Foundation for Statistical Computing, Vienna, Austria).

Results

Study characteristics

Study characteristics of the 19 studies that were included in the meta-regression are listed in Table 1. The 104 risk estimates extracted from these 19 studies and corresponding CIs are shown in Fig. 1. From this figure, it is apparent that the risk estimates vary substantially even at the lower end of the cumulative exposure range. Of all risk estimates, 38 (37 %) were assessed at a cumulative exposure level of 40 f-y/ml or less, and 10 (10 %) at a cumulative exposure level of 4 f-y/ml or less.

Predictions

Table 2 shows the predicted risks based on the different exposure–response relationships. In all models, inclusion of cumulative exposure as an explanatory variable

significantly reduced model deviance. Compared to the linear model, a significantly better fit was observed for the model including a natural spline (for the explanatory variable) when a random intercept and slope was fitted (model 2A: deviance = 105.6, model 1A: deviance = 111.7; χ^2 test (1 *df*), $p = 0.01$). The natural spline suggested a nearly linear increase in the relative lung cancer risk at low levels as a function of exposure (Fig. 2). The slope slightly decreased after exposure of 150 f-y/ml. Based on this model, the relative lung cancer risk for 4 and 40 f-y/ml was estimated to be 1.502 (95 % CI: 1.173–1.922) and 1.680 (95 % CI: 1.317–2.142), respectively. After correction for the common estimated intercept, these RRs were $RR = 1.013$ for 4 f-y/ml and $RR = 1.133$ for 40 f-y/ml. Similarly, when fitting regression models without intercept, a significantly better fit was observed for the spline model over the linear model (χ^2 test (1 *df)*, $p = \langle 0.001 \rangle$. The RR that we predicted based on the natural spline model (model 2B) was 1.027 (95 % CI: 1.020–1.034) for 4 f-y/ml and 1.301 (95 % CI: 1.215–1.392) for 40 f-y/ml cumulative exposure.

Sensitivity analyses

A jackknife analysis, leaving one study at a time, was applied to the natural spline with intercept model (model 2A). Exclusion of the Ontario study (#11) resulted in the highest slope estimates, whereas the Pennsylvania study (#17) resulted in the lowest. However, their influence on

Fig. 1 Scatter plot of the relative risk estimates and their 95 % confidence intervals extracted from the 19 studies included in the meta-regression (a full range of cumulative exposures and b lower range of cumulative exposure $<$ 50 f-y/ml)

Table 2 Comparison of predicted risk at different exposure levels

	Deviance $(df)^a$	Intercept ^b (95 $%$ CI)	RR 4 f-y/ml ^b (95 % CI)	RR 40 f-y/ml ^b (95 % CI)
Models ^c				
1A. Linear model	111.7(100)	1.580 (1.243–2.008)	$1.592(1.252 - 2.023)$	$1.701(1.338 - 2.164)$
Corrected for intercept		$1.000(0.787-1.271)$	$1.007(0.793 - 1.280)$	$1.077(0.847 - 1.370)$
1B. Linear model without intercept	806.9 (102)	1.000 (1.000-1.000)	$1.017(1.009-1.024)$	$1.182(1.096-1.274)$
2A. Natural spline	105.6(99)	$1.483(1.157-1.900)$	$1.502(1.173 - 1.922)$	$1.680(1.317 - 2.142)$
Corrected for intercept		1.000 (0.780–1.281)	$1.013(0.791 - 1.296)$	$1.133(0.888 - 1.444)$
2B. Natural spline without intercept	703.6 (101)	1.000 (1.000-1.000)	$1.027(1.020-1.034)$	$1.301(1.215-1.392)$

RR relative risk, df degrees of freedom calculated as the number of data points minus the number of coefficients estimated

^a Fitted using ML estimation

^b Fitted using REML estimation

 \degree The deviance of the empty and intercept-only model was 3,433.9 and 309.9, respectively.

the predicted risks was negligible after correction for the intercept (data not shown). After correction for the intercept, the predicted risks increased most, and considerably, upon exclusion of the Quebec study (#1) from the analysis $(RR = 1.019$ for 4 f-y/ml and $RR = 1.211$ for 40 f-y/ml), and decreased most, but only slightly, upon exclusion of the South Carolina study (H4) (RR = 1.010 for 4 f-y/ml) and $RR = 1.103$ for 40 f-y/ml).

When models were fitted on exclusively risk estimates corresponding to exposures of 100 f-y/ml or less, a nonsignificant better fit was observed for the spline model as compared to a linear model (Supplementary data Table S1). Based on this sensitivity analyses, predicted risk ranged from $RR = 1.012$ for 4 f-y/ml to $RR = 1.152$ for 40 f-y/ml, which are comparable to the estimates based on the full range. Moreover, the predictive risk was about three times higher in studies that used a latency time of 10 years (RR = 1.030 for 4 f-y/ml and RR = 1.329 for 40 f-y/ml) compared to studies that used no latency $(RR = 1.012$ for 4 f-y/ml and $RR = 1.126$ for 40 f-y/ml) between lung cancer and exposure (Supplementary data Table S2, after correction for intercept).

Fiber type

After stratification of the results by fiber type, we observed a non-significant three- to fourfold higher combined RR for studies investigating exposure to mixed and amphibole fibers compared to studies investigating exposure predominantly to chrysotile fibers (Table 3, model 2A after correction for intercept). Additional analyses showed that these potency differences decreased to about twofold at higher exposures (Supplementary data Table S3, model 2A). The relative potencies across the exposure range are also shown in Fig. 3. When spline regressions were fitted without intercept (Table 3, model 2B), amphiboles had an 8–12-fold increased risk compared to chrysotile which was statistically significant. However, the exposure–response relationship based on the spline without intercept seems to be unrealistic and uncertain at higher cumulative exposures for amphiboles since the risk decreased after exposure of 150 f-y/ml (Fig. 3).

The predicted risk for chrysotile at low cumulative exposure ranges was heavily influenced by the Quebec and South Carolina studies. For exposures of 4 f-y/ml, the

RR relative risk

Fig. 3 Predicted exposure–response relationship over an exposure range of 0–400 f-y/ml stratified by fiber type (based on a spline regression model fitted with and without intercept)

exclusion of Quebec and the South Carolina study yielded a corrected predicted risk of 1.016 and 1.001, respectively, as compared to the overall estimate of 1.006. When both studies were excluded, this risk was estimated to be 1.004. The estimated risks for amphiboles were largely driven by the Wittenoom study (#6) and the New Jersey study (#7). When both studies were removed from the analyses, the risk for exposures of 4 f-y/ml dropped from 1.022 to 1.005. For mixed fibers, the predicted risk was most heavily influenced by the Belgian study (#14). Upon exclusion of the Belgian study, the risk increased from 1.018 to 1.027 for mixed exposures of 4 f-y/ml. Removing these five most influential studies from the analyses resulted in a 1.3–7 fold higher combined risk for studies investigating exposure to amphibole and mixed fibers compared to studies investigating exposure predominantly to chrysotile fibers.

Table 4 Overview of predicted risk based on previously published meta-analyses

RR relative risk, K_L the excess relative risk per unit of fiber year, PCM phase-contrast microscopy and measures fibers of longer than 5 μ m, thicker than approximately 0.25 μ m, and with an aspect (length-to-width) ratio >3 . TEM transmission electron microscopy

 a Estimates were based on a random effect model by combining K_L values that were derived by fitting an additive linear risk model with a variable intercept to each study.

 b Overall estimates were based on a random effect model by combining K_L values that were derived by fitting an additive linear risk model to</sup> each study (the K_L values and the intercepts were assumed to have a log normal distribution).

^c Estimates were based on exposure-risk relationships across cohorts by calculating an average exposure and an excess risk for each cohort.

 $d K_L$ values shown in the table are based on moderate or higher exposures. For low exposures, risks were calculated by applying the sub-linear model: RR = 1.6*cumulative exposure^1.3 for amphiboles and RR = 0.028*cumulative exposure^1.3 for chrysotile (as assessed by the authors for the best fitted model). No model for low exposures of mixed fibers was assessed.

^e Estimates were based by fitting K_L values and matching fiber type and size dimensions (as determined by TEM). The K_L values were derived by fitting an additive linear risk model with a variable intercept (with a maximum of $RR = 2$) to each study.

^f The K_L value based on PCM was assessed by fitting a metric with fibers of >0.2 µm width in which the relative potencies of long fibers versus short fibers and chrysotile versus amphibole were restricted to 1.

Comparison with risk estimates from other metaanalysis

An overview of the risk estimates based on previously published meta-analyses for cumulative exposure estimates of 4 and 40 f-y/ml is shown in Table 4. Our overall point estimates were higher compared to the risks that we calculated based on the meta- K_L value presented for the same 19 studies in the study of Lenters et al. [4]. Under a random linear effect model, they observed a K_L value (*100) of 0.13 (95 % CI: 0.04–0.22) with an intercept of 1.47 when all 19 studies were considered. Also, they showed that the meta- K_L value was higher for studies with a better exposure measurement strategy. When studies with two or more limitations in the exposure assessment component were excluded, their meta- K_L value was about two times higher. An ad hoc analysis showed that predictions based on our model also yielded higher risk estimates for studies with fewer limitations in the exposure assessment component (Supplementary data Table S4).

Estimates based on the overall meta- K_L value from the meta-analysis of Lash et al. [5] were similar albeit slightly lower. Overall estimates based on the analyses of Berman and Crump were comparable to our estimates [2]. However, they used a proxy for PCM measurements based on transmission electron microscopy (TEM), which complicates direct comparisons. If analyses would have been performed with results based on PCM measurements, their estimates would have been considerably lower (Supplemental Material, Table 6 of the meta-analysis of Lenters et al. [4]). Although we observed higher or comparable overall risks, we observed a much lower potency difference between amphiboles and chrysotile compared to those observed by Berman and Crump, and Hodgson and Darnton [2, 3].

Discussion

We used all available quantitative exposure–response data from observational epidemiological studies to assess the association between cumulative asbestos exposure and the risk of lung cancer. Our estimates for low-level exposures are of particular interest to predict the impact of exposures on individuals occupationally exposed to low levels and the general population. We estimated the RR for lung cancer to be 1.013 (95 % CI: 0.791–1.296) for 4 f-y/ml and 1.133 (95 % CI: 0.888–1.444) for 40 f-y/ml cumulative exposure. These predictions were based on a natural spline model that best fitted our data. When no intercept was fitted, significantly higher RRs were observed ranging from 1.027 (95 % CI: 1.020–1.034) for 4 f-y/ml to 1.301 (95 % CI: 1.215–1.392) for 40 f-y/ml. Our most conservative predicted risks were equal or higher than estimates based on additive linear relative risk models applied in previously published meta-analysis [4, 5]. Furthermore, our results indicated a moderately higher increased risk at low exposure in studies investigating amphiboles and mixed fibers compared to studies investigating chrysotile. These potency differences, however, were strongly influenced by a few studies. In general, we observed a lower potency difference between fiber types compared to those observed in previous meta-analyses [2, 3].

We adjusted the predicted estimates for the intercept. This assumes that the intercept fully represents a difference in baseline risk, and in practice, this may not be true. In fact, the observed intercepts suggest a very high excess risk (about 50 %) among workers compared to the general population that is attributable to other factors than asbestos exposure. Besides differences in risk factors between the exposed and unexposed population, systematic and random measurement errors can lead to an intercept greater than one [13]. In the study of Lenters et al. [4], a critical review

was performed on the quality of the exposure assessment methodology of the included studies. Here, it was shown that only a few studies had few limitations in the exposure assessment component and were of high quality. Furthermore, studies with lower quality had on average higher intercepts. Therefore, it is reasonable to assume that the observed intercept above $RR = 1$ is at least partly due to measurement error. Therefore, we also showed results of the natural spline fitted without intercept, since one might suggest that fitting a line through the origin $(ln(RR) = 0$ at zero exposure) would be more appropriate in the case of random measurement error. Our results also showed a lower intercept for the studies that included a latency time compared to those that did not include a latency time. Including no latency between exposure and lung cancer could also lead to measurement error in the exposure assessment when it incorrectly reflects the etiological time window of exposure.

Potential publication bias may have affected our risk estimates. However, this effect seemed minimal in previous analyses (see Supplemental Material of the study of Lenters et al. [4]). Moreover, the studies included in the meta-analyses differ from each other in terms of effect measure (ORs, RRs, and SMRs), place, time, follow-up, and exposure assessment. To limit the effect of differences across studies, our analysis are based on exposure–response relationship fitted within studies allowing for a random intercept and slope. However, we still observed a substantial variability in the exposure–response relationship which might be explained by various factors. For instance, the included studies might not have fully adjusted for differences in covariates between internal exposure categories. This may especially hold for studies in which the effect measure is adjusted only for age and gender. However, the effect of differences in background rates between exposed and non-exposed individuals on the exposure– response relationship can be addressed to a certain extend by adding an intercept to the model. Also, differences in smoking prevalence and mean length of follow-up per person could have resulted in different exposure–response relationships. The limited information available on these aspects did not allow further exploration of the potential bias they may incur.

The large heterogeneity between individual study results motivated the use of a random intercept and slope model consistent with previous meta-analyses [2–5]. The natural spline model provided the best fit to the data. After retransforming the results to the original scale, our results substantiated the evidence that the RR increases virtually linear with increasing exposure. Our findings are in contrast to data of Hodgson and Darnton suggesting a sublinear relationship [3]. One might also have expected a more supra-linear effect based on substantial high risks observed at very low exposures in a population-based study [7]. Although this population-based study was included in the current meta-regression, our results were statistically compatible with a more-or-less linear exposure–response model. The advantage, however, of our new method is that it provides a more accurate estimate of the lung cancer risk at low exposure since all available information could be used, and estimates did not need to be based on extrapolations below the study-specific exposure range. Moreover, our predictions are not heavily dependent on estimates at high exposure levels which are vulnerable to measurement error [18]. Substantially higher risk at low exposure has been observed in population-based studies with semiquantitative results [19, 20]. Although estimates from these studies are quite high, they are in the range of our results when we included only high-quality studies.

Low cumulative exposures are associated with all kind of occupations if duration of exposure is short. However, low cumulative exposures have been particularly observed in the general population due to downstream use of asbestos [7]. Like other meta-analyses, we could not determine whether risks might differ by exposure intensities, since intensities could mostly not be distinguished from reported cumulative exposures. Information on intensities is especially important if a threshold exists for asbestos-related lung cancer. However, no threshold of exposure intensity has been delineated for asbestos-related lung cancer. Moreover, a study by Frost et al. [21] showed that long-term asbestos removal workers had a significant increased risk of lung cancer compared to short-term workers indicating that cumulative exposure is an important measure if persons are exposed to low intensities.

The degree to which different types of asbestos have different potencies is a topic of ongoing debate [2, 22]. Berman and Crump showed a nine times higher increased risk for long amphiboles compared to long chrysotile fibers of all widths, and had even higher estimates for specific diameters (a ratio of 16:1 for long amosite versus long chrysotile for fibers with widths $\langle 4 \mu m \rangle$ [2, 11]. Hodgson and Darnton estimated the risk difference between chrysotile and amphibole fibers for lung cancer to be between 10 and 50 [3]. In the study of Lenters et al., a difference in risk ratio of a factor 8 was observed when all 19 studies were included (i.e., without adjusting for quality) [4]. In our analyses, we observed a non-significant three- to fourfold difference in potency between chrysotile and amphibole fibers.

Various explanations exist for the higher potency differences observed in previous meta-analyses compared to our results. Firstly, we used nonlinear regressions and estimated the overall slope from a distribution of study slopes. This resulted in shrinkage of study-specific slopes to the overall combined slope as well as less weight of point estimates at high exposures, and therefore, our analyses are less influenced by extreme results. Secondly, among the amphibole studies, we observed very high intercepts for the Wittenoom and New Jersey studies (i.e., intercepts of 2.8 and 3.8, respectively). These high intercepts were partly due to very high risks observed at relatively low exposures: the Wittenoom study observed a risk of 2.6 for 0.11 f-y/ml and the New Jersey study a risk of 2.8 for 3 f-y/ml. In the metaanalysis by Berman and Crump, these high intercepts were truncated at 2 [2, 12]. Therefore, our estimated risk for amphiboles is likely to be lower compared to risks estimated by Berman and Crump. When we fitted a natural spline without intercept, we observed a significant increased risk for amphiboles. In this case, the ratio of potency for amphobiles versus chrysotile was estimated to lie between 8:1 and 12:1, which was comparable to the ratios observed in the analyses of Berman and Crump for long fibers. However, the observed exposure–response relationship for amphiboles based on the spline without intercept was uncertain at higher cumulative exposure levels. Thirdly, Berman and Crump controlled for different fiber sizes in their meta-analysis [2]. Several studies showed that relatively longer and thinner fibers are stronger associated with lung cancer [2, 23, 24]. Since chrysotile fibers are generally longer and thinner than amphiboles, this might also explain the higher potency ratio between fibers types observed by Berman and Crump. Finally, Hodgson and Darnton used a different methodology to estimate the asbestos-related lung cancer risk [3]. They derived exposure-risk relationships across cohorts by calculating an average exposure and an excess risk for each cohort to avoid the effect of random measurement error. However, when for example, misclassification is more severe in lower exposure categories, the method applied does not necessarily completely eliminate the effect of exposure misclassification. Furthermore, mean levels do not reflect actual exposure levels accurately when observations are skewed. Also, it is expected that extraneous risk factors are differential distributed across study cohorts, which can have influenced their results.

From our results, it was apparent that the Quebec mine study and South Caroline textile study had a significant impact on the risk estimates for chrysotile. Upon removing the Quebec study, the RR for chrysotile increased considerably, whereas the exclusion of the South Carolina resulted in lower risks for chrysotile. The combined estimate of the three other studies involving chrysotile exposure also showed relatively low risks. The differences between the Quebec mine and South Carolina textile studies have been discussed extensively. A recent study by Berman concluded that the characteristics of the fiber can potentially explain the differences in lung cancer potency observed between these cohorts [25]. In that study, it was shown that the South Carolina textile workers were exposed to longer asbestos structures compared to the

Quebec miners and millers. The PCM-counted structures in textile factory dusts were virtually 100 % asbestos and 100 % asbestiform. In contrast, at least one-third of the structures counted by PCM in chrysotile mine and mill dusts were not asbestos. Additional limitations of PCM measurements have been discussed elsewhere [26]. Interestingly, the South Carolina study was classified as one with no limitations and the Quebec study as one with several limitations in the exposure assessment component as assessed by Lenters et al. [4] They showed that better quality studies yielded higher meta-estimates. This pattern was also observed with our spline regression model suggesting that observed fiber-specific potency differences at low cumulative exposure might also be partly due to differences in quality. Moreover, Lenters et al. showed that when analysis is restricted to only studies with few quality limitations of the exposure assessment component, the epidemiological evidence base is too sparse to draw deductions about potency differences per fiber type. Therefore, in light of the quality, we could not easily ascertain the magnitude of the potency differences between different fibers at low cumulative exposure.

Conclusion

Our results showed relative lung cancer risks for asbestos exposures of 4 and 40 f-y/ml to be between 1.013 and 1.027, and 1.13 and 1.30, respectively. Although we could not unequivocally determine potency differences between different fiber types at very low exposure levels of asbestos, the collected evidence suggests a threefold difference in risk between chrysotile and amphibole asbestos. This potency difference was not significant and lower than the generally held consensus. The flexible spline regression model we applied indicated that for low cumulative exposures, the increase in relative risk of lung cancer due to asbestos exposure may be larger than expected from previous results. This would suggest that, in general, a larger fraction of lung cancer incidence may be attributable to (many individuals having) relatively low cumulative exposure levels than previously estimated and might have important implications in developed nations. Additional research is required, in particular among removal workers and the general population in developed countries or low-exposed industrial cohorts using quantitative estimates of asbestos exposure, to further substantiate this notion.

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Conflict of interest The authors declare that they have no conflict of interest.

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