CoDatMo: Creating a Common Epidemiological Framework with Bayesian Models

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Who

- Breck Baldwin: Project Manager Laplace Lab @Columbia statistics
- Andrew Gelman: @Columbia statistics
- Simon Maskell: PI Big Hypotheses Lab at U of Liverpool

How

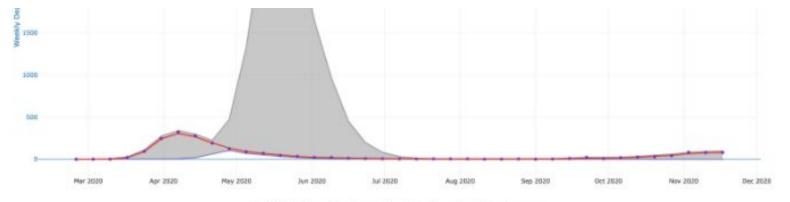
- UKRI Big Hypotheses EP/R018537/1
- NSF RAPID grant 2055251

Outline

- Overview CoDatMo
- Brief intro to Bayesian Modeling and Stan
- Go over 'model reproduction checklist'
- Resources

CoDatMo Goals and History

- 1. August 2020:
 - a. Stan--a popular language for modeling COVID-19 epidemiology ~ 20 models
 - b. The UK Joint Biosecurity Center (JBC) uses many models but similarities/differences
 - c. Liverpool/Columbia decide to create an online resource for comparative COVID modeling
 - i. Improve the models
 - ii Share ideas
- 2. November 15, 2020: NSF RAPID grant 2055251 for COVID awarded
- 3. December 2, 2020: https://github.com/codatmo up and running
 - a. Model example
 - b. Data
- Submissions:
 - a. December 2, 2020: Liverpool
 - b. December 8, 2020: Ye Old Lancaster
 - c. December 20, 2020: Simple SIR
 - d.Rethinking appeal of CoDatMo to external contributors
 - e. March 14, 2020: UNINOVE Sao Paulo
- 5. April 14, 2020: This presentation



Death incidence in nhs sandwell and west birmingham ccg

home about

Welcome to the CoDatMo site

CoDatMo (COvid DATa MOdel) exists to replicate and host COVID-19 models written in the Bayesian modeling languages like Stan or PyMC. The goals include:

- Reproduce COVID-19 models with an eye to:
 - Get the word out on important COVID-19 models by offering well documented examples approachable by statiticians, programmers and researchers.
 - Increase the impact of the original researchers and get them additional exposure.
 - Help researchers improve their models via the replication process
- Provide a common language for implementing various models.
- Facillitate information exchange between model developers.
- Provide public access to models.

CombineR

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UNINOVE_Sao_Paulo

Covid Modeling work at UNINOVE Brazil

170 (1) 0 Updated 3 days ago

Data

Collection of open-source data sources for the United Kingdom and the rest of the world. To be used for testing of the various models.

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Liverpool

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Top languages

HTML R Stan

People



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Invite someone

Over 50 models for COVID-19 written in Stan: Why?

- Hierarchical models
- Noise tolerant models with small to medium data sets
- Lots of room for integration of prior knowledge
- Can be very interpretable models to support counterfactual reasoning etc...
- Very robust uncertainty handling
- Huge effort on inference safety
 - Bayesian workflow
 - Inference diagnostics
 - Self monitoring gradient crawler
- Generative models
 - Mechanistic
 - Human interpretable
 - Simulation

Deep Learning

Repos in CoDatMo

https://codatmo.github.io/Simple SIR/

https://github.com/codatmo/UNINOVE_Sao_Paulo

Resources

- Codatmo project: https://codatmo.github.com
- Slides: https://codatmo.github.com//documentation/talks/columbia_4_2021.pdf
- Modern Bayesian modeling packages:
 - Stan: https://mc-stan.org
 - R, Python, Scala, https://mc-stan.org/users/interfaces/
 - High level interfaces in lme4 tradition (R only)
 - brms: https://paul-buerkner.github.io/brms/
 - RStanArm: https://mc-stan.org/rstanarm/
 - PyMC Python HMC/NUTS: https://pypi.org/project/pymc/
 - TensorFlow/PyTorch
- Statistical Rethinking, Second Edition, by Richard McElreath
 - Youtube lectures https://youtube.com/playlist?list=PLDcUM9US4XdNM4Edgs7weiylguLSToZRI
 - brms implementation of 1st Edition at: https://bookdown.org/content/4857/
- Bayesian Statistics for the Social Sciences GR5065 in http://www.qmss.columbia.edu/:
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