

Modeling Historic Arsenic Exposures and Spatial Risk for Bladder Cancer

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Abstract

Arsenic is a bladder carcinogen though less is known regarding the specific temporal relationship between exposure and bladder cancer diagnosis. In this study, we modeled time-varying mixtures of arsenic exposures at many historic temporal windows to evaluate their association with bladder cancer risk in the New England Bladder Cancer Study. We used arsenic exposure estimates up to 60 years prior to study entry and compared the goodness of fit of models using these mixtures to those using summary measures of arsenic exposures. We used the Bayesian index low rank kriging multiple membership model (LRK-MMM) to estimate the associations of these mixtures with bladder cancer and estimate cumulative spatial risk for bladder cancer using participants' residential histories. We found consistent evidence that modeling arsenic exposures as a time-varying mixture provided better fit to the data than using a single arsenic exposure summary measure. We estimated several positive though not significant associations of the time-varying arsenic mixtures with bladder cancer having odds ratios (ORs) of 1.03-1.14 and identified many significant and positive associations for an interaction among those who consumed water from a private dug well (ORs 1.28-1.60). Arsenic exposures 40-50 years before study entry received elevated importance weights in these mixtures. Additionally, we found two small areas of elevated cumulative spatial risk for bladder cancer in southern New Hampshire and in south central Maine. These results emphasize the importance of considering time-varying mixtures of exposures for diseases with long latencies such as bladder cancer.

Keywords Arsenic \cdot Mixture analysis \cdot Spatial analysis \cdot Residential history \cdot Bladder cancer \cdot Bayesian

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1 Introduction

Bladder cancer is a relatively common cancer and particularly so in industrialized nations [1]. It is expected to be the fourth-leading cancer in terms of incidence and eighth in terms of mortality among males in the United States in 2023 [2]. While the primary risk factor for bladder cancer is tobacco smoking [3], environmental and occupational exposures are other important risk factors, the latter of which accounts for 20–25% of all bladder cancer cases in men [4]. One notable environmental exposure for bladder cancer is arsenic, a naturally occurring metalloid that has been established as a bladder carcinogen by the International Agency for Research on Cancer [5]. Humans are exposed to arsenic primarily through drinking water in addition to exposure from contaminated soil and air.

Several studies have examined the arsenic-bladder cancer relationship across a variety of settings and study designs. For example, a large prospective cohort study in Chile identified significant relative risks for bladder cancer among residents of a region whose drinking water had high levels of arsenic [6]. Additionally, the New England Bladder Cancer Study (NEBCS) was a population-based case-control study conducted in Maine, New Hampshire and Vermont that was designed to identify the reasons for the elevated bladder cancer mortality rates in northern New England since at least the mid-twentieth century [7]. Arsenic was a natural candidate risk factor to investigate in this region, as its geology leads to low-to-moderate levels of arsenic in well water and a large proportion of the population use private wells as their primary source of drinking water. The NEBCS found a statistically significant trend in risk occurring with increasing drinking water intake from all sources and among participants with exclusive use of shallow dug wells before 1960 [8]. Further, lagging exposure 40 years led to stronger estimated associations with bladder cancer for cumulative arsenic exposure (average arsenic concentration multiplied by daily intake multiplied by duration of exposure). This long latency period was similar to that estimated in the Chile study [6], suggesting that evaluating arsenic exposures at different times may be informative for understanding its temporal relationship with bladder cancer.

Two classes of statistical techniques—mixture analysis and spatial analysis—can address this accumulation of exposures over space and time. Mixture analysis has emerged as a powerful technique to assess the effects of mixtures of exposures on health outcomes. This class of methods evaluates how the totality of a mixture, often comprised multiple chemicals, may be associated with an outcome, in contrast to analyzing the chemicals separately. This is in line with the exposome concept [9, 10], which holds that health outcomes result from a variety of exposure sources accumulating over the individual life course. Five methods for mixture analysis that have been developed in recent years are weighted quantile sum (WQS) regression [11, 12], Bayesian kernel machine regression (BKMR) models [13, 14], quantile g-computation [15], functional logistic regression [16], and the Bayesian group index model [17]. While these methods vary regarding implementation and estimation details, they share a goal of estimating the health effect of a mixture of environmental exposures as well as how each exposure varies in its contribution to the mixture effect. Additionally, mixture analysis can consider the timing of mixture exposures in relation to health outcomes. Recent applications of WQS and the Bayesian index model have assessed how exposure to mixtures of metals at different times is associated with rapid visual processing in children [18] and how exposure to neighborhood deprivation at historic time lags is associated with risk for non-Hodgkin lymphoma [19], respectively. These methods have extended distributed lag models [20, 21] to identify critical temporal exposure windows for mixtures in contrast to single chemicals. Identifying important temporal windows from time-varying mixtures provides additional information regarding how exposures influence health over time.

Statistical analyses of diseases with long latencies such as bladder cancer have begun to use residential histories to assess how cumulative spatial exposure to potentially unmeasured risk factors may also associate with disease risk. This approach embodies the exposome framework as well by considering accumulating unmeasured exposures for which residential histories are proxies. For example, the convolution multiple membership model (MMM) aggregates study participants' residential histories to larger administrative units and has been applied to model cumulative spatial risk for mesothelioma [22]. Also, the low-rank kriging (LRK) MMM uses the precise points in space from residential histories to estimate cumulative spatial risk and has been applied to case-control analyses of bladder cancer [23] and non-Hodgkin lymphoma (NHL) [24, 25]. The applications of this model to the NHL data also included weighted indices of environmental exposures to estimate mixture effects, the importance of each component in the mixture, and cumulative spatial risk simultaneously in a model termed the Bayesian index LRK-MMM. These models can identify geographic areas where residence is associated with elevated risk for disease and can motivate follow-up investigations into the causes of such risk, such as previously unmeasured environmental exposures.

Following the above developments in mixture and spatial modeling, we conducted an analysis in the NEBCS using historic arsenic exposure measurements and residential histories to estimate the associations between arsenic and bladder cancer risk over time. We used the Bayesian index LRK-MMM owing to its ability to flexibly estimate mixture effects and importance weights for components in the mixture and also estimate cumulative spatial risk for disease with residential histories. A primary goal of our analysis was to assess whether modeling arsenic exposures as a time-varying mixture provided better model fit to NEBCS data rather than doing so with a similar summary measure [8]. Our hypothesis was that modeling time-varying arsenic exposures as a weighted index would provide better model fit. We considered many different annual arsenic exposures up to a 60-year period prior to study entry, which would provide insight regarding the importance of arsenic exposures at different times for bladder cancer risk. We also estimated the residual cumulative spatial risk surface to evaluate whether any areas in the study region conferred significantly elevated cumulative spatial risk for bladder cancer that was unexplained by arsenic exposures or other covariates.

2 Methods

2.1 Study Population

The New England Bladder Cancer Study (NEBCS) is a population-based case-control study in Maine, New Hampshire, and Vermont. Rates of bladder cancer incidence and mortality have been elevated in this region for decades, and the NEBCS sought to identify risk factors responsible for the elevated incidence. The details of this study have been described previously [8, 26]. Briefly, cases were all newly diagnosed cases of bladder between 2001 and 2004 cancer among residents of the study region, and controls were randomly selected from driver's license registration (for controls under 65 years) or Centers for Medicare and Medicaid Services (for controls 65–79 years) records. Controls were frequency matched to cases by state, sex, and approximate age at diagnosis. There was a 65% participation rate in the study among both cases and controls. Residential histories were collected from participants based on in-person interviews with each subject that used a standardized questionnaire. In our analysis, we used the residential histories of long-term residents who had been living in Maine, New Hampshire, or Vermont for a minimum of 25 years prior to diagnosis (reference date for controls) (500 cases and 602 controls, which was 43% of the interviewed and eligible totals for both cases and controls), and considered residential locations for subjects from 1970 to 1986, to reflect a latency period that has been estimated for bladder cancer occupational carcinogens [27]. We made this restriction to focus on historic exposures within the study region. We adjusted our models for smoking status (former, current/occasional, don't know, versus reference never), high-risk occupation [28] (ever/ never), sex, age group (<55 or 55–64 or 65–74 or 75+), race, French-Canadian ancestry, ethnicity (Hispanic or don't know versus not Hispanic), educational attainment (high school degree, vocational or some college, college degree, postgraduate versus less than high school degree), cumulative total trihalomethanes (THM) intake from age 15+, average daily nitrate intake from public water supplies and private wells in 1970 or later, and drinking from a shallow dug well in the study area before 1960. Arsenic exposures were estimated at participant residential locations using drinking water sampling and statistical modeling, the details of which have been described previously [29]. In Table 1 we summarize the characteristics of the analysis population, as well as of a subset that had complete arsenic exposure measurements for the 60 years prior to study entry. This involved first excluding 85 cases and 95 controls who were less than 60 years old and then excluding 279 cases and 350 controls without complete estimated arsenic exposures over this time period, meaning that 33% of cases and 31% of controls who were 60 years or older had complete arsenic exposure histories. Overall, this sub-sample comprised 27% of case long-term residents and 26% of control long-term residents.

2.2 Model Specification

We considered three classes of models to assess the associations of arsenic exposure and cumulative unmeasured exposures with bladder cancer risk using the Bayesian

Variable	Long-term residents		Participants > = 60 years of age with complete arsenic measurements	
	Cases	Controls	Cases	Controls
	(N=500)	(N=602)	(N=136)	(N=157)
Study state				
Maine	260 (52)	336 (56)	80 (59)	96 (61)
Vermont	78 (16)	87 (14)	19 (14)	23 (15)
New Hampshire	162 (32)	179 (30)	37 (27)	38 (24)
Sex				
Female	127 (25)	159 (26)	57 (42)	71 (45)
Male	373 (75)	443 (74)	79 (58)	86 (55)
Age group				
< 55 years	42 (8)	49 (8)	0 (0)	0 (0)
55–64 years	126 (25)	141 (23)	33 (24)	33 (21)
65-4 years	203 (41)	253 (42)	74 (54)	79 (50)
>75 years	129 (26)	159 (26)	29 (21)	45 (29)
Race				
White	471 (94)	578 (96)	126 (93)	150 (96)
Native American/White only	26 (5)	19 (3)	9 (7)	6 (4)
Other	3 (1)	4 (1)	1 (1)	1(1)
Don't know	0 (0)	1 (0)	0 (0)	0 (0)
Hispanic ethnicity				
No	496 (99)	592 (98)	134 (99)	155 (99)
Yes	4 (1)	9 (1)	2 (1)	2 (1)
Don't know	0 (0)	1 (0)	0 (0)	0 (0)
Education				
<high degree<="" school="" td=""><td>123 (25)</td><td>122 (20)</td><td>39 (29)</td><td>46 (29)</td></high>	123 (25)	122 (20)	39 (29)	46 (29)
High School degree	189 (38)	229 (38)	66 (49)	77 (49)
Vocational	40 (8)	52 (9)	8 (6)	10 (6)
Some college	60 (12)	76 (13)	12 (9)	14 (9)
College graduate	52 (10)	57 (9)	11 (8)	9 (6)
Postgraduate	36 (7)	66 (11)	0 (0)	1 (1)
Smoking history				
Never smoked	77 (15)	207 (34)	22 (16)	71 (45)
Former smoker	273 (55)	306 (51)	71 (52)	70 (45)
Current/occasional smoker	149 (30)	89 (15)	42 (31)	16 (10)
Don't know	1 (0)	0 (0)	1 (1)	0 (0)
Ever held high risk occupation				
No	233 (47)	406 (67)	64 (47)	115 (73)
Yes	261 (52)	189 (31)	68 (50)	39 (25)
Never worked	6(1)	7(1)	4 (3)	3 (2)

Table 1 Summary of NEBCS analysis population

Variable	Long-term res	Long-term residents		Participants > = 60 years of age with complete arsenic measurements	
	Cases	Controls (N=602)	Cases (N=136)	Controls $(N=157)$	
	(N=500)				
Cumulative lagged arsenic in	take ^a				
<3.52 mg	81 (16)	114 (19)	18 (13)	15 (10)	
(3.52, 8.77] mg	118 (24)	136 (23)	28 (21)	33 (21)	
(8.78, 22.42] mg	110 (22)	156 (26)	36 (26)	50 (32)	
(22.42, 83.53] mg	119 (24)	128 (21)	41 (30)	48 (31)	
(83.53, 124.80] mg	17 (3)	14 (2)	5 (4)	5 (3)	
>124.80 mg	26 (5)	18 (3)	8 (6)	6 (4)	
Missing	29 (6)	36 (6)	0 (0)	0 (0)	
Average arsenic concentration	1 ^b				
<0.53 ug/L	153 (31)	165 (27)	46 (34)	45 (29)	
(0.53, 0.95] ug/L	85 (17)	138 (23)	24 (18)	43 (27)	
(0.95, 2.06] ug/L	121 (24)	126 (21)	35 (26)	35 (22)	
(2.06, 6.99] ug/L	94 (19)	108 (18)	24 (18)	28 (18)	
> 6.99 ug/L	22 (4)	36 (6)	7 (5)	6 (4)	
Missing	25 (5)	29 (5)	0 (0)	0 (0)	
French-Canadian ancestry					
No	328 (66)	414 (69)	86 (63)	110 (70)	
Yes	172 (34)	188 (31)	50 (37)	47 (30)	
Dug well drinker ^c					
No	334 (67)	409 (68)	82 (60)	96 (61)	
Yes	141 (28)	164 (27)	54 (40)	61 (39)	
Missing	25 (5)	29 (5)	0 (0)	0 (0)	

Table 1 (continued)

Quantities presented in table are frequency (percentage). Some percentages may not sum exactly to 100 due to rounding

^aFrom residential and workplace water, lagged 40 years

^bFrom residential and workplace water

^cEver drank from unconsolidated well at home or work in study area before 1960

index low-rank kriging multiple membership model [24] (LRK-MMM), a type of hierarchical Bayesian regression model. We chose to use this model due to its ability to accommodate measured mixtures and cumulative spatial risk using residential histories. The three classes of models we used varied in their specification and timing of arsenic exposures and are given below. For all models we assumed a binary outcome variable Y_i denoting case membership that was distributed as a Bernoulli random variable with probability of being a case p_i , and we modeled the log-odds of the probability of being a case.

In Class 0,

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 * \operatorname{arsenic}_i + \sum_{b=1}^B \theta_b x_{ib} + \sum_{j \in A_{(i)}} w_{ij} \sum_{m=1}^{n_K} \psi_m C\left[\left|s_{ij} - k_m\right|/\rho\right]$$

In Class 1,

$$\log\left(\frac{p_{i}}{1-p_{i}}\right) = \beta_{0} + \beta_{1} * \sum_{t=1}^{T} \omega_{t}q_{it} + \sum_{b=1}^{B} \theta_{b}x_{ib} + \sum_{j \in A_{(i)}} w_{ij} \sum_{m=1}^{n_{K}} \psi_{m}C\Big[\Big|s_{ij} - k_{m}\Big|/\rho\Big]$$

In Class 2,

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \left(\beta_1 + \beta_2 * \text{DugWell}_i\right) * \sum_{t=1}^T \omega_t q_{it} + \sum_{b=1}^B \theta_b x_{ib} + \sum_{j \in A_{(i)}} w_{ij} \sum_{m=1}^{n_k} \psi_m C\Big[\left| s_{ij} - k_m \right| / \rho \Big]$$

Several components are common to models. Namely, β_0 is an intercept term, and we control for covariate vector x_i with coefficient vector θ . Additionally, we estimate cumulative spatial risk with the right-most term in all models. For the *i*th participant, define the set of all locations they have lived in their residential history to be $A_{(i)}$. For *j*th location s_{ij} , the spatial risk they experience at that location is estimated with a sum of spatial random effects ψ weighted by the spatial covariance $C[\cdot]$ between the residential location and the knot locations where the spatial random effects are evaluated $(k_m, m = 1, ..., n_K)$. We use the Matern covariance function that simplifies to $C[d] = (1 + d/\rho)e^{-d/\rho}$ for distance *d* and spatial range parameter ρ when fixing parameters of the Matern family of *m* and ν to 1 and 3/2, respectively. We choose to use the Matern family of covariance functions in our models due to its popularity in geostatistical models [30, 31], flexibility, and smoothness with respect to distance. Finally, the term w_{ij} represents the proportion of the study period that the *i*th participant resided at their *j*th location, so $\sum_{i=1}^{J} w_{ij} = 1$.

The specification of the arsenic term is different in each class of models. In Class 0, we estimate the association of a summary arsenic exposure (cumulative arsenic intake from residential and workplace water lagged 40 years (meaning a sum of these arsenic exposures ending 40 years before study entry), or average arsenic concentration from residential and workplace water) with bladder cancer risk using regression coefficient β_1 . In Class 1, we replace the summary arsenic exposure with a time-varying mixture of arsenic exposures. Over a time period of T years, the quantized arsenic exposure for the *i*th individual in the *t*th year is q_{it} , t = 1, ..., T. We quantized the arsenic exposures to accommodate high correlations between arsenic measurements over time and to account for uncertainty in the measurements, using quartiles (four groups) in all models. The importance weight for the tth year in the mixture was ω_t , and $\sum_{t=1}^T \omega_t = 1$ for interpretability. We used a variety of windows of temporal exposure for the arsenic mixture (Table 2). In Class 2, we used the same set of temporal windows as in Class 1 but added an additional interaction term ($\beta_2 * \text{Dug Well}_i$) to evaluate how the health effect of the arsenic mixtures could vary for study participants who had consumed water from shallow dug wells. We included this class of models due to a previous finding that water intake was significantly associated with bladder cancer risk among study participants who drank

Model	Exposure years	Number of weights
1, 2, 3, 4	[60,0)	60, 30, 20, 15
5, 6, 7	[60,15)	45, 15, 9
8, 9, 10	[60, 30)	30, 15, 10
11, 12	[60, 45)	15, 5
13, 14, 15	[45, 0)	45, 15, 9
16, 17, 18	[30, 0)	30, 15, 10
19, 20	[15, 0)	15, 5

In the Exposure Years column, the pair [x, y) denotes arsenic exposures beginning x (inclusive) and ending y (exclusive) years before study entry. If the number of weights ν in a given model is less than x - y, the first entry in the exposure index is the median of the exposures in years $x, \ldots, x - (x - y)/\nu + 1$, and so on

from shallow dug wells prior to 1960 [8]. Exponentiating the regression coefficients β_{i} , j = 1, 2 gives the odds ratio for bladder cancer for the arsenic exposures.

Some study participants had missing arsenic exposure estimates at certain time lags, and increasingly so many decades prior to study entry. In the case of missing arsenic exposures, we imputed missing values from a log-normal distribution using the non-missing logged estimated arsenic measurements in the given year. In a sensitivity analysis, we fit all models described above to a subset of the analysis sample who had complete arsenic exposure estimates for the 60 years prior to study entry. We conducted the sensitivity analysis to evaluate the impact of imputing missing arsenic measurements on model estimated odds ratios for the arsenic mixtures and on the importance of historic arsenic exposures.

2.3 Knot Selection

The Bayesian index LRK-MMM reduces the dimensionality of the spatial risk model component through a set of n_K knots, which are locations where the spatial random effects are estimated. This requires specification of the number and location of the knots. Previous research has suggested that the Teitz and Bart heuristic [32], originally developed to address the location-allocation problem, chooses knot locations in a way that enables good sensitivity and power to detect regions of elevated spatial risk for disease [33]. Therefore, we use this heuristic to choose knot locations in our models. Briefly, Teitz and Bart begins with a random set of knot locations and iteratively changes knot location points to candidate ones if doing so decreases the objective function of the total distance between cases and their nearest knot location. Additionally, we use $n_K = 60$ knots in all models for comparability with a previous LRK-MMM analysis of NEBCS data [23].

 Table 2
 Summary of temporal

 windows and number of weights
 for arsenic exposure index

We fitted models in a Bayesian framework using Markov chain Monte Carlo (MCMC) methods. We specified the following priors: for the adjustment covariates, $\theta_b \sim Normal(0, \tau_b)$, where $\tau_b = 1/\sigma_b^2$ and $\sigma_b \sim Uniform(0, 100)$. The intercept and arsenic exposure coefficients received a similar prior $\beta_j \sim Normal(0, \tau_j)$, with $\tau_j = 1/\sigma_j^2$ and $\sigma_j \sim Uniform(0, 100), j = 0, 1, 2$. The arsenic index importance weight vector $\boldsymbol{\omega}$ received a Dirichlet prior with parameter vector $\boldsymbol{\alpha} = (\alpha_1, \dots, \alpha_T)$ to assure that each weight in the index was between 0 and 1 and $\sum_{t=1}^{T} \omega_t = 1$. The spatial random effect vector $\boldsymbol{\psi}$ received a multivariate normal prior $MVN(0, \tau_s \Omega^{-1})$, with covariance matrix $\boldsymbol{\Omega} = [C[|k_a - k_b|/\rho]], \tau_s = 1/\sigma_s^2$, and $\sigma_s \sim Uniform(0, 100)$. Finally, the spatial range parameter received a uniform prior on the range $(0, d_{max})$, where d_{max} represents the maximum distance between a knot location and a residential location.

For model estimation, we used Just Another Gibbs Sampler (JAGS) [34], in the software R, version 4.1.0 [35], using two chains that each had a burn in period of 60,000 iterations and retained 40,000 iterations for sampling from the joint posterior distribution. We monitored convergence of model parameters using the Gelman-Rubin statistic [36], where a parameter was considered to have converged if its statistic was less than 1.1, using the coda package [37] in R. We compared model goodness of fit using the deviance information criterion (DIC), which is a common method to compare model fit that penalizes model complexity [38]. Smaller DIC values indicate a better fit to the data, and differences of greater than 5 in DIC may indicate meaningfully better fitting models [39]. We summarized associations with the arsenic exposures with posterior mean and 95% credible interval for the odds ratio.

We assessed spatial risk over a 6 km by 6 km grid covering the study region, predicting spatial risk at each grid cell with the posterior estimates of the spatial random effects at the knot locations and the covariance function between the grid cell and the knot locations. We identified grid cells as being significantly elevated or lowered in risk using exceedance probabilities [40], which estimate how frequently the spatial odds at a location (θ_i) exceed the null value ($\theta_i = 1$). The exceedance probabilities use the posterior distribution of spatial odds at the *i*th location ($\theta_{i,m+1}, \ldots, \theta_{i,m+G}$), where *m* represents the burn-in and *G* represents the number of posterior samples after the burn-in, and are calculated as $\hat{q}_i = 1/G \sum_{g=m+1}^{m+G} I(\theta_{i,g} > 1)$. We determined significance of spatial risk using 95% exceedance probabilities.

3 Results

According to the DIC values, modeling the effect of arsenic exposure as time-varying rather than as a single summary measure consistently provided better model fit to the data (Table 3). Where the DIC values for the Class 0 models with summary arsenic measures were 1439 and 1440, the DIC values for the Class 1 time-varying

Class 0		Class 1		Class 2	
Model	DIC	Model	DIC	Model	DIC
A	1439	1	1433	1	1428
В	1440	2	1431	2	1424
		3	1434	3	1429
		4	1433	4	1429
		5	1436	5	1434
		6	1434	6	1432
		7	1435	7	1428
		8	1433	8	1432
		9	1433	9	1432
		10	1434	10	1431
		11	1435	11	1431
		12	1432	12	1428
		13	1433	13	1429
		14	1433	14	1430
		15	1432	15	1431
		16	1432	16	1430
		17	1433	17	1429
		18	1433	18	1429
		19	1433	19	1427
		20	1433	20	1427

DIC stands for Deviance Information Criterion, where smaller values indicate better-fitting models. In Class 0 that does not fit a time-varying arsenic exposure index, Model A and B use summary arsenic exposures cumulative arsenic intake from residential and workplace water lagged 40 years and average arsenic concentration from residential and workplace water, respectively

arsenic exposure index models were between 1431 and 1436. Therefore, 14 of the 20 Class 1 models provided considerably better model fit than the Class 0 models. Including the additional interaction term for the dug well drinkers in the Class 2 models provided even better model fit. The DIC values for models in Class 2 ranged from 1424 to 1434, meaning that 19 of the 20 models fit considerably better than the Class 0 models.

Comparing the estimated odds ratios for the arsenic exposure index provides additional insight into the health effect of arsenic exposure at different times on bladder cancer risk. In the Class 1 models, the odds ratio for the arsenic exposure index ranged from 1.03 to 1.14 (Fig. 1). The four best fitting models in this class included arsenic exposures up to 30 years (one model), 45 years (one model), and 60 years (two models) prior to study entry. In the Class 2 models, the odds ratio for the arsenic index ranged from 1.28 to 1.60 among dug well drinkers and was significantly elevated for 16 of the 20 models (Fig. 2). These results provide consistent evidence for the association between historic time-varying arsenic exposures and

 Table 3
 Summary of model

 goodness of fit by model class



Fig. 1 Summary of model fit and arsenic index odds ratio, Class 1 models. Models are ranked from top to bottom indicating best to worst fit. DIC values are displayed in blue text. The posterior mean and 95% credible interval for the arsenic exposure index odds ratio are displayed in black text (Color figure online)



Fig. 2 Summary of model fit and arsenic index odds ratio for dug well drinkers, Class 2 models. Models are ranked from top to bottom indicating best to worst fit. DIC values are displayed in blue text. The posterior mean and 95% credible interval for the arsenic exposure index odds ratio among dug well drinkers are displayed in black text (Color figure online)

bladder cancer risk among those who drank water from shallow dug wells. The six best fitting models in this class included four models with arsenic exposures up to 60 years before study entry and two models with arsenic exposures in the 15 years prior to study entry.



Fig. 3 Summary of yearly importance weights among best fitting models in Classes 1 and 2. Class 1 and 2 models are in the left and right panel, respectively. The value on the y-axis represents the importance weight of the arsenic exposure at the given time lag. Weights in each index sum to one

Evaluating the estimated importance weights in different indices supports the variation in the importance of arsenic exposures at different times for bladder cancer risk (Fig. 3). In the Class 1 models, there were several elevated importance weights at time lags between 60 and 45 years before study entry for several models, and in three of the four models the largest estimated importance weight corresponded to arsenic exposure at least 45 years prior to study entry. For the Class 2 models, there were many elevated importance weights between 60 and 45 years prior to study entry as well as a few elevated importance weights in the 10 years before study entry.

The sensitivity analysis found that results changed little when including only study participants having 60 years of arsenic exposures. Relative to the best fitting Class 0 models, the Class 1 models provided improvements in DIC of 10–22, and the Class 2 models provided improvements of 14–33 (Supplemental Material Table S1). In the Class 1 models, the estimated odds ratio for the arsenic exposure index was close to the null and not significant for any model (Supplemental Material Fig. S1). In the Class 2 models, the estimated odds ratio for the arsenic exposure index among dug well drinkers was elevated for all models and significant for 16 of 20 models (Supplemental Material Figure S2). Therefore, the sensitivity analysis suggests that imputing missing arsenic exposures did not bias conclusions in the main analysis.

After modeling the time-varying arsenic exposure mixtures and adjusting for individual-level covariates, there was little additional variation in the cumulative spatial risk surface. In the main analysis, nowhere in the study region constituted a clustering of significantly elevated cumulative spatial risk. In the Class 1 models from the sensitivity analysis, three models identified a small circular region of elevated risk in southern New Hampshire near Rochester that had approximate



Fig. 4 Class 1 models finding significantly elevated cumulative spatial risk for bladder cancer, only using data from long-term residents with 60 years of arsenic exposures. Red grid cells denote areas with significantly elevated cumulative spatial risk (Color figure online)



Fig. 5 Class 2 models finding significantly elevated cumulative spatial risk for bladder cancer, only using data from long-term residents with 60 years of arsenic exposures. Red grid cells denote areas with significantly elevated cumulative spatial risk (Color figure online)

radius 15 km, and one of these models identified a region of similar size in south central Maine near Augusta (Fig. 4). In the Class 2 models from the sensitivity analysis, five models identified the region in southern New Hampshire, and four of these models identified the region in south central Maine (Fig. 5).

4 Discussion/Conclusion

In this study, we estimated the associations between mixtures of historic arsenic exposures and bladder cancer risk in the NEBCS. Our analytical approach used the Bayesian index LRK-MMM, which simultaneously estimates cumulative spatial risk using residential histories and mixture effects. We compared the goodness of fit of these models to those that used a 40-year lagged cumulative measure of arsenic exposure in addition to estimating cumulative spatial risk (standard LRK-MMMs). We found that modeling arsenic exposures as a time-varying mixture consistently provided better goodness of fit than using the summary measure. In these time-varying mixtures, we estimated many positive though not significant associations with bladder cancer risk. However, an interaction term for the arsenic exposure mixture among dug well drinkers was positively and significantly associated with bladder cancer risk in almost every model fit, which provides insight into how arsenic exposures affect cancer risk differentially over time and by source of drinking water. In these models, arsenic exposures between 45 and 60 years before study entry received large importance weights in the mixture, and a few models estimated larger importance weights for more recent exposures. These results communicate the complexity and time-varying nature of arsenic exposures for bladder cancer risk. Additionally, modeling time-varying arsenic exposures explained one area of elevated spatial risk for bladder cancer identified from a previous analysis that used LRK-MMMs and a summary measure of arsenic exposure [23]. We found that our results changed little when restricting the analysis sample to include only residents with complete arsenic exposure histories in contrast to the imputation we performed in the main analysis. We also identified small areas of elevated spatial risk for bladder cancer in the sensitivity that could warrant additional hypothesis generation and analyses.

Findings in our study contribute to the literature on arsenic and bladder cancer. Though arsenic has been recognized as a carcinogen for bladder cancer [5], less is known regarding the exact timing of the relationship between exposure and diagnosis. A previous analysis of arsenic in the NEBCS found that lagging arsenic exposure 40 years provided stronger and significant associations with bladder cancer risk compared with shorter exposure lags [8]. This latency was approximately equal to that found in the prospective study in Chile that identified significantly elevated risks for bladder cancer risks up to 40 years after exposure reduction [6, 41]. One notable difference between the sample in the Chile study and the NEBCS is that many residents in the former study were exposed to very high levels of arsenic in the earliest decades of the exposure range (1950s to 1970s). Contrastingly, most participants in the NEBCS were exposed to low-tomoderate levels of arsenic that were lower than for many participants in the Chile study. The large estimated importance weights in many of our models for arsenic exposures occurring 45–50 years before study entry (Fig. 3) suggest a somewhat longer latency period for these more moderate exposures.

There are several strengths to our study. First is its use of complete residential histories and historic arsenic exposure estimates [29] that enable assessment of how historic unmeasured and estimated exposures respectively influence risk for bladder cancer, a disease with a long latency period. Modeling exposure over the life course requires [42, 43] a large effort. Our results illustrate the importance of estimating these historic exposures. Additionally, our analytical framework employed the Bayesian index LRK-MMM, which developed from the Bayesian index model and has demonstrated the ability to accurately and consistently estimate mixture effects as well as high power to detect regions of elevated unmeasured spatial risk for disease [17, 24]. Finally, we considered a wide range of exposure windows for the time-varying arsenic mixtures to assess how exposures at different times vary in importance for explaining bladder cancer risk. The methods we used in this analysis can motivate future studies that model the associations of time-varying exposures with disease risk.

The limitations of our study should also be considered. First, although the historic arsenic exposure measurements were a product of extensive water sampling, characterization of the geology of the aquifers, and statistical modeling, there is a possibility that the arsenic estimates may not have represented the true exposures of some participants, and particularly for certain historical residential locations. For a subset of drilled wells, this method of arsenic exposure assignment offered only moderate agreement, sensitivity, and specificity for a binary classification of less or greater than 2 µg/L of arsenic [29]. Thus, it is possible that the variation in the arsenic exposure estimates could have led to non-differential exposure misclassification and a subsequent underestimation of true risk [44]. Second, the prevalence of use of arsenical pesticides and dug wells decreased from the mid-twentieth century to the present [8, 45]. Therefore, these factors may be less relevant for future studies and populations. Third, it is possible that some other water contaminants demonstrated different time-varying associations and strengths of association with bladder cancer risk. A previous analysis of data in this study found that high exposures to drinking water nitrates (average concentration above the 95th percentile) was associated with elevated risk for bladder cancer [46]. It is possible that exposures at different times to this contaminant could be important for bladder cancer risk. Fourth, nonparticipation occurred in this study, with a 65% response rate for both case and control subjects. Therefore, it is possible that selection bias could have impacted our findings. However, at least regarding rurality and well use, this does not appear to have been an issue in this study. For both cases and controls there were a similar proportion of participants and nonparticipants living outside a designated census place, which is a common indicator for having a private well [8], so nonparticipation does not appear to have influenced the composition of the sample in regards to this important factor for bladder cancer in this region. Additionally, while this is not necessarily a limitation, we note that we used one statistical method—the Bayesian index LRK-MMM—given the nature of our data and goal to estimate mixture effects and spatial risk simultaneously. It is possible that other statistical methods could provide new insights into arsenic exposures and bladder cancer. Finally, we are unable to establish causality due to the retrospective nature of the study, and despite adjusting for many demographic, occupational, and environmental covariates, we cannot rule out the possibility of residual confounding.

In conclusion, we found that modeling historic arsenic exposures as a time-varying mixture provided better fit to the data than modeling a single summary lagged exposure in a population-based case–control study of bladder cancer in New England. We estimated significant and positive associations for arsenic mixtures and bladder cancer among dug well drinkers across several historic windows of exposure. We found evidence for the importance of arsenic exposures 40–50 years before study entry in these time window mixtures. Our results support the importance of historic environmental exposures particularly in the context of diseases with long latency periods. Although the prevalence of dug wells and arsenical pesticides have decreased over time, our results provide additional evidence for the latency period between arsenic and bladder cancer and motivate the use of time-varying mixtures in future health research. Future studies should continue to investigate how geospatial analysis and exposure assignment can benefit health research, particularly over historic time periods.

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