

Combining multiple sources of information with Markov melding

Examples, challenges, and (partial) solutions

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Model criticism - 1

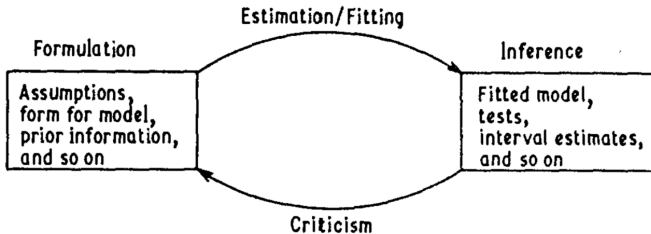


Figure 1.2.1 Schematic outline of model fitting. Adapted from Box (1979, 1980)

- From Cook and Weisberg (1982), but really from Box (1979; 1980)

Model criticism - 2

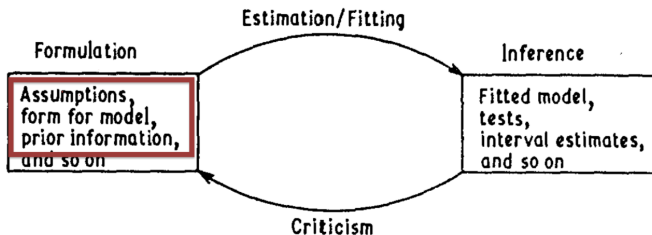


Figure 1.2.1 Schematic outline of model fitting. Adapted from Box (1979, 1980)

- From Cook and Weisberg (1982), but really from Box (1979; 1980)
- Non-linear increase in difficulty when $n, p \rightarrow \infty$

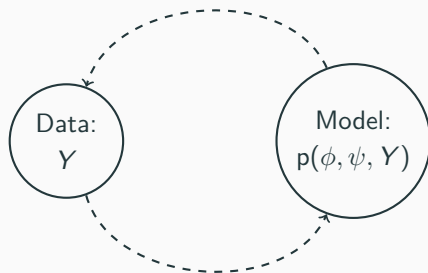
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SYNTHESISING EVIDENCE TO ESTIMATE PANDEMIC (2009) A/H1N1 INFLUENZA SEVERITY IN 2009–2011¹

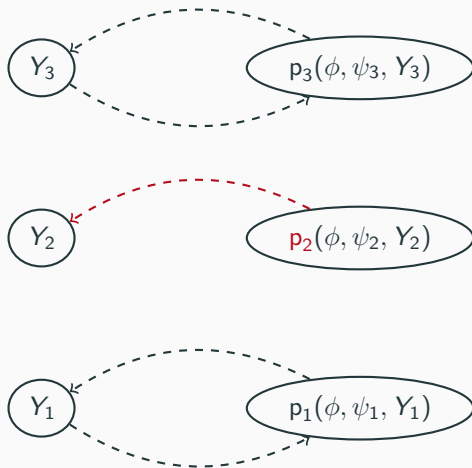
BY ANNE M. PRESANIS^{*}, RICHARD G. PEBODY[†], PAUL J. BIRRELL^{*},
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DOUGLAS FLEMING[‡] AND DANIELA DE ANGELIS^{*}

Medical Research Council Biostatistics Unit^{}, Public Health England[†]
and Royal College of General Practitioners[‡]*

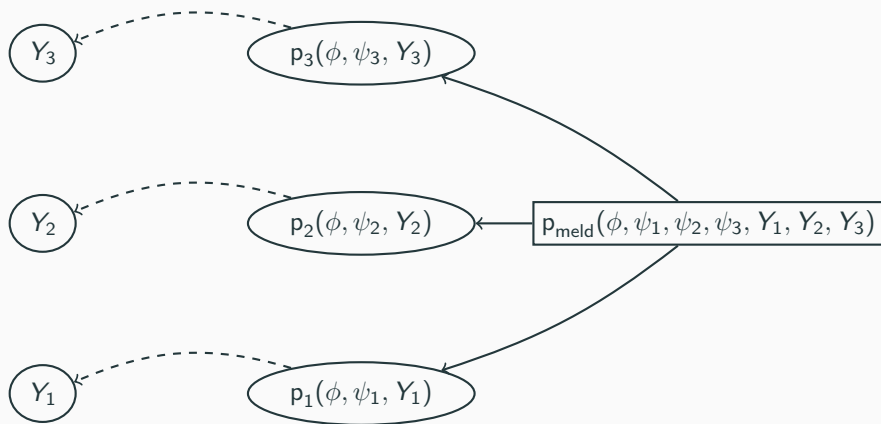
- 5 sources of data, each idiosyncratic.
 - Some known to be biased in particular directions
 - Different levels of aggregation, national / hospital specific data
- Specifying and critiquing a joint model for all of this?
- Will return to a simplified version



Statistical submodelling



Melding based statistical modelling?



Specification: Markov melding

- Markov melding (Goudie et al. 2019) is a method for joining models with common component: ϕ
- Ideally we would specify the generative model (For $M = 2$ submodels):

$$p_{\text{meld}}(\phi, \psi_1, \psi_2, Y_1, Y_2) = p_{\text{pool}}(\phi) p_1(\psi_1, Y_1 \mid \phi) p_2(\psi_2, Y_2 \mid \phi)$$

- Submodels developed in isolation, re-express as:

$$p_{\text{meld}}(\phi, \psi_1, \psi_2, Y_1, Y_2) = p_{\text{pool}}(\phi) \frac{p_1(\phi, \psi_1, Y_1)}{\mathbf{p}_1(\phi)} \frac{p_2(\phi, \psi_2, Y_2)}{\mathbf{p}_2(\phi)}$$

- Analytic form of $\mathbf{p}_m(\phi)$ often unknown
 - Can cause numerical issues in the model fitting process
 - Requires accurate estimation in low probability regions
 - We will revisit

Specification: Notes on the pooled prior

$p_{\text{pool}}(\phi)$ addresses two issues:

1. Necessary because we have two priors for ϕ (Poole and Raftery 2000)
2. $p_{\text{pool}}(\phi)$ should be appropriate for ϕ under the melded model *and* both submodels
 - Suggests defining $p_{\text{pool}}(\phi) = g(p_1(\phi), p_2(\phi))$
 - Similar problem encountered when experts encode their opinions as priors (O'Hagan et al. 2006)
 - **Log:** $p_{\text{pool}}(\phi) = p_1(\phi)^{w_1} p_2(\phi)^{w_2}$
 - Linear: $p_{\text{pool}}(\phi) = w_1 p_1(\phi) + w_2 p_2(\phi)$
 - Product-of-experts: $p_{\text{pool}}(\phi) = p_1(\phi) p_2(\phi)$ (Hinton 2002)
 - Dictatorial: $p_{\text{pool}}(\phi) = p_m(\phi)$ for an $m \in \{1, 2\}$

Estimation: Multi-stage sampler

- Need to sample the melded posterior:

$$\begin{aligned} p_{\text{meld}}(\phi, \psi_1, \psi_2 \mid Y_1, Y_2) &\propto p_{\text{meld}}(\phi, \psi_1, \psi_2, Y_1, Y_2) \\ &= p_{\text{pool}}(\phi) \frac{p_1(\phi, \psi_1, Y_1)}{p_1(\phi)} \frac{p_2(\phi, \psi_2, Y_2)}{p_2(\phi)} \end{aligned}$$

- Submodels can be complex \rightarrow sample in stages
 - Multi-stage sampling & approximations are common in practice
 - Staged sampling targets components of $p_{\text{meld}}(\phi, \psi_1, \psi_2 \mid Y_1, Y_2)$ in a cumulative manner
 - Can reuse other implementations of $p_m(\phi, \psi_m, Y_m)$
 - Potentially more efficient

Estimation: Stage one acceptance probability

- Say we choose to target, $p_{\text{meld},1}(\phi, \psi_1 \mid Y_1)$:

$$\begin{aligned} p_{\text{meld}}(\phi, \psi_1, \psi_2 \mid Y_1, Y_2) &\propto p_{\text{pool}}(\phi) \frac{p_1(\phi, \psi_1, Y_1)}{p_1(\phi)} \frac{p_2(\phi, \psi_2, Y_2)}{p_2(\phi)} \\ &\rightarrow p_{\text{meld},1}(\phi, \psi_1 \mid Y_1) \end{aligned}$$

- Given our stage one target, the stage one sampler has the following acceptance probability:

$$\alpha((\phi^*, \psi_1^*), (\phi, \psi_1)) = \frac{p_1(\phi^*, \psi_1^*, Y_1)p_1(\phi)Q(\phi, \psi_1 \mid \phi^*, \psi_1^*)}{p_1(\phi, \psi_1, Y_1)p_1(\phi^*)Q(\phi^*, \psi_1^* \mid \phi, \psi_1)}$$

- (asymptotically) Produces samples from the stage one target

Estimation: Stage two acceptance probability

- Stage two target, the melded posterior:

$$p_{\text{meld}}(\phi, \psi_1, \psi_2 \mid Y_1, Y_2) \propto p_{\text{pool}}(\phi) \frac{p_1(\phi, \psi_1, Y_1)}{p_1(\phi)} \frac{p_2(\phi, \psi_2, Y_2)}{p_2(\phi)}$$

- Use stage one samples of ϕ, ψ_1 as the proposal distribution in a Metropolis-within-Gibbs update for $\phi, \psi_1 \mid \psi_2$:

$$\alpha((\phi^*, \psi_1^*), (\phi, \psi_1)) = \frac{p_{\text{pool}}(\phi^*)p_1(\phi^*, \psi_1^*, Y_1)p_2(\phi^*, \psi_2, Y_2)p_1(\phi)p_2(\phi)}{p_{\text{pool}}(\phi)p_1(\phi, \psi_1, Y_1)p_2(\phi, \psi_2, Y_2)p_1(\phi^*)p_2(\phi^*)} \frac{p_1(\phi, \psi_1, Y_1)p_1(\phi^*)}{p_1(\phi^*, \psi_1^*, Y_1)p_1(\phi)}$$

- Stage one terms cancel:

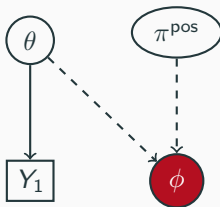
$$\alpha(\phi^*, \phi) = \frac{p_{\text{pool}}(\phi^*)p_2(\phi^*, \psi_2, Y_2)p_2(\phi)}{p_{\text{pool}}(\phi)p_2(\phi, \psi_2, Y_2)p_2(\phi^*)}$$

- Generic Metropolis-Hastings update for $\psi_2 \mid \phi, \psi_1$

Example: Simplified version of Presanis et. al. (2014)

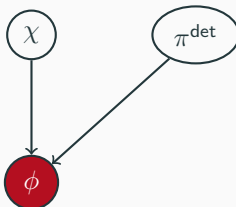
- We will now consider a simplified version of Presanis et. al. (2014)
- Submodel details are not overly important for our purpose
- Consider 2 submodels:
 1. Models number of patients in the ICU with H1N1
 2. Collapses the other 4 data sources into an informative prior structure

Example: Submodel 1 - ICU submodel



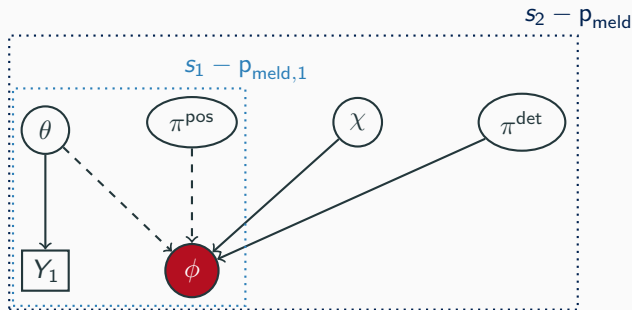
- Notation: $\psi_1 = (\theta, \pi^{\text{pos}})$
- $p_1(\theta, Y)$ is a thinned Poisson process as an observational model for Y
 - Can derive total number of people in ICU with influenza
- π^{pos} derived from virological data
 - Informs the proportion of people with influenza who are positive for H1N1
- Submodel deterministically defines $\phi = f(\theta, \pi^{\text{pos}})$.

Example: Submodel 2 - Simplified severity model



- Notation: $\psi_2 = (\chi, \pi^{\text{det}})$
- $\phi \sim \text{Bin}(\chi, \pi_{\text{det}})$ to account for known underestimation of ϕ
- Informative prior for χ summarises other components of severity model

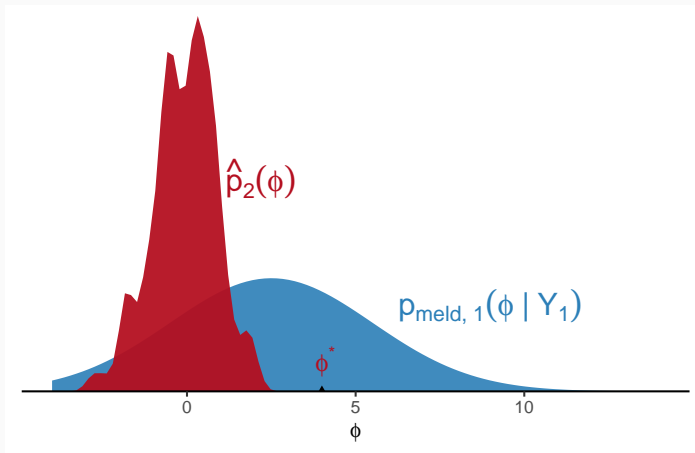
Example: Multi-stage sampler - in DAG form



- After stage 2 (s_2), every node's distribution incorporates information from Y_1 and χ .

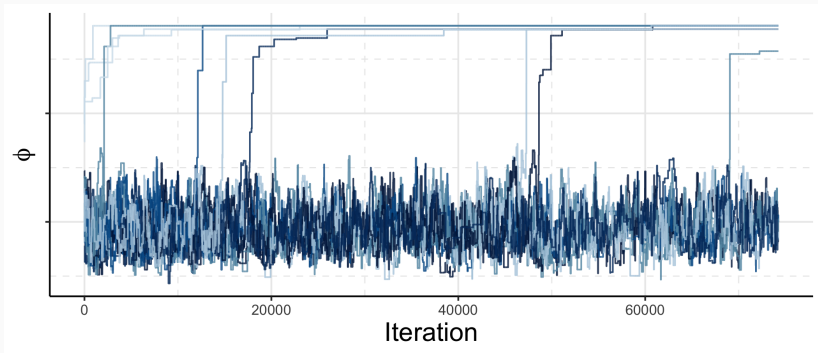
Issue 1: Things don't always work!

$$\alpha(\phi^*, \phi) = \frac{p_{\text{pool}}(\phi^*)p_2(\phi^*, \psi_2, Y_2)}{p_{\text{pool}}(\phi)p_2(\phi, \psi_2, Y_2)} \cdot \frac{\hat{p}_2(\phi)}{\hat{p}_2(\phi^*)}$$



Issue 1: KDE error in action

$$\alpha(\phi^*, \phi) = \frac{p_{\text{pool}}(\phi^*)p_2(\phi^*, \psi_2, Y_2)}{p_{\text{pool}}(\phi)p_2(\phi, \psi_2, Y_2)} \cdot \frac{\hat{p}_2(\phi)}{\hat{p}_2(\phi^*)}$$



Method 1: Self density ratios

We only interact with unknown marginal via self-density ratio (Hiraoka, Hamada, and Hori 2018):

$$r(\phi_{\text{nu}}, \phi_{\text{de}}) = \frac{p(\phi_{\text{nu}})}{p(\phi_{\text{de}})}$$

Weighted-sample self-density ratio estimation (WSRE) (Manderson and Goudie 2020) Intuition:

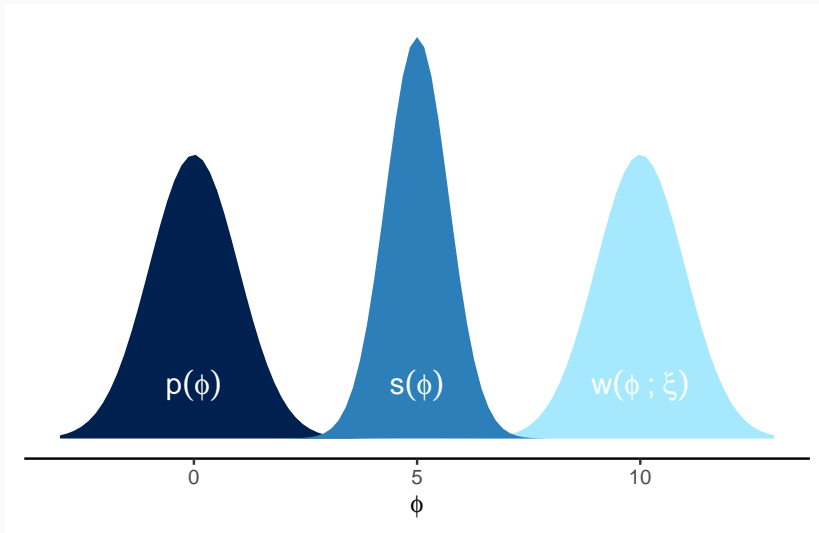
- Sampling $\phi \sim p(\phi, \psi, Y)$ admits $\phi \sim p(\phi)$
- Instead, sample:
 $\phi \sim p(\phi, \psi, Y)w(\phi; \xi) \rightarrow \phi \sim \frac{1}{Z}p(\phi)w(\phi; \xi) = s(\phi)$
- Use the weighted sample density estimator of Jones (1991):

$$\hat{p}(\phi) = \frac{1}{ZNh} \sum_{i=1}^N w(\phi; \xi)^{-1} K_h(\phi - \phi_i)$$

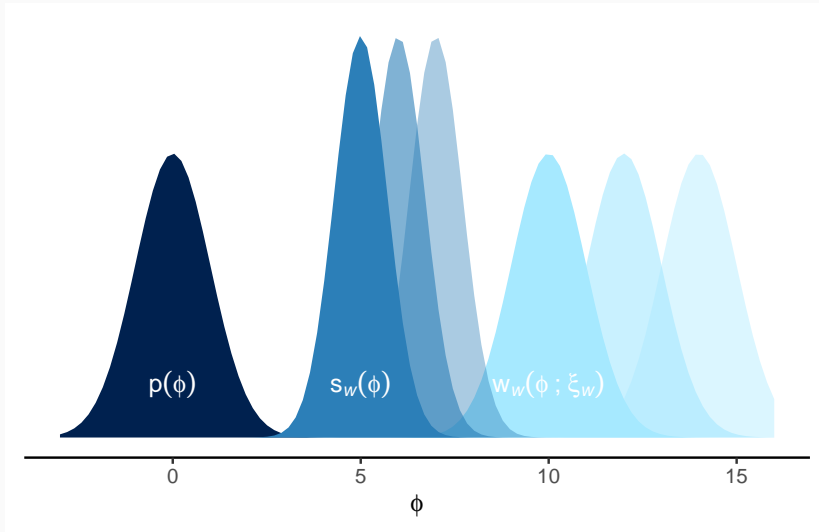
- Ratio estimator is then:

$$\hat{r}(\phi_{\text{nu}}, \phi_{\text{de}}) = \frac{\hat{p}(\phi_{\text{nu}})}{\hat{p}(\phi_{\text{de}})}$$

Method 1: WSRE intuition - 1

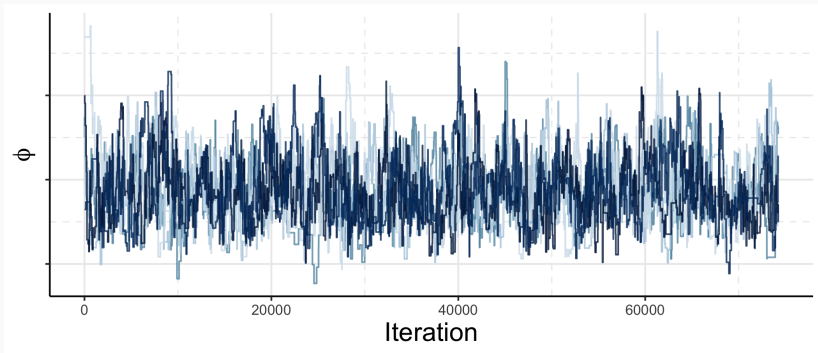


Method 1: WSRE intuition - W

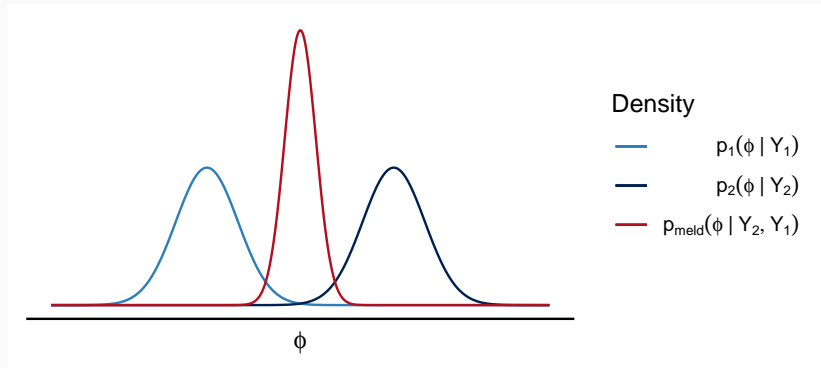


Method 1: H1N1 example - 2

$$\alpha(\phi^*, \phi) = \frac{p_{\text{pool}}(\phi^*) p_2(\phi^*, \psi_2, Y_2)}{p_{\text{pool}}(\phi) p_2(\phi, \psi_2, Y_2)} \cdot \hat{r}(\phi, \phi^*)$$

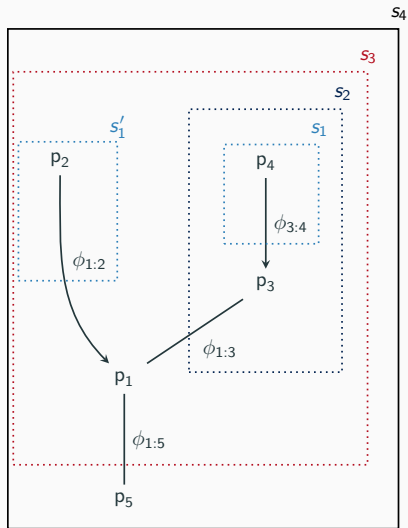


Issue 2: Submodel conflict



- Last issue arose (primarily) due to differences in scale
- What about more general model conflict?
 - There is a philosophical issue here
- *Partial solution:* Automatic, multiple importance sampling (Paananen et al. 2019)

Issue 3: Multiple ϕ ?



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A trio of inference problems that could win you a Nobel Prize in statistics (if you help fund it)

Xiao-Li Meng

Department of Statistics

Harvard University, Cambridge, MA

- Multi-source inference is extremely difficult in practice (Meng 2014) (along with multi-resolution and multi-phase)

Should you also wish to estimate self-density ratios:

- <https://github.com/hhau/wsre>

The traceplots are from this example:

- <https://github.com/hhau/full-melding-example>

This talk:

**TODO: Push this branch to
github / link to it**

-

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