

Al buio non si trova

Biostatistics in the 21st century^a

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^aAvailable from <https://github.com/maxbiostat/presentations/>

Le dirò con due parole, chi son

Personal

- ⊙ Born and raised in Petrópolis-RJ;
- ⊙ Eldest of three kids;
- ⊙ Married and father of a daughter;
- ⊙ Mais Querido supporter.

Academic

- ⊙ BSc in Microbiology & Immunology (UFRJ, 2012);
- ⊙ PhD Evolutionary Biology (Edinburgh, 2018);
- ⊙ Post doctoral researcher at ENSP/Fiocruz (2019);
- ⊙ Lecturer (Assistant Professor) at EMap since Jan/2020.

Applications of Statistics/Mathematics

Applications in Epidemiology, (Molecular) Biology, Ecology, Psychology, Linguistics, etc.

Applied Statistics

Markov Chain Monte Carlo, Model combination and selection, Statistical Phylogenetics.

That's like... Your opinion, man

Logarithmic Opinion pooling, aka log-linear mixtures¹

$$\pi(\theta \mid \boldsymbol{\alpha}) = t(\boldsymbol{\alpha}) \prod_{i=1}^K f_i(\theta)^{\alpha_i}, \alpha_i \geq 0, \sum_{i=1}^K \alpha_i = 1.$$

Applications: combining forecast models, robust Bayesian inference.

- ⊙ Hierarchical approach: which prior $\pi_A(\boldsymbol{\alpha})$ will induce nice behaviour on both $p_A(\boldsymbol{\alpha} \mid \mathbf{y})$ and $p_T(\theta \mid \mathbf{y})$?
- ⊙ Bayesian predictive synthesis: can we solve

$$\int_{\mathfrak{X}} \gamma(y \mid \mathbf{x}) \prod_{i=1}^K f_i(x_i) d\mu(x_i) = \prod_{i=1}^J f_i(y)^{\alpha_j}?$$

¹<https://arxiv.org/abs/1502.04206>

Next-generation epidemiological surveillance²

$$\tilde{f}(\mathbf{y}) = \mathbb{H} \left(\mathbf{M}(\mathbf{y}), \boldsymbol{\alpha} \right),$$

with $\mathbf{M}(\mathbf{y}) = \{m_1(\mathbf{y}), \dots, m_K(\mathbf{y})\}$ a collection of forecasts.

- ⊙ Can we build a unified framework to aggregate models from different groups, places, types?
- ⊙ What form should the combining operator \mathbb{H} take?
- ⊙ Which proper scoring rules³ to use?

²https://github.com/maxbiostat/EBDS_2021

³https://github.com/maxbiostat/proper_scoring_rules

Normalised power prior⁴

$$\tilde{\pi}(\theta, a_0 \mid \mathbf{y}_0) = \frac{L(\mathbf{y}_0 \mid \theta)^{a_0} \pi(\theta \mid \eta) \pi_A(a_0 \mid \phi)}{c(a_0; \eta, \phi)}$$

Applications: clinical trials, quality control, policy-making.

- ⊙ How pick π_A such that prediction error (say) is minimised?
- ⊙ How to **efficiently** compute

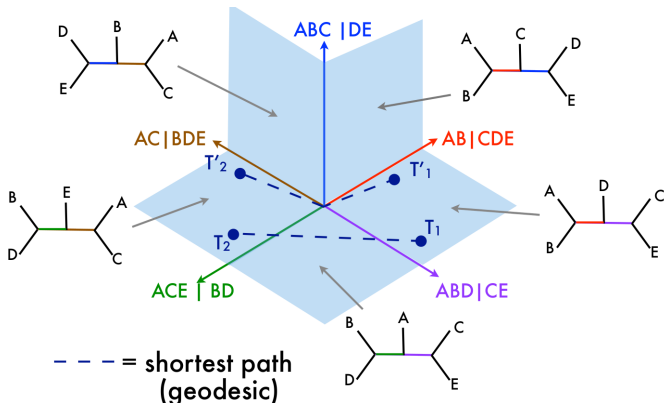
$$c(a_0; \eta, \phi) = \int_{\Theta} L(\mathbf{y}_0 \mid t)^{a_0} \pi(t \mid \eta) d\mu(t)$$

by leveraging its special properties as function of a_0 ?

⁴<https://doi.org/10.1002/sim.9124>

This place is weird

Traversing cubic complexes efficiently⁵



Applications: Molecular Epidemiology, Evolutionary Biology.

⁵<https://youtu.be/h9bWRQ6aeKA>

THE
END