# Using statistical methods and reproducible tools to gain new insights from biomedical and public health data

#### Ariel Mundo Ortiz

Centre de Recherches Mathematiques, Université de Montréal

### MfPH Next Generation Seminar Series

3/15/23



#### Introduction

Data is the core of research. However, data is not information, as it needs to be processed before we can get information from it.

#### Introduction

- Data is the core of research. However, data is not information, as it needs to be processed before we can get information from it.
- This is specially true in the case of health research: public health, or biomedical data can be complex, and decisions along the analysis can result in different interpretations.

#### Introduction

- Data is the core of research. However, data is not information, as it needs to be processed before we can get information from it.
- This is specially true in the case of health research: public health, or biomedical data can be complex, and decisions along the analysis can result in different interpretations.
- In this talk I will focus on two examples that showcase how we can get more insight from data

The Case of Public Health Data

#### The Case of Public Health Data

■ COVID-19 vaccination has been an important component of public health strategies aimed at managing the pandemic.

<sup>&</sup>lt;sup>1</sup>Nafilyan et al. 2021.

<sup>&</sup>lt;sup>2</sup>Gerretsen et al. 2021.

- COVID-19 vaccination has been an important component of public health strategies aimed at managing the pandemic.
- However, COVID-19 vaccination has not been equal across different population segments.

<sup>&</sup>lt;sup>1</sup>Nafilyan et al. 2021.

<sup>&</sup>lt;sup>2</sup>Gerretsen et al. 2021.

- COVID-19 vaccination has been an important component of public health strategies aimed at managing the pandemic.
- However, COVID-19 vaccination has not been equal across different population segments.

<sup>&</sup>lt;sup>1</sup>Nafilyan et al. 2021.

<sup>&</sup>lt;sup>2</sup>Gerretsen et al. 2021.

- COVID-19 vaccination has been an important component of public health strategies aimed at managing the pandemic.
- However, COVID-19 vaccination has not been equal across different population segments.
- Individuals with lower income, and those belonging to a racial/ethnic minority have had lower vaccination uptake<sup>1,2</sup>.

<sup>&</sup>lt;sup>1</sup>Nafilyan et al. 2021.

<sup>&</sup>lt;sup>2</sup>Gerretsen et al. 2021.

The Fields Institute collected some very nice data regarding COVID-19 vaccination in Ontario, the Survey of COVID-19 related Behaviours and Attitudes.

- The Fields Institute collected some very nice data regarding COVID-19 vaccination in Ontario, the Survey of COVID-19 related Behaviours and Attitudes.
  - The survey ran between late 2021 and early 2022 and collected socio-demographic information along with self-reported vaccination status ("Have you received the first dose of the Covid vaccine?")

Table 1: Selected socio-economic factors from the survey

Variable	Levels
Age group Income bracket (CAD) Race/ethnicity	16-34,35-54,55 and over under 25,000, 25,000-59,999, 60,000 and above Arab/Middle Eastern, Black, East Asian/Pacific Islander, Indigenous, Latin American, Mixed, South Asian, White Caucasian, Other

The Case of Public Health Data

#### COVID-19: The Case of Ontario

 Other studies have analyzed the dependency on vaccination status using socio-economic data.

- Other studies have analyzed the dependency on vaccination status using socio-economic data.
- We could do the same, but what other information could we get from this data?

- Other studies have analyzed the dependency on vaccination status using socio-economic data.
- We could do the same, but what other information could we get from this data?
- From a Public Health Perspective, there have been some relatively recent developments in Ontario.

The Case of Public Health Data

#### COVID-19: The Case of Ontario

 However, Ontario adopted in late 2019 the Health Regions for healthcare and phased out the Local Health Integration Network (LHIN) approach.

- However, Ontario adopted in late 2019 the Health Regions for healthcare and phased out the Local Health Integration Network (LHIN) approach.
- The change is relatively new, and therefore, geographical data can be used to analyze data within the different Health Regions.

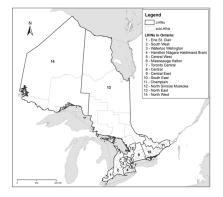


Figure 1: Ontario LHINs (Crighton et al. 2015)

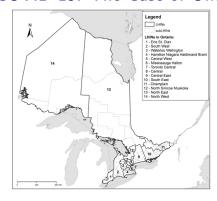


Figure 1: Ontario LHINs (Crighton et al. 2015)



Figure 2: Ontario Health Regions (Ontario Business Health Plan 2022-2023)

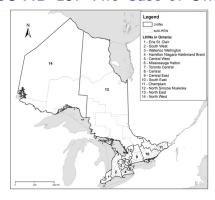


Figure 1: Ontario LHINs (Crighton et al. 2015)



Figure 2: Ontario Health Regions (Ontario Business Health Plan 2022-2023)

Data cleaning is important

■ Therefore, we decided to integrate the different Health Regions in our analysis to determine the odds of vaccination.

$$\log \left( \frac{p(\text{vac})}{1 - p(\text{vac})} \right) = \beta_0 + \beta_1(\text{Age group}) + \beta_2 \text{ Race} + \beta_3 \text{ Health Region} + \beta_4 \text{ Income} + (1)$$

$$\beta_5$$
(Health Region × Race) +  $\beta_6$  (Income × Race)

#### Results

Table 2: Selected Multivariable Regression Results

Characteristic	OR	95% CI	p-value
Income (CAD)			
60000 and above			0.011
25000-59999 under 25000	0.59 0.37	0.39, 0.89 0.25, 0.56	0.011 <0.001
Race	0.37	0.23, 0.30	<0.001
White/Caucasian	_	_	
Arab/Middle Eastern	0.31	0.14, 0.69	0.004
Black	0.32	0.17, 0.60	< 0.001
East Asian/Pacific Islander	1.15	0.50, 2.66	0.7
Indigenous	0.44	0.19, 1.02	0.056
Latin Aamerican	0.28	0.11, 0.67	0.004
Mixed Other	0.64 0.22	0.25, 1.65 0.12, 0.41	< 0.4
South Asian	0.91	0.49, 1.69	0.801
Health Region			
Toronto		—	
Central	1.47	0.92, 2.35	0.11
East West	1.42 1.55	0.90, 2.23 1.05, 2.30	0.13 0.029
Income and Race	1.55	1.05, 2.50	0.029
25000-59999 * Arab/Middle Eastern	1.79	0.67, 4.83	0.2
under 25000 * Arab/Middle Eastern	3.05	1.26, 7.39	0.013
25000-59999 * Black	1.34	0.59, 3.05	0.5
under 25000 * Black	3.19	1.45, 6.99	0.004
25000-59999 * East Asian/Pacific Islander	0.42	0.17, 1.05	0.062
under 25000 * East Asian/Pacific Islander	1.16	0.47, 2.86	0.8
25000-59999 * Indigenous	1.36	0.48, 3.89	0.6
under 25000 * Indigenous	1.45	0.55, 3.80	0.5
25000-59999 * Latin American	1.24	0.45, 3.43	0.7

#### Results

Characteristic	OR	95% CI	p-value	
under 25000 * Latin American	2.80	1.04, 7.51	0.041	
25000-59999 * Mixed	0.85	0.32, 2.26	0.7	
under 25000 * Mixed	1.10	0.37, 3.27	0.9	
25000-59999 * Other	6.93	2.65, 18.1	< 0.001	
under 25000 * Other	4.59	2.33, 9.05	< 0.001	
25000-59999 * South Asian	1.20	0.51, 2.85	0.7	
under 25000 * South Asian	2.00	0.93, 4.30	0.077	
Race and Health Region				
Arab/Middle Eastern * Central	0.66	0.26, 1.70	0.4	
Black * Central	0.44	0.19, 0.98	0.046	
East Asian/Pacific Islander * Central	0.98	0.38, 2.53	>0.9	
Mixed * East	0.91	0.28, 3.03	0.9	
other * East	1.05	0.39, 2.83	>0.9	
South Asian * East	0.52	0.19, 1.45	0.2	
Arab/Middle Eastern * West	1.00	0.37, 2.73	>0.9	
Black * West	0.76	0.32, 1.80	0.5	
East Asian/Pacific Islander * West	0.52	0.20, 1.34	0.2	
Indigenous <sup>*</sup> West	0.39	0.14, 1.09	0.073	
Latin American * West	0.94	0.32, 2.72	>0.9	
Mixed * West	0.37	0.12, 1.16	0.089	
Other * West	0.41	0.18, 0.93	0.032	
South Asian * West	0.41	0.18, 0.95	0.037	
1 OR — Odds Ratio, CI — Confidence Interval				

 Our results show that there were disparities in vaccination uptake in Ontario.

<sup>&</sup>lt;sup>3</sup>Hawkins 2020.

- Our results show that there were disparities in vaccination uptake in Ontario.
- People in certain racial minority groups had lower odds of vaccination than White/Caucasian individuals.

<sup>&</sup>lt;sup>3</sup>Hawkins 2020.

- Our results show that there were disparities in vaccination uptake in Ontario.
- People in certain racial minority groups had lower odds of vaccination than White/Caucasian individuals.
- However, individuals that identified with a racial/ethnic minority and that were in a low household income bracket (<60k CAD) had higher odds of vaccination than individuals with a high household income.

<sup>&</sup>lt;sup>3</sup>Hawkins 2020.

- Our results show that there were disparities in vaccination uptake in Ontario.
- People in certain racial minority groups had lower odds of vaccination than White/Caucasian individuals.
- However, individuals that identified with a racial/ethnic minority and that were in a low household income bracket (<60k CAD) had higher odds of vaccination than individuals with a high household income.
- This is likely caused by the type of occupation: people in racial minorities, and those with a low household income work in essential occupations<sup>3</sup>, and thus potentially got the vaccine to be able to work.

<sup>&</sup>lt;sup>3</sup>Hawkins 2020.

But there are also intra-provincial differences in vaccine uptake within the Health Regions:

- But there are also intra-provincial differences in vaccine uptake within the Health Regions:
  - For example, South Asian individuals in the West Health Region had lower odds of vaccination that in other Health Regions.

- But there are also intra-provincial differences in vaccine uptake within the Health Regions:
  - For example, South Asian individuals in the West Health Region had lower odds of vaccination that in other Health Regions.
  - These results provide a more comprehensive assessment of COVID-19 vaccination rates within Ontario, as they showed that certain minority groups within specific income brackets and certain Health Regions had differences in vaccination.

The Case of Biomedical Data

#### The Case of Biomedical Data

Biomedical studies often collect longitudinal data to see the effect of an intervention over time:

- Biomedical studies often collect longitudinal data to see the effect of an intervention over time:
  - How a chemotherapy treatment changes the metabolism of a tumor

- Biomedical studies often collect longitudinal data to see the effect of an intervention over time:
  - How a chemotherapy treatment changes the metabolism of a tumor
  - How the concentration of a drug changes over time in the blood

- Biomedical studies often collect longitudinal data to see the effect of an intervention over time:
  - How a chemotherapy treatment changes the metabolism of a tumor
  - How the concentration of a drug changes over time in the blood
- How is this data typically analyzed?

#### Linear Models

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt} \colon$  is the response for subject i in treatment group j at time t

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt}$ : is the response for subject i in treatment group j at time t  $\beta_0$ : the mean group value

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt}$ : is the response for subject i in treatment group j at time t

 $\beta_0$ : the mean group value

 $time_t$ ,  $treatment_j$ : fixed effects

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt}$ : is the response for subject i in treatment group j at time t

 $\beta_0$ : the mean group value

 $time_t$ ,  $treatment_i$ : fixed effects

 $\beta_1,\beta_2$  and  $\beta_3$ : linear slopes of the fixed effects.

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt}$ : is the response for subject i in treatment group j at time t

 $\beta_0$ : the mean group value

 $time_t$ ,  $treatment_i$ : fixed effects

 $\beta_1,\beta_2$  and  $\beta_3$ : linear slopes of the fixed effects.

 $\varepsilon_{ijt}$ : error, assumed to be  $\sim N(0,\sigma^2)$ 

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + \beta_2 \times time_t + \beta_3 \times time_t \times treatment_j + \varepsilon_{ijt}$$
(2)

where,

 $y_{ijt}$ : is the response for subject i in treatment group j at time t

 $\beta_0$ : the mean group value

 $time_t$ ,  $treatment_i$ : fixed effects

 $\beta_1, \beta_2$  and  $\beta_3$ : linear slopes of the fixed effects.

 $\varepsilon_{ijt}$ : error, assumed to be  $\sim N(0,\sigma^2)$ 

A LMEM follows the same exact structure, only incorporates a random effect  $\alpha_{ij}$ , which allows for different intercepts.

### Trends Over Time

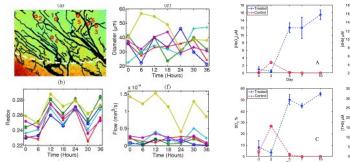


Figure 3: Tumor imaging data (Skala et al. 2010)

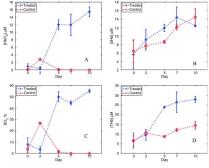


Figure 4: Tumor oxygenation data (Vishwanath et al. 2009)

#### Trends Over Time

The issue in those data is that the trends are not linear, and therefore, a linear model will miss changes in the signal where some metabolic or physiological relevant change is taking place.

<sup>&</sup>lt;sup>4</sup>Beck and Jackman 1998.

### Trends Over Time

- The issue in those data is that the trends are not linear, and therefore, a linear model will miss changes in the signal where some metabolic or physiological relevant change is taking place.
- Polynomial effects can be used, but they create biases at the boundaries of the covariates<sup>4</sup>.

<sup>&</sup>lt;sup>4</sup>Beck and Jackman 1998.

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + f(time_t \mid \beta_j) + \varepsilon_{ijt}$$
 (3)

■ The change of  $y_{ijt}$  over time is represented by the smooth  $function \ f(time_t \mid \beta_j)$  with inputs as the covariates  $time_t$  and parameters  $\beta_j$ .

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + f(time_t \mid \beta_j) + \varepsilon_{ijt}$$
 (3)

■ The change of  $y_{ijt}$  over time is represented by the smooth  $function \ f(time_t \mid \beta_j)$  with inputs as the covariates  $time_t$  and parameters  $\beta_j$ .

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + f(time_t \mid \beta_j) + \varepsilon_{ijt}$$
 (3)

- $\blacksquare$  The change of  $y_{ijt}$  over time is represented by the smooth function  $f(time_t \mid \beta_j)$  with inputs as the covariates  $time_t$  and parameters  $\beta_j.$
- We can use a basis function to estimate the smooth function.

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + f(time_t \mid \beta_j) + \varepsilon_{ijt}$$
 (3)

- $\blacksquare$  The change of  $y_{ijt}$  over time is represented by the smooth function  $f(time_t \mid \beta_j)$  with inputs as the covariates  $time_t$  and parameters  $\beta_j.$
- We can use a basis function to estimate the smooth function.

$$y_{ijt} = \beta_0 + \beta_1 \times treatment_j + f(time_t \mid \beta_j) + \varepsilon_{ijt}$$
 (3)

- The change of  $y_{ijt}$  over time is represented by the smooth  $function \ f(time_t \mid \beta_j)$  with inputs as the covariates  $time_t$  and parameters  $\beta_j$ .
- We can use a *basis function* to estimate the smooth function.
- Splines are helpful as basis functions: Thin plate regression splines (TPRS) are computationally efficient, and the underlying principle is that of polynomial pieces "joined" together

### How GAMs work

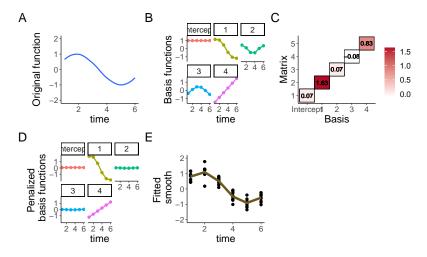


Figure 5: Fitting process of a GAM.

### An Example

 Simulated data from a study on radiotherapy in a mouse model of cervical cancer<sup>5</sup>.

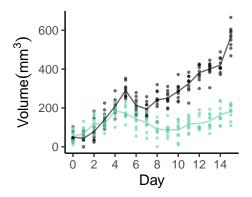


Figure 6: Tumor volume in two groups of tumors under radiotherapy

<sup>&</sup>lt;sup>5</sup>Zheng et al. 2019.

- Beck, Nathaniel and Simon Jackman (Apr. 1998). "Beyond Linearity by Default: Generalized Additive Models". In: *American Journal of Political Science* 42.2, p. 596. DOI: 10.2307/2991772. URL: https://doi.org/10.2307/2991772.
- Gerretsen, Philip et al. (Nov. 2021). "Individual determinants of COVID-19 vaccine hesitancy". In: *PLOS ONE* 16.11. Ed. by Leeberk Raja Inbaraj, e0258462. DOI: 10.1371/journal.pone.0258462. URL: https://doi.org/10.1371/journal.pone.0258462.
- Hawkins, Devan (June 2020). "Differential occupational risk for COVID-19 and other infection exposure according to race and ethnicity". In: *American Journal of Industrial Medicine* 63.9, pp. 817–820. DOI: 10.1002/ajim.23145. URL: https://doi.org/10.1002/ajim.23145.

- Nafilyan, Vahe et al. (July 2021). "Sociodemographic inequality in COVID-19 vaccination coverage among elderly adults in England: a national linked data study". In: *BMJ Open* 11.7, e053402. DOI: 10.1136/bmjopen-2021-053402. URL: https://doi.org/10.1136/bmjopen-2021-053402.
- Zheng, Xianchuang et al. (Sept. 2019). "A Near-Infrared Phosphorescent Nanoprobe Enables Quantitative, Longitudinal Imaging of Tumor Hypoxia Dynamics during Radiotherapy". In: Cancer Research 79.18, pp. 4787–4797. DOI: 10.1158/0008-5472.can-19-0530. URL: https://doi.org/10.1158/0008-5472.can-19-0530.