Modeling Linear Combinations of Multiple Pollutants

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Research Aims

We propose to develop a fully Bayesisan implementation of the *groupwise additive index model* (GAIM) (Masselot et al. 2020). Sequential quadratic programming is used in Masselot et al. (2020) to optimize the GAIM in the frequentist setting, but there is no discussion of a Bayesian implementation. Putting priors aside, for response distribution D from the exponential family with a d-dimensional parameter $\theta=(\theta_1,\dots,\theta_d)$ and canonical link function g, the GAIM is,

$$\begin{split} Y_t|\theta_t &= D(\theta), \\ g(\theta) &= X\beta + \sum_{i=1}^I s_i(\alpha_1\gamma_{1,t} + \ldots + \alpha_{n_i}\gamma_{n_i,t}) + \sum_{k=1}^K f_k(\eta_k) \end{split}$$

In the above, X is a regression matrix and β a vector of fixed effects. The second sum consists of K smoothing functions f_1,\ldots,f_K applied to K potential confounders $\eta_k=(\eta_1,\ldots,\eta_{n_k})$. The novelty lies in the first sum, which fits smooth functions s_i to a weighted sum of potentially time-varying covariates.

The GAIM is a flexible and extensible statistical model with a wide variety of potential applications, as the researcher can choose the number I of smooth functions s_i and the number of covariates n_i in each s_i . For example, the researcher may specify a model with one smooth function s_1 and two covariates, where $\gamma_{1,t}$ is the observed levels of the pollutant ozone (O₃) and $\gamma_{2,t}$ is the observed levels of particulate matter less than 2.5 μg (PM_{2.5}). In this case, the 2 weights give the *relative* contribution of O₃ and PM_{2.5} to Y_t , the health outcome of interest. If the researcher added 1-day lagged observations of O₃ and PM_{2.5}, the 4 weights would give the *relative* contribution of the 0-day and 1-day lagged O₃ and PM_{2.5} levels. If the researcher instead added three more pollutants, nitrogen dioxide (NO₂), sulfur dioxide (SO₂), and particulate matter less than 10 μg (PM₁₀), the 5 weights would give the *relative* contribution of each pollutant.

To explore the benefits the GAIM provides over other possible model specifications, we next compare it to a standard regression model and one where the s_i have more general form. To ease notation, let us assume there are no confounders and that we are only considering a linear combination of two terms, $\gamma_{1,t}$ and $\gamma_{2,t}$. Then, these three models have link functions,

$$g(\theta) = X\beta + \alpha_1\gamma_1 + \alpha_2\gamma_2 \tag{1}$$

$$g(\theta) = X\beta + s_1(\alpha_1\gamma_1 + \alpha_2\gamma_2) \tag{2}$$

$$g(\theta) = X\beta + s_1^*(\gamma_1, \gamma_2) \tag{3}$$

The model with link function (1) is a regression model with two fixed effects, α_1 and α_2 . The GAIM has link function (2) with weights α_1 and α_2 , while link function (3) specifies s_1^* as a 2-dimensional smoothing function. The GAIM is more flexible than a regression model, as the inclusion of s_1 allows it to capture nonlinearities in the relationship between $\gamma_{1,t}$, $\gamma_{2,t}$ and the outcome Y_t . Compared to the 2-dimensional smoothing function, the GAIM is less computationally demanding – it does *not* suffer from the curse of dimensionality. The linear combination $\alpha_1\gamma_{1,t}+\alpha_2\gamma_{2,t}$ is always 1-dimensional, and remains so regardless of how many terms are in its sum. This constrasts with s_1^* , whose dimensionality scales with its number of arguments. The second advantage of the GAIM is that it is more interpretable, as the *weights* α_1 and α_2 reflect the *relative* contribution of $\gamma_{1,t}$ and $\gamma_{2,t}$, respectively. In comparison, $s_1^*(\gamma_1,\gamma_2)$ has no such weights; indeed, s_1^* may not have *any* directly interpretable parameters.

We will develop a fully Bayesian implementation of the GAIM in Stan, a widely used statistical modeling language that allows for the rapid iteration and development of Bayesian models with inference performed using Hamiltonian Monte Carlo (Carpenter et al. 2017). No such Bayesian implementation currently exists; developing a robust implementation and releasing the software would be a significant contribution towards having accessible and interpretable dimension-reducing models. Developing a robust Bayesian implementation includes, among other things, identifying suitable priors for the weights α and suitable priors for imposing a variety of shape constraints, such as monotonicity and convexity, on the s_i . Kimeldorf and Wahba (1970) describes a natural correspondence between random walks and smoothing splines, such that random walks are the natural smoothing function for Bayesians. The second goal is to develop non-MCMC estimation methods, and to characterize the performance of this implementation under a variety of different different response distributions.

The Research Questions

We will demonstrate the computational and interpretation benefits of the GAIM in the context of the health effects of multiple air pollutants. Recent years have seen increased interest in modeling the joint effect of two or more pollutants in health outcomes (Dominici et al. 2010; Billionnet et al. 2012; Davalos et al. 2017); Bayesian approaches can be seen in (Blangiardo et al. 2019; Bobb, Dominici, and Peng 2013; Huang, Lee, and Scott 2018). We will consider two research questions in the course of our inquiry, namely,

- 1. What is the combined effect of multiple pollutants on various daily mortality outcomes?, and
- 2. What is the relationship between daily COVID-19 mortality and air pollution?

Question 1

The workhorse of the (predominantly frequentist) air pollution literature is the one-pollutant log-linear Poisson regression model. This model accounts for confounders using fixed effects and smooth functions, such as the natural cubic spline (Samet et al. 2000; Dominici et al. 2002; Liu et al. 2019). Let the average rate an outcome Y_t occurs on day t be denoted by λ_t . Then, a typical one-pollutant model is,

$$\begin{split} Y_t | \lambda_t &= \mathrm{Poisson}(\lambda_t), \\ \log(\lambda_t) &= X\beta + \gamma_1 P_{1,t} + \sum_{k=1}^K f_k(\eta_k). \end{split}$$

Here, Y_t is the health outcome of interest, such as respiratory mortality or morbidity. The design matrix X contains day-of-the-week effects and seasonal terms, and the $f_i(\eta_{k,*})$ are smooth functions of potential confounders such as time and temperature. An extension of this model would fit $P_{1,t}$ to its own smooth function s_1 . More generally, we could fit N pollutants to N smooth functions, s_1,\ldots,s_N . A third alternative, which we propose here, is to model N pollutants using the GAIM.

We have previously discussed the computational and interpretability advantages of the GAIM. For one, people are exposed to the mixture (a weighted sum) of *all* the pollutants present in their area. When extending the one pollutant model to account for this, it is important to note that common pollutants are highly correlated (Huang, Brown, and Shin 2020). When fitting a single pollutant model, health effects that are attributed to this one pollutant may very well be caused by another correlated pollutant(s). This has been observed in Franklin and Schwartz (2008), where the authors found that the effect of ozone

on non-accidental mortality was "substantially reduced" after adjusting for particle sulfate. In Liu et al. (2019), the authors found statistically significant differences in the percentage change in all-cause mortality attributable to $PM_{2.5}$ and PM_{10} when adjusting for either of NO_2 or SO_2 .

Therefore, it is is crucial to extend the one-pollutant model so that we may more accurately attribute the health effects to the *correct* pollutants. If we want to know the health impact of the mixture of pollutants being modeling, it is only natural to model the pollutants as a mixture. Therefore, the most informative extension of the one pollutant model is one that conducts inference on a mixture of pollutants in the model. Using a nonlinear function of the weighted sum has advantages, as there is evidence that some health outcomes are nonlinearly related to pollution levels (Feng et al. 2016), and that synergistic effects occur when more than one pollutant is present at higher levels (Xia and Tong 2006). We propose using the GAIM in the multiple pollutant setting because it is particularly well-suited to estimating nonlinear effects of weighted sums: the value of smooth function s_i is the (log) effect of its weighted sum on the health outcome of interest, and its weights α give the relative contribution of each component in the sum.

A wide variety of statistical methods have been advanced for modeling multiple pollutants. Five appraches are detailed in Davalos et al. (2017). Of these, adding additive main effects can present biased estimates in the presence of highly correlated variables, while nonparameteric methods are less interpretable than desired. The research teams and related stake-holders involved in air pollution research are often interdisciplinary, such that it is crucial that model parameters be easily interpretable and the estimation results be very communicable. While the unsupervised dimension reduction methods (such as principle components analysis and k-means clustering) indentified in Davalos et al. (2017) are difficult to interpret, some of the supervised methods consider easier-to-interpret weighted sums of pollutant concentrations. In Pachon et al. (2012), the authors specify weights from data rather than estimating them, while Roberts and Martin (2006) introduces a model that is equivalent to one that assumes s_1 is linear. While are all viable statistical methods, they lack the GAIM's combination of extensibility and interpretability.

Davalos et al. (2017) also discusses non-parametric methods, including Bayesian Kernel Machine Regression (BKMR). BKMR allows for estimation and variable selection. It was introduced in Bobb et al. (2015), and an R package was released in 2018 (Bobb et al. 2018).

Question 2

The relationship between daily coronavirus deaths and air pollution levels has become an active area of research in recent months. For instance, Wu et al. (2020) finds that a 1 μ g increase in long-term exposure to ambient PM_{2.5} increases the coronavirus death rate by 15%. Additional studies that examine this relationship include Conticini, Frediani, and Caro (2020), Sciomer et al. (2020), and Setti et al. (2020).

However, much work remains to be done. For instance, non-COVID-19 daily mortality data is generally not yet available, such that we do not have an accurate measure of *excess* deaths attributable to COVID-19. These excess deaths could be attributable to under-reported COVID-19 case and death counts (due to limited testing), restricted access to care for patients with other health conditions, or potential reporting delays. Moreover, cumulative COVID-19 mortality will likely continue to rise for some time, making the question of excess deaths due to COVID-19 best suited for an ongoing inquiry that help inform public health responses.

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Anticipated roles of trainees (students and post-doctoral fellows)

Kamal will develop the Bayesian implementation of the multiple pollutant models in Stan. This includes exploring determining appropriate prior distributions for the weights α and developing visualizations that communicate model results in a simple and interpretable way. He will be responsible for producing paper(s) summarizing the results of this model when run on Canadian air pollution and mortality data. To facilitate team communication and cohesion, he will also split time between Toronto (at the Centre for Global Health Research) and Ottawa (at the University of Ottawa), and use the proximity to Quebec to visit the collaborators there.

The University of Toronto PhD student will compare the results from the Bayesian random walk models to those obtained from non-MCMC methods. For example, these could include frequentist methods that fit (natural cubic) splines or Bayesian inference using R-INLA (Rue, Martino, and Chopin 2009). The University of Laval/University of Ottawa PhD student will compare the results from the Stan implementation to those obtained by a case-crossover model.

Plans for dissemination and communication

The results and findings of this multiple pollutant inquiry will be shared with Health Canada and the Institut National de Santé Publique du Québec. The lead investigators have a track record of publishing their research results in statistical and epidemiolgical journals, and aim to publish the results of this project in high-impact journals. They (or the trainees) will also attend appropriate conferences to present the work while it in progress.

Suggested reviewers

Ask Patrick?

Five CVs

• Patrick, Fateh, Hwashin, Meredith (?), Cindy

Preliminary budget description

The CANSSI Collaborative Research Team (CRT) grant is for \$180,000 over 3 years. We propose the budget:

- 1. \$30,000/year to support post-doctoral funding; matched by the Centre for Global Health Research.
- 2. \$12,000/year to support a Laval University or University of Ottawa PhD student.
- 3. \$12,000/year to support a University of Toronto PhD student.
- 4. \$6,000/year to support travel to/from the cities of the lead investigators Toronto, Ottawa, and Quebec and annual team meetings held around the Statistical Society of Canada conference.