joint work with K. Kamari, K. Mengersen and C. Robert april 2016, Toulouse Hypothesis testing

- central problem of statistical inference
- witness the recent ASA's statement on *p*-values (Wasserstein, 2016)
- dramatically differentiating feature between classical and Bayesian paradigms
- wide open to controversy and divergent opinions, includ. within the Bayesian community
- non-informative Bayesian testing case mostly unresolved, witness the Jeffreys–Lindley paradox

Berger (2003), Mayo & Cox (2006), Gelman (2008)

 $\mathfrak{B}$ 

Standard Bayesian approach to testing : consider two families of models, one for each of the hypotheses under comparison,

 $\mathfrak{M}_1$ :  $x \sim f_1(x|\theta_1), \ \theta_1 \in \Theta_1$  and  $\mathfrak{M}_2$ :  $x \sim f_2(x|\theta_2), \ \theta_2 \in \Theta_2$ ,

**Priors**  $\theta_1 \sim \pi_1(\theta_1)$  and  $\theta_2 \sim \pi_2(\theta_2)$ ,

 $m_1(x) = \int_{\Omega} f_1(x|\theta_1) \pi_1(\theta_1) d\theta_1$  and  $m_2(x) = \int_{\Omega} f_2(x|\theta_2) \pi_1(\theta_2) d\theta_2$ 

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$$\begin{split} \mathfrak{M}_1: \ x \sim f_1(x|\theta_1) \,, \ \theta_1 \in \Theta_1 \quad \text{and} \quad \mathfrak{M}_2: \ x \sim f_2(x|\theta_2) \,, \ \theta_2 \in \Theta_2 \,, \\ \mathbf{Priors} \ \theta_1 \sim \pi_1(\theta_1) \quad \text{and} \quad \theta_2 \sim \pi_2(\theta_2) \,, \end{split}$$

$$m_1(x) = \int_{\Theta_1} f_1(x|\theta_1) \, \pi_1(\theta_1) \, d\theta_1 \quad \text{and} \quad m_2(x) = \int_{\Theta_2} f_2(x|\theta_2) \, \pi_1(\theta_2) \, d\theta_2$$

either through Bayes factor or posterior probability, respectively :

$$\mathfrak{B}_{12}=\frac{m_1(x)}{m_2(x)},\quad \mathbb{P}(\mathfrak{M}_1|x)=\frac{\omega_1m_1(x)}{\omega_1m_1(x)+\omega_2m_2(x)};$$

the latter depends on the prior weights  $\omega_i = \pi(\Theta_i)$ 

Bayesian decision step

- $\bullet$  comparing Bayes factor  $\mathfrak{B}_{12}$  with threshold value of one or
- comparing posterior probability  $\mathbb{P}(\mathfrak{M}_1|x)$  with bound 1/2

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When comparing more than two models, model with *highest posterior probability* is the one selected, but highly dependent on the prior modelling.

• eliminates choice of  $\pi(\Theta_0)$ 

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- Jeffreys' scale of evidence :
  - if  $\log_{10}(\mathfrak{B}_{10}^{\pi})$  between 0 and 0.5, evidence against  $H_0$  weak,
  - if  $\log_{10}(\mathfrak{B}_{10}^{\pi})$  0.5 and 1, evidence substantial,
  - if  $\log_{10}(\mathfrak{B}^{\pi}_{10})$  1 and 2, evidence *strong* and
  - if  $\log_{10}(\mathfrak{B}_{10}^{\pi})$  above 2, evidence *decisive*

Quite arbitrary really ! : consequence of 0-1 loss function

- Bayesian model selection as comparison of k potential statistical models towards the selection of model that fits the data "best"
- mostly accepted perspective : it does not primarily seek to identify which model is "true", but compares fits

- Bayesian model selection as comparison of k potential statistical models towards the selection of model that fits the data "best"
- mostly accepted perspective : it does not primarily seek to identify which model is "true", but compares fits
- tools like Bayes factor naturally include a penalisation addressing model complexity, mimicked by Bayes Information (BIC) and Deviance Information (DIC) criteria .
- Under quite genreal conditions : consistent criterion for testing or model selection

- long-lasting impact of prior modeling, i.e., choice of prior distributions on parameters of both models, despite overall consistency proof for Bayes factor
- discontinuity in valid use of <u>improper priors</u> since they are not justified in most testing situations, leading to many alternative and *ad hoc* solutions, where data is either used twice or split in artificial ways [or further tortured into confession]
- binary (accept vs. reject) outcome more suited for immediate decision (if any) than for model evaluation, in connection with rudimentary loss function 0-1 [atavistic remain of Neyman-Pearson formalism]

- related impossibility to ascertain simultaneous misfit or to detect outliers
- no assessment of uncertainty associated with decision itself besides posterior probability
- difficult computation of marginal likelihoods in most settings with further controversies about which algorithm to adopt
- time for a paradigm shift?

New proposal for a paradigm shift ( !) in the Bayesian processing of hypothesis testing and of model selection

- convergent and naturally interpretable solution
- more extended use of improper priors

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Simple representation of the testing problem as a two-component mixture estimation problem where the weights are formally equal to 0 or 1

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- Approach inspired from consistency result of Rousseau and Mengersen (2011) on estimated overfitting mixtures
- Mixture representation not directly equivalent to the use of a posterior probability
- Calibration of posterior distribution of the weight of a model, moving from the notion of posterior probability of a model

Idea : Given two statistical models,

 $\mathfrak{M}_{1}: x_{i} \overset{\textit{ind.}}{\sim} f_{1}(x_{i}|\theta_{1}), \ \theta_{1} \in \Theta_{1} \quad \text{and} \quad \mathfrak{M}_{2}: x_{i} \overset{\textit{ind.}}{\sim} f_{2}(x|\theta_{2}), \ \theta_{2} \in \Theta_{2}, i \leq n$ 



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 $\mathfrak{M}_{\alpha}: x_{i} \overset{ind.}{\sim} \alpha f_{1}(x|\theta_{1}) + (1-\alpha)f_{2}(x|\theta_{2}), \ 0 \leq \alpha \leq 1, \quad i \leq n$ (1)

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Note : Both models correspond to special cases of (1), one for  $\alpha = 1$  and one for  $\alpha = 0$ Draw inference on mixture representation (1), as if each observation was individually and independently produced by the mixture model Sounds like approximation to the real model, but several definitive advantages to this paradigm shift :

- Bayes estimate of the weight  $\alpha$  replaces posterior probability of model  $\mathfrak{M}_1$ , equally convergent indicator of which model is "true", while avoiding artificial prior probabilities on model indices,  $\omega_1$  and  $\omega_2$
- interpretation of estimator of  $\alpha$  at least as natural as handling the posterior probability, while avoiding zero-one loss setting : proportion of individuals from each model
- $\alpha$  and its posterior distribution provide measure of proximity to the models, while being interpretable as data propensity to stand within one model
- further allows for alternative perspectives on testing and model choice, like predictive tools, cross-validation, and information indices

# Computational motivations

- avoids highly problematic computations of the marginal likelihoods, since standard algorithms are available for Bayesian mixture estimation
- straightforward extension to a finite collection of models, with a larger number of components, which considers all models at once and eliminates least likely models by simulation
- eliminates difficulty of label switching that plagues both Bayesian estimation and Bayesian computation, since components are no longer exchangeable
- posterior distribution of  $\alpha$  evaluates more thoroughly strength of support for a given model than the single figure outcome of a posterior probability
- $\bullet$  variability of posterior distribution on  $\alpha$  allows for a more thorough assessment of the strength of this support

## Noninformative motivations

- additional feature missing from traditional Bayesian answers : a mixture model acknowledges possibility that, for a finite dataset, *both* models or *none* could be acceptable
- standard (proper and informative) prior modeling can be reproduced in this setting, but non-informative (improper) priors also are manageable therein, provided both models first reparameterised towards shared parameters, e.g. location and scale parameters
- in special case when all parameters are common

$$\mathfrak{M}_{\alpha}: \ x \sim \alpha f_1(x|\theta) + (1-\alpha)f_2(x|\theta), 0 \leq \alpha \leq 1$$

if heta is a location parameter, a flat prior  $\pi( heta) \propto 1$  is available

## Weakly informative motivations

- using the *same* parameters or some *identical* parameters on both components highlights that opposition between the two components is not an issue of enjoying different parameters
- those common parameters are nuisance parameters, to be integrated out *[unlike Lindley's paradox]*
- prior model weights  $\omega_i$  rarely discussed in classical Bayesian approach, even though linear impact on posterior probabilities. Here, prior modeling only involves selecting a prior on  $\alpha$ , e.g.,  $\alpha \sim \mathscr{B}(a_0, a_0)$
- while  $a_0$  impacts posterior on  $\alpha$ , it always leads to mass accumulation near 1 or 0, i.e. favours most likely model
- sensitivity analysis straightforward to carry
- approach easily calibrated by parametric boostrap providing reference posterior of  $\alpha$  under each model
- natural Metropolis-Hastings alternative

• choice betwen Poisson  $\mathscr{P}(\lambda)$  and Geometric  $\mathscr{G}eo(p)$  distribution

- choice betwen Poisson  $\mathscr{P}(\lambda)$  and Geometric  $\mathscr{G}eo(p)$  distribution
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$$\mathfrak{M}_{lpha}$$
:  $lpha \mathscr{P}(\lambda) + (1 - lpha) \mathscr{G} eo(1/1 + \lambda)$ 

Allows for Jeffreys prior since resulting posterior is proper

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$$\mathfrak{M}_{lpha}: \ lpha \mathscr{P}(\lambda) + (1-lpha) \mathscr{G} eo(1/1+\lambda)$$

Allows for Jeffreys prior since resulting posterior is proper

• independent Metropolis–within–Gibbs with proposal distribution on  $\lambda$  equal to Poisson posterior (with acceptance rate larger than 75%)

When  $\alpha \sim \mathscr{B}e(a_0, a_0)$  prior, full conditional posterior

$$\alpha \sim \mathscr{B}e(n_1(\zeta) + a_0, n_2(\zeta) + a_0)$$

Exact Bayes factor opposing Poisson and Geometric

$$\mathfrak{B}_{12} = n^{n\bar{x}_n} \prod_{i=1}^n x_i! \, \Gamma\left(n+2+\sum_{i=1}^n x_i\right) / \Gamma(n+2)$$

although arbitrary from a purely mathematical viewpoint

#### Parameter estimation : $\lambda$ then $\alpha$



Posterior means of  $\lambda$  and medians of  $\alpha$  for 100 Poisson  $\mathscr{P}(4)$  datasets of size n = 1000, for  $a_0 = .0001, .001, .01, .1, .2, .3, .4, .5$ . Each posterior approximation is based on  $10^4$  Metropolis-Hastings iterations.

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# Consistency



Posterior means (*sky-blue*) and medians (*grey-dotted*) of  $\alpha$ , over 100 Poisson  $\mathscr{P}(4)$  datasets for sample sizes from 1 to 1000.

#### Behaviour of Bayes factor



Comparison between  $\mathbb{P}(\mathfrak{M}_1|x)$  (red dotted area) and posterior medians of  $\alpha$  (grey zone) for 100 Poisson  $\mathscr{P}(4)$  datasets with sample sizes *n* between 1 and 1000, for  $a_0 = .001, .1, .5$ 

• comparison of a normal  $\mathscr{N}(\theta_1,1)$  with a normal  $\mathscr{N}(\theta_2,2)$  distribution

## Normal-normal comparison

- comparison of a normal  $\mathscr{N}(\theta_1,1)$  with a normal  $\mathscr{N}(\theta_2,2)$  distribution
- mixture with identical location parameter  $\theta$  $\alpha \mathcal{N}(\theta, 1) + (1 - \alpha) \mathcal{N}(\theta, 2)$
- Jeffreys prior  $\pi(\theta) = 1$  can be used, since posterior is proper

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- Jeffreys prior  $\pi(\theta) = 1$  can be used, since posterior is proper
- Reference (improper) Bayes factor

$$\mathfrak{B}_{12} = 2^{n-1/2} / \exp \frac{1}{4} \sum_{i=1}^{n} (x_i - \bar{x})^2$$

# Consistency



Posterior means (wheat) and medians of  $\alpha$  (dark wheat), compared with posterior probabilities of  $\mathfrak{M}_0$  (gray) for a  $\mathscr{N}(0,1)$  sample, derived from 100 datasets for sample sizes equal to 15, 50, 100, 500. Each posterior approximation is based on 10<sup>4</sup> MCMC iterations.

## Comparison with posterior probability



Plots of ranges of  $\log(n) \log(1 - \mathbb{E}[\alpha|x])$  (gray color) and  $\log(1 - p(\mathfrak{M}_1|x))$  (red dotted) over 100  $\mathcal{N}(0, 1)$  samples as sample size *n* grows from 1 to 500. and  $\alpha$  is the weight of  $\mathcal{N}(0, 1)$  in the mixture model. The shaded areas indicate the range of the estimations and each plot is based on a Beta prior with  $a_0 = .1, .2, .3, .4, .5, 1$  and each posterior approximation is based on  $10^4$  iterations.

- convergence to one boundary value as sample size *n* grows
- impact of hyperarameter *a*<sub>0</sub> slowly vanishes as *n* increases, but present for moderate sample sizes
- when simulated sample is neither from  $\mathcal{N}(\theta_1, 1)$  nor from  $\mathcal{N}(\theta_2, 2)$ , behaviour of posterior varies, depending on which distribution is closest

- binary dataset, R dataset about diabetes in 200 Pima Indian women with body mass index as explanatory variable
- comparison of logit and probit fits could be suitable. We are thus comparing both fits via our method

$$\mathfrak{M}_1 : y_i \mid \mathbf{x}^i, \theta_1 \sim \mathscr{B}(1, p_i) \quad \text{where} \quad p_i = \frac{\exp(\mathbf{x}^i \theta_1)}{1 + \exp(\mathbf{x}^i \theta_1)}$$
  
 $\mathfrak{M}_2 : y_i \mid \mathbf{x}^i, \theta_2 \sim \mathscr{B}(1, q_i) \quad \text{where} \quad q_i = \Phi(\mathbf{x}^i \theta_2)$ 

#### Common parameterisation

Local reparameterisation strategy that rescales parameters of the probit model  $\mathfrak{M}_2$  so that the MLE's of both models coincide. choudhuty et al., 2007

$$\Phi(\mathbf{x}^{i}\theta_{2}) pprox rac{\exp(k\mathbf{x}^{i} heta_{2})}{1+\exp(k\mathbf{x}^{i} heta_{2})}$$

and use best estimate of k to bring both parameters into coherency

$$(k_0, k_1) = \left(\widehat{\theta_{01}}/\widehat{\theta_{02}}, \widehat{\theta_{11}}/\widehat{\theta_{12}}\right),$$

reparameterise  $\mathfrak{M}_1$  and  $\mathfrak{M}_2$  as

$$\mathfrak{M}_1 : y_i \mid \mathbf{x}^i, \theta \sim \mathscr{B}(1, p_i) \quad \text{where} \quad p_i = rac{\exp(\mathbf{x}^i heta)}{1 + \exp(\mathbf{x}^i heta)}$$
  
 $\mathfrak{M}_2 : y_i \mid \mathbf{x}^i, \theta \sim \mathscr{B}(1, q_i) \quad \text{where} \quad q_i = \Phi(\mathbf{x}^i(\kappa^{-1} heta)),$ 

with  $\kappa^{-1}\theta = (\theta \mathbf{0}/k_{\mathbf{0}}, \theta_{\mathbf{1}}/k_{\mathbf{1}}).$ 

Under default g-prior

$$\theta \sim \mathscr{N}_2(0, n(X^{\mathsf{T}}X)^{-1})$$

full conditional posterior distributions given allocations

$$\pi(\theta \mid \mathbf{y}, X, \zeta) \propto \frac{\exp\left\{\sum_{i} \mathbb{I}_{\zeta_{i}=1} y_{i} \mathbf{x}^{i} \theta\right\}}{\prod_{i;\zeta_{i}=1} [1 + \exp(\mathbf{x}^{i} \theta)]} \exp\left\{-\theta^{T} (X^{T} X) \theta / 2n\right\}} \\ \times \prod_{i;\zeta_{i}=2} \Phi(\mathbf{x}^{i} (\kappa^{-1} \theta))^{y_{i}} (1 - \Phi(\mathbf{x}^{i} (\kappa^{-1} \theta)))^{(1-y_{i})}$$

hence posterior distribution clearly defined

#### Results

		Logistic		Probit	
a <sub>0</sub>	$\alpha$	$\theta_0$	$\theta_1$	$\frac{\theta_0}{k_0}$	$\frac{\theta_1}{k_1}$
.1	.352	-4.06	.103	-2.51	.064
.2	.427	-4.03	.103	-2.49	.064
.3	.440	-4.02	.102	-2.49	.063
.4	.456	-4.01	.102	-2.48	.063
.5	.449	-4.05	.103	-2.51	.064



Histograms of posteriors of  $\alpha$  in favour of logistic model where  $a_0 = .1, .2, .3, .4, .5$  for (a) Pima dataset, (b) Data from logistic model, (c) Data from probit model

27/40

# Survival analysis

Testing hypothesis that data comes from a

- $\ 0 \ \, \log\text{-Normal}(\phi,\kappa^2),$
- **2** Weibull $(\alpha, \lambda)$ , or
- $\ \ \, \hbox{log-Logistic}(\gamma,\delta)$

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- log-Normal( $\phi, \kappa^2$ ),
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distribution

Corresponding mixture given by the density

$$\alpha_{1} \exp\{-(\log x - \phi)^{2}/2\kappa^{2}\}/\sqrt{2\pi}x\kappa + \\ \alpha_{2}\frac{\alpha}{\lambda} \exp\{-(x/\lambda)^{\alpha}\}((x/\lambda)^{\alpha-1} + \\ \alpha_{3}(\delta/\gamma)(x/\gamma)^{\delta-1}/(1 + (x/\gamma)^{\delta})^{2}$$

where  $\alpha_1 + \alpha_2 + \alpha_3 = 1$ 

Looking for common parameter(s) :

$$\phi = \mu + \gamma\beta = \xi$$
  
$$\sigma^2 = \pi^2\beta^2/6 = \zeta^2\pi^2/3$$

where  $\gamma \approx 0.5772$  is Euler-Mascheroni constant.

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Allows for a noninformative prior on the common location scale parameter,

$$\pi(\phi,\sigma^2) = 1/\sigma^2$$

Recovery



Boxplots of the posterior distributions of the Normal weight  $\alpha_1$  under the two scenarii : truth = Normal *(left panel)*, truth = Gumbel *(right panel)*,  $a_0$ =0.01, 0.1, 1.0, 10.0 *(from left to right in each panel)* and n = 10,000 simulated observations.

Posterior consistency holds for mixture testing procedure [under minor conditions]

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Two different cases

- $\bullet$  the two models,  $\mathfrak{M}_1$  and  $\mathfrak{M}_2,$  are well separated
- model  $\mathfrak{M}_1$  is a submodel of  $\mathfrak{M}_2$ .

I. Posterior concentration rate :  $f_{\theta,\alpha} = \alpha f_{1,\theta_1} + (1-\alpha) f_{2,\theta_2}$ 

Let  $\pi$  be the prior and  $\mathbf{x}^n = (x_1, \cdots, x_n)$  a sample with true density  $f^*$  proposition

Assume that, for all c > 0, there exist  $\Theta_n \subset \Theta_1 \times \Theta_2$  and B > 0 such that

 $\pi \left[\Theta_n^c\right] \le n^{-c}, \quad \Theta_n \subset \{\|\theta_1\| + \|\theta_2\| \le n^B\}$ 

and that there exist  $H \ge 0$  and  $L, \delta > 0$  such that, for j = 1, 2,

$$\begin{split} \sup_{\theta, \theta' \in \Theta_{\mathbf{n}}} \|f_{j,\theta_{j}} - f_{j,\theta'_{j}}\|_{1} &\leq L n^{H} \|\theta_{j} - \theta'_{j}\|, \quad \theta = (\theta_{1}, \theta_{2}), \, \theta' = (\theta'_{1}, \theta'_{2}), \\ \forall \|\theta_{j} - \theta^{*}_{j}\| &\leq \delta; \quad \mathsf{KL}(f_{j,\theta_{j}}, f_{j,\theta^{*}_{j}}) \lesssim \|\theta_{j} - \theta^{*}_{j}\|. \end{split}$$

Then, when  $f^* = f_{ heta^*, \alpha^*}$ , with  $\alpha^* \in [0, 1]$ , there exists M > 0 such that

$$\pi \left| (\alpha, \theta); \| f_{\theta, \alpha} - f^* \|_1 > M \sqrt{\log n/n} |\mathbf{x}^n| = o_p(1).$$

32/40

II. Recovery of the parameters : Separated models –  $f_{ heta,lpha} = lpha f_{1, heta_1} + (1-lpha) f_{2, heta_2}$ 

Assumption : Models are separated, i.e. identifiability holds :

$$\forall \alpha, \alpha' \in (0, 1), \quad \forall \theta_j, \theta_j', j = 1, 2 \quad f_{\theta, \alpha} = f_{\theta', \alpha'} \quad \Rightarrow \alpha = \alpha', \quad \theta = \theta'$$

Further

$$\begin{split} \inf_{\substack{\theta_1 \in \Theta_1 \\ \theta_2 \in \Theta_2}} \inf_{\substack{\theta_2 \in \Theta_2 \\ \theta_1 \in \Theta_2}} \|f_{1,\theta_1} - f_{2,\theta_2}\|_1 > 0 \\ \end{split}$$
 and, for  $\theta_j^* \in \Theta_j$ , if  $P_{\theta_j}$  weakly converges to  $P_{\theta_j^*}$ , then  $\theta_j \longrightarrow \theta_j^*$ 

in the Euclidean topology

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#### theorem

Under above assumptions, then for all  $\epsilon > 0$ ,

 $\pi\left[|\alpha - \alpha^*| > \epsilon | \mathbf{x}^n\right] = o_p(1)$ 

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#### theorem

#### lf

•  $\theta_j \rightarrow f_{j,\theta_j}$  is  $\mathscr{C}^2$  around  $\theta_j^*$ , j = 1, 2, •  $f_{1,\theta_1^*} - f_{2,\theta_2^*}, \nabla f_{1,\theta_1^*}, \nabla f_{2,\theta_2^*}$  are linearly independent in y and • there exists  $\delta > 0$  such that  $\nabla f_{1,\theta_1^*}, \nabla f_{2,\theta_2^*}, \sup_{|\theta_1 - \theta_1^*| < \delta} |D^2 f_{1,\theta_1}|, \sup_{|\theta_2 - \theta_2^*| < \delta} |D^2 f_{2,\theta_2}| \in L_1$ 

then

$$\pi\left[|\alpha - \alpha^*| > M\sqrt{\log n/n} | \mathbf{x}^n\right] = o_p(1).$$

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$$\forall \alpha, \alpha' \in (0, 1), \quad \forall \theta_{j}, \theta_{j}^{'}, j = 1, 2 \quad f_{\theta, \alpha} = f_{\theta^{'}, \alpha^{'}} \quad \Rightarrow \alpha = \alpha^{'}, \quad \theta = \theta^{'}$$

theorem allows for interpretation of  $\alpha$  under the posterior : If data  $\mathbf{x}^n$  is generated from model  $\mathfrak{M}_1$  then posterior on  $\alpha$  concentrates around  $\alpha = 1$ 

Here  $\mathfrak{M}_1$  is a submodel of  $\mathfrak{M}_2$ , i.e.

$$\theta_2 = (\theta_1, \psi)$$
 and  $\theta_2 = (\theta_1, \psi_0 = 0)$ 

corresponds to  $f_{2,\theta_2} \in \mathfrak{M}_1$ Same posterior concentration rate

 $\sqrt{\log n/n}$ 

for estimating  $\alpha$  when  $\alpha^* \in (0, 1)$  and  $\psi^* \neq 0$ .

- Case where  $\psi^* =$  0, i.e.,  $f^*$  is in model  $\mathfrak{M}_1$
- Two possible paths to approximate  $f^*$  : either  $\alpha$  goes to 1 (path 1) or  $\psi$  goes to 0 (path 2)
- New identifiability condition :  $P_{\theta,\alpha} = P^*$  only if

$$\alpha=1, \ \theta_1=\theta_1^*, \ \theta_2=(\theta_1^*,\psi) \quad \text{or} \quad \alpha\leq 1, \ \theta_1=\theta_1^*, \ \theta_2=(\theta_1^*,0)$$

- Case where  $\psi^* =$  0, i.e.,  $f^*$  is in model  $\mathfrak{M}_1$
- Two possible paths to approximate f\* : either α goes to 1 (path 1) or ψ goes to 0 (path 2)
- New identifiability condition :  $P_{\theta,\alpha} = P^*$  only if

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Prior

$$\pi(\alpha, \theta) = \pi_{\alpha}(\alpha)\pi_1(\theta_1)\pi_{\psi}(\psi), \quad \theta_2 = (\theta_1, \psi)$$

with common (prior on)  $\theta_1$ 

• [B1] Regularity : Assume that  $\theta_1 \to f_{1,\theta_1}$  and  $\theta_2 \to f_{2,\theta_2}$  are 3 times continuously differentiable and that

$$F^*\left(\frac{\sup_{|\theta_1-\theta_1^*|<\delta}|D^rf_{1,\theta_1^*}|^s}{\underline{f}_{1,\theta_1^*}^s}\right)<+\infty, \quad r\leq 3$$

• [B2] Integrability :  $\exists \mathscr{S}_0 \subset \mathscr{S} \cap \{ |\psi| > \delta_0 > 0 \}$  s.t. Leb $(\mathscr{S}_0) > 0$ , and s.t.  $\forall \psi \in \mathscr{S}_0$ ,

$$F^*\left(\frac{\sup_{|\theta_1-\theta_1^*|<\delta}f_{2,\theta_1,\psi}}{f_{1,\theta_1^*}^4}\right)<+\infty,\quad F^*\left(\frac{\sup_{|\theta_1-\theta_1^*|<\delta}f_{2,\theta_1,\psi}^3}{\underline{f}_{1,\theta_1^*}^4}\right)<+\infty,$$

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#### Assumptions

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[B3] Stronger identifiability : Set

$$\nabla f_{2,\theta_{\mathbf{1}}^*,\psi^*}(x) = \left(\nabla_{\theta_{\mathbf{1}}}f_{2,\theta_{\mathbf{1}}^*,\psi^*}(x)^\mathsf{T}, \nabla_{\psi}f_{2,\theta_{\mathbf{1}}^*,\psi^*}(x)^\mathsf{T}\right)^\mathsf{T}.$$

Then for all  $\psi \in \mathscr{S}$  with  $\psi \neq 0$ , if  $\eta_0 \in \mathbb{R}$ ,  $\eta_1 \in \mathbb{R}^{d_1}$ 

$$\eta_{0}(f_{1,\theta_{1}^{*}}-f_{2,\theta_{1}^{*},\psi})+\eta_{1}^{\mathsf{T}}\nabla_{\theta_{1}}f_{1,\theta_{1}^{*}}=0 \quad \Leftrightarrow \eta_{1}=0, \ \eta_{2}=0$$

#### theorem

In the model :  $f_{\theta_1,\psi,\alpha} = \alpha f_{1,\theta_1} + (1-\alpha) f_{2,\theta_1,\psi}$  and  $\mathbf{x}^n = (x_1, \cdots, x_n) \stackrel{i.i.d}{\sim} f_{1,\theta_1^*}$ , If B1 - B3 hold , then  $\pi \left[ (\alpha, \theta); \| f_{\theta,\alpha} - f^* \|_1 > M \sqrt{\log n/n} |\mathbf{x}^n \right] = o_p(1).$ If  $\alpha \sim \mathscr{B}(a_1, a_2)$ , with  $a_2 < d_2$ , and if the prior dens.  $\pi_{\theta_1,\psi}$  is  $C^o$  and > 0 at  $(\theta_1^*, 0)$ , then  $M_n \longrightarrow \infty$ 

 $\pi \left[ \alpha < 1 - M_n (\log n)^{\gamma} / \sqrt{n} | \mathbf{x}^n \right] = o_p(1), \quad \gamma = \max((d_1 + a_2) / (d_2 - a_2), 1) / 2,$ 

When the true model behind the data is neither of the tested models, what happens  $? \end{tabular}$ 

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- issue mostly bypassed by classical Bayesian procedures
- ${\scriptstyle \bullet}$  theoretically produces an  $\alpha^*$  away from both 0 and 1
- possible (recommended ?) inclusion of a Bayesian non-parametric model within alternatives

And if we have to make a decision?

soft consider behaviour of posterior under prior predictives

- or posterior predictive [e.g., prior predictive does not exist]
- boostrapping behaviour
- comparison with Bayesian non-parametric solution

hard rethink the loss function

Thank You



40/40