



# Data Integration Model for Air Quality: A Hierarchical Approach to the Global **Estimation of Exposures to Ambient Air Pollution**

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### **OUTLINE**

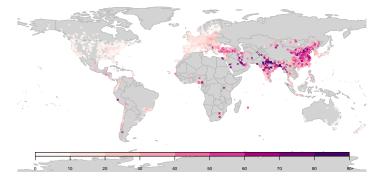
- ▶ Introduction
- ► DIMAQ
- ► Results
- ► Conclusions

#### INTRODUCTION

- ▶ Air pollution has been identified as a global health priority.
- ▶ In 2016, the World Health Organisation (WHO) estimated that over 3 million deaths can be attributed to ambient air pollution.
- ▶ The Global Burden of Disease (GBD) project estimate that in 2015 ambient air pollution was in the top ten leading risks to global health.
- ▶ Burden of disease calculations require accurate estimates of population exposure for each country.

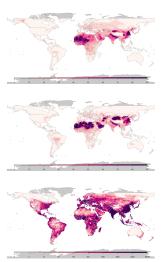
## ESTIMATING PM<sub>2.5</sub>

- ▶ Accurate estimates of exposure to air pollution are required
  - at global, national and local levels
  - with associated measures of uncertainty.
- ▶ While networks are expanding, ground monitoring is limited in many areas of the world.



## ESTIMATING PM<sub>2.5</sub>

- Can utilise information from other sources
  - satellite remote sensing
  - atmospheric models
  - population estimates
  - ▶ land use
  - local network characteristics.
- Result of modelling and will be subject to uncertainties and biases.



### DATA INTEGRATION MODEL FOR AIR QUALITY

- ▶ Developed to the Data Integration Model for Air Quality (DIMAO).
- ▶ DIMAQ calibrates ground measurements to estimates
  - satellite remote sensing,
  - specific components of chemical transport models
  - land use
  - population.
- ▶ The coefficients in the calibration model are estimated by country.
- Model allows borrowing from higher aggregations and if information is not available on a country level.
- Exploits a geographical nested hierarchy.
- Achieved using hierarchical random effects.

#### **REGIONS**

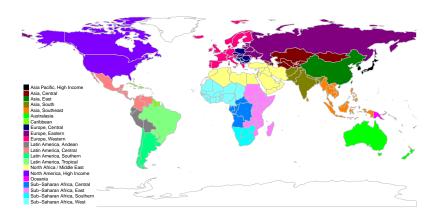


Figure: Map of regions.

### **SUPER-REGIONS**

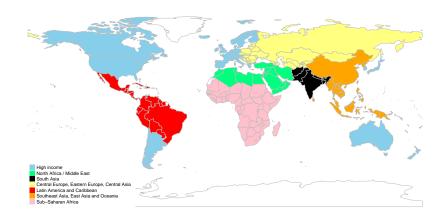


Figure: Map of super-regions.

## DATA INTEGRATION MODEL FOR AIR QUALITY

- Ground measurements at point locations, s, within grid cell, l, country, i, region, j, and super–region, k are denoted by  $Y_{sliik}$ .
- ▶ The model consists of a set of fixed and random effects, for both intercepts and covariates, and is given as follows,

$$\begin{split} \log(Y_{slijk}) &= \tilde{\beta}_{0,lijk} &+ \sum_{p \in P} \beta_p X_{p,slijk} \\ &+ \sum_{q \in Q} \tilde{\beta}_{q,lijk} X_{slijk} \\ &+ \epsilon_{slijk} \; . \end{split}$$

#### HIERARCHICAL RANDOM EFFECTS

▶ The random effect terms have contributions from the country, the region and the super-region.

$$\tilde{\beta}_{q,ijk} = \beta_q + \beta_{q,ijk}^{C} + \beta_{q,jk}^{R} + \beta_{q,k}^{SR}$$

▶ The intercept also having a random effect for the cell representing within-cell variation in ground measurements.

$$\tilde{\beta}_{0,lijk} = \beta_0 + \beta_{0,lijk}^G + \beta_{0,ijk}^C + \beta_{0,jk}^R + \beta_{0,k}^{SR}$$

#### RANDOM EFFECTS STRUCTURE

▶ The coefficients for super-regions are distributed with mean equal to the overall mean ( $\beta_0$ , the fixed effect) and variance representing the between super-region variation,

$$\beta_k^{SR} \sim N(\beta, \sigma_{SR}^2)$$

▶ The coefficients for regions are distributed with mean equal to the coefficient for the super-region with variance representing the between region variation,

$$\beta_{jk}^R \sim N(\beta_k^{SR}, \sigma_{R,k}^2)$$

▶ The coefficients for a country is distributed with mean equal to the coefficient for the region with variance representing the between country variation,

$$\beta_{ijk}^{C} \sim N(\beta_{jk}^{R}, \sigma_{C,jk}^{2})$$

#### INFERENCE

- Approximate Bayesian inference, such as Integrated Nested Laplace Approximations (INLA), provide fast and efficient methods for modelling with latent Gaussian models.
- ▶ INLA performs numerical calculations of posterior densities using Laplace Approximations hierarchical latent Gaussian models:

$$p(\theta_k|\mathbf{y}) = \int p(\mathbf{\theta}|\mathbf{y})d\mathbf{\theta}_{-k} \quad p(z_j|\mathbf{y}) = \int p(z_j|\mathbf{\theta},\mathbf{y})p(\mathbf{\theta}|\mathbf{y})d\mathbf{\theta}$$

▶ Latent Gaussian models allows for sparse matrices, and therefore efficient computation.

#### COMPUTATION

- R-INLA was used to implement DIMAQ.
- ▶ Unable to run this model on standard computers (4-8GB RAM).
- Required the use of a High-Performance Computing (HPC) service.
  - Balena cluster at University of Bath.
  - ▶  $2 \times 512$ GB RAM nodes ( $32 \times 32$ GB RAM cores).
- ▶ Took an iterative approach to prediction.

#### **EVALUATION: CROSSVALIDATION**

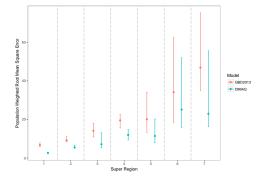


Figure: Summaries of predictive ability of the GBD2013 model and DIMAQ, for each of seven super-regions: 1, High income; 2, Central Europe, Eastern Europe, Central Asia; 3, Latin America and Caribbean; 4, Southeast Asia, East Asia and Oceania; 5, North Africa / Middle East; 6, Sub-Saharan Africa; 7, South Asia. For each model, population weighted root mean squared errors  $(\mu \mathrm{gm}^{-3})$  are given with dots denoting the median of the distribution from 25 training/evaluation sets and the vertical lines the range of values.

### **PREDICTIONS**

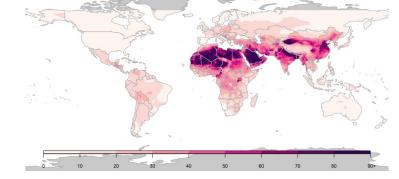


Figure: Median estimates of annual averages of  $PM_{2.5}$  ( $\mu gm^{-3}$ ) for 2014 for each grid cell (0.1°  $\times$  0.1° resolution) using DIMAQ.

### **UNCERTAINTY**

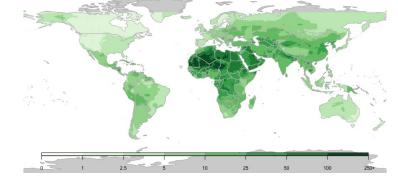


Figure: Half the width of 95% posterior credible intervals for 2014 for each grid cell  $(0.1^{o} \times 0.1^{o}$  resolution) using DIMAQ.

### POSTERIOR DISTRIBUTIONS

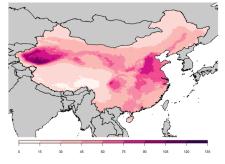


Figure: Medians of posterior distributions for estimates of annual mean  $PM_{2.5}$  concentrations  $(\mu gm^{-3})$  for 2014, in China.

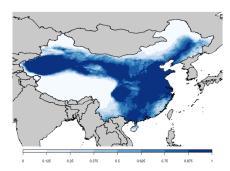


Figure: Probability of exceeding 35  $\mu$ gm<sup>-3</sup> using a Bayesian hierarchical model for each grid cell  $(0.1^{\circ} \times 0.1^{\circ} \text{ resolution})$  for 2014, in China.

## POPULATION EXPOSURES TO PM<sub>2.5</sub>

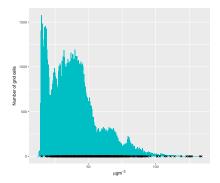


Figure: Estimated annual average concentrations of PM<sub>2.5</sub> by grid cell ( $0.1^{o} \times 0.1^{o}$  resolution). Black crosses denote the annual averages recorded at ground monitors.

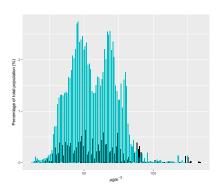
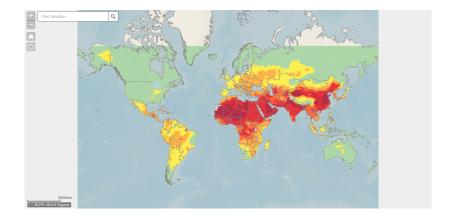


Figure: Estimated population level exposures (blue bars) and population weighted measurements from ground monitors (black bars).

#### CONCLUSION

- DIMAQ integrates data from multiple sources with producing high-resolution estimates of concentrations of ambient particulate matter.
- ▶ Estimates used by the WHO and GBD in burden of disease calculations.
- ► Future Developments
  - Higher resolution estimates
  - Within country variability
  - Allowing for errors and biases in covariates
  - Use data at native resolutions
- Possible approaches to address these issues
  - Statistical downscaling
  - Bayesian melding.

### INTERACTIVE MAP



#### REFERENCES

#### ▶ DIMAQ Paper:

http://onlinelibrary.wiley.com/doi/10.1111/rssc.12227/full

#### ▶ WHO Report:

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http://who.int/phe/publications/
air-pollution-global-assessment/en/
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#### ▶ GBD Paper:

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http://www.thelancet.com/journals/lancet/article/
PIIS0140-6736(16)31679-8/abstract
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#### ▶ Interactive Map:

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http://maps.who.int/airpollution/
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# ANY QUESTIONS?

