Markov Melding Notes

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0. Notation

• Captial X is a random variable, \mathbf{X} is a vector of random variables and x and \mathbf{x} are their realisations

1. Background

1.1 Monte Carlo

(Johansen, 2018)

For the following samplers targeting density f and starting with $x^{(0)} := (x_1^{(0)}, \dots, x_p^{(0)})$, iterate for $t = 1, 2, \dots$

1.1.1. Metropolis-Hastings sampler

- 1. Draw $x \sim q\left(\cdot|x^{(t-1)}\right)$
- 2. With probability min $\left\{1, \frac{f(x) \cdot q(x^{(t-1)}|x)}{f(x^{(t-1)}) \cdot q(x|x^{(t-1)})}\right\}$ set $x^{(t)} = x$, else set $x^{(t)} = x^{(t-1)}$

Note that if the proposal q is symmetric (as in random-walk metropolis-hastings) then the acceptance probability simplifies to min $\left\{1, \frac{f(x)}{f(x^{(t-1)})}\right\}$.

1.1.2. (Random scan) Gibbs sampler

- 1. Draw $j \sim \text{Unif}\{1, \dots, p\}$ 2. Draw $x_j^{(t)} \sim f_{x_j|x_{-j}}\left(\cdot | x_1^{(t-1)}, \dots, x_{j-1}^{(t-1)}, x_{j+1}^{(t-1)}, \dots, x_p^{(t-1)}\right)$, and set $x_i^{(t)} := x_i^{(t-1)}$ for all $i \neq j$

1.1.3. (Random scan) Metropolis-within-Gibbs

- 1. Draw $j \sim \text{Unif}\{1,\ldots,p\}$
- 2. a) Draw $x_j \sim q_j\left(\cdot|x^{(t-1)}\right)$ and set $x = \left(x_1^{(t-1)}, \dots, x_j, \dots, x_p^{(t-1)}\right)$ b) With probability min $\left\{1, \frac{f(x) \cdot q\left(x^{(t-1)}|x\right)}{f\left(x^{(t-1)}\right) \cdot q\left(x|x^{(t-1)}\right)}\right\}$ set $x^{(t)} = x$, else set $x^{(t)} = x^{(t-1)}$

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1.2. Meta-analysis, evidence synthesis, combining expert opinion etc.

1.2.1 (O'Hagan 2006) Chapter 9: Multiple Experts

- Want to obtain a single distribution which encapsulates the beliefs of several experts
- Two approaches
 - Mathematical aggregation: elicit distribution from each expert individually and independently then mathematically combine
 - Behavioural aggregation: Create an interaction beween the group of experts through which a single distribution is elicited from the group as a whole
- In reference to Markov melding, it seems as though Behavioural approaches are not possible
- Each of a group of n experts asked individually for her beliefs about some unknown quantity θ , eliciting distributions $f_i(\theta)$ for i = 1, ..., n
- Formal Bayesian perspective (DM is a supra-Bayesian): DM begins with his own prior $f(\theta)$ for θ and has posterior $f(\theta|D)$ after incorporating the experts opinions $D = \{f_1(\theta), \dots, f_n(\theta)\}$. This is difficult as DM must construct likelihood $f(D|\theta)$
- Simpler and widely used technique is opinion pooling where a consensus distribution $f(\theta)$ is obtained as some function of the individual distributions $\{f_1(\theta), \ldots, f_n(\theta)\}$
- Linear opinion pool $f(\theta) \propto \sum_{i=1}^n w_i f_i(\theta)$ where the weights w_i sum to one
 - Could weight all experts equally $w_i = 1/n$ for all i
 - Alternatively, give more weight to some expert
 - Coherent marginalisation
 - This approach is not externally Bayesian: after recieving new information, updating the priors then pooling the result is not the same as updating the pooled prior
 - Not consistent with regard to judgements of independence
- Logarithmic opinion pool $f(\theta) \propto \prod_{i=1}^n f_i(\theta)^{w_i}$
 - Again can weight opinions as wishes
 - Externally Bayesian and consistent about independence
 - However, unlike linear, no coherent marginalisation (no pooling satisfies both externally Bayesian and coherent marginalisation)
- Product of Experts (special case of logarithmic pooling) $f(\theta) \propto \prod_{i=1}^n f_i(\theta)$ To-do: look at Hinton 2002
- Unlike supra-Bayesian approach, the result of opinion pooling does not represent the actual beliefs of any individual as so may not behave as one would expect a probability distribution to
- "In general, while the linear opinion pool has been quite widely used in practice, the logarithmic opinion pool has been largely ignored, perhaps because it is perceived to lead to unrealistically strong aggregated beliefs"
- Dictatorial pooling $f(\theta) = f_i(\theta)$ for some i = 1, ..., n (not mentioned in book)
- Cooke's method
- For more about opinion pooling, could look at (Clemen, 1999)

1.3. Graphical models

1.3.1 (Smith, 2010) Chapter 7: Bayesian networks

1.3.1.1 Relevance, informativeness and independence

- Client believes that measurement X is irrelevant for predicting Y given the measurement Z, written $Y \perp X | Z$ if she believes now that once she learns the value of Z then then measurement of X will provide her with no extra useful information with which to predict the value of Y.
- In this case, if she is a Bayesian, then she could write her conditional density p(y|x,z) = p(y|z) so that it did not depend on the value of x for all possible values of (x,y,z). Equivalently the joint mass function can be factorised as p(x,y,z) = p(y|z)p(x|z)p(z)
- Two most important and universally applicable rules:
 - Symmetry $Y \perp X|Z \iff X \perp Y|Z$
 - Perfect composition $X \perp (Y,Z)|W \iff X \perp Y|(W,Z) \iff X \perp Z|W$
- To-do: look at some work of Pearl?

1.3.1.2. Bayesian networks and DAGs

- Bayesian network is a simple and convenient way of representing a factorisation of a joint pdf of a vector of random variables $\mathbf{X} = (X_1, X_2, \dots, X_n)$.
- Always the case that $p(\mathbf{x}) = p(x_1)p(x_2|x_1)p(x_3|x_2,x_1) \times \cdots \times p(x_n|x_1,x_2,\ldots,x_{n-1})$
- Often many of the functions $p(x_i|x_1, x_2, ..., x_{i-1})$ are explicit functions of components of X whose indices lie in a proper subset $Q_i \subset \{1, 2, ..., i-1\}$ so that $p(x_i|x_1, x_2, ..., x_{i-1}) = p(x_i|\mathbf{x}_{Q_i})$
- Now $p(\mathbf{x}) = p(x_1) \prod_{i=2}^n p(x_i | \mathbf{x}_{Q_i})$
- Let the remainder set $R_i = \{1, 2, ..., i-1\}$ Q_i then the above bullet point is equivalent to the set of n-1 irrelevance statements $X_i \perp \mathbf{X}_{R_i} | \mathbf{X}_{Q_i}, 2 \leq i \leq n$
- Definition: A directed acyclic graph (DAG) $\mathcal{G} = (\mathcal{V}, \mathcal{E})$ with set of vertices \mathcal{V} and set of directed edges \mathcal{E} is a directed graph having no directed cycles
- Definition: A Bayesian network (BN) on the set of measurements $\{X_1, X_2, ..., X_n\}$ is a set of the n-1 conditional irrelevance statements together with a DAG \mathcal{G} . The set of vertices $\mathcal{G} = \{X_1, X_2, ..., X_n\}$ and a directed edge from X_i to X_j is in \mathcal{E} if and only if $i \in Q_j$

1.3.2 Factor graphs

• Factor graphs (Bishop 2016): "Factor graphs make this decomposition explicit by introducing additional nodes for the factors themselves in addition to the nodes representing the variables."

1.4. Misc. reading

1.4.1. (Spiegelhalter, 1995)

- Meta analysis, also known as systematic overview, is a statistical procedure in which the results of several independent studies are integrated. Aim is to resolve issues that cannot be concluded from a single study alone
- Note: Gelman blog post about why fixed and random effects terminology is confusing
- Fixed-effect analysis: a common effect across studies is estimated
- Random-effects model: a probability model for individual study effects is assumed
- "In contrast to the pooled estimate arising from a fixed-effect model, a random-effects metaanalysis assumes that the true effects in each trial are not necessarily equal, but are random observations drawn from some common population distribution"

Let r_i^C denote the number of infections in the control group in trial i, arising from n_i^C cases each assumed to have probability p_i^C of developing an infection. Adopt equivalent notation for the treatment group and assume that δ_i is the true treatment effect on a log-odds scale such that

$$\delta_i = \operatorname{logit}(p_i^T) - \operatorname{logit}(p_i^C).$$

The "average" infection rate in the ith trial is

$$\mu_i = \frac{1}{2}(\operatorname{logit}(p_i^T) + \operatorname{logit}(p_i^C)).$$

Individual trial effects are drawn from some Gaussian population with mean d and variance σ^2 , giving the full model as

$$r_i^C \sim \text{Binomial}\left(p_i^C, n_i^C\right)$$

$$r_i^T \sim \text{Binomial}\left(p_i^T, n_i^T\right)$$

$$\text{logit}(p_i^C) = \mu_i - \delta_i/2$$

$$\text{logit}(p_i^T) = \mu_i + \delta_i/2$$

$$\delta_i \sim \text{Normal}(d, \sigma^2)$$

- Empirical Bayes' methods estimate d and σ^2 from the data by moment-matching, then make inferences conditional on these estimates \hat{d} , $\hat{\sigma}^2$ but this does not propigate uncertainty about the estimates. Instead full Bayes puts priors on the unknown parameters
- Based on suitable DAG, see Figure 1., the joint distribution is $p(V) = \prod_{v \in V} p(v|pa(v))$
- Use Gibbs sampling (BUGS) to perform inference

2. (Goudie, 2018) Markov Melding

2.1. Introduction to Markov melding

- Motivation: by using all available data typically get:
 - More precise estimates
 - More accurate reflection of true uncertainty
 - Minimise risk of selection-type biases
- Modular approaches
 - 1. Plug in a point estimate
 - Very easy and fast
 - Underestimates uncertainty
 - 2. Plug in an approximation to the posterior
 - Quite easy and fast
 - Assumptions made can be unclear
 - 3. Integrate the models
 - All uncertainty propagated
 - All assumptions explicit

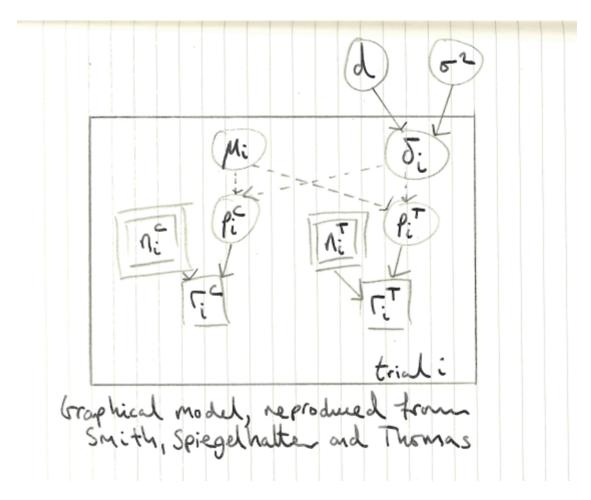


Figure 1: To-do: reproduce this and similar models in <math>Tikz Solid line is stochastic dependence, dashed line is logical dependence

- Probably tricky to do
- Aims of work:
 - 1. Join submodels p_m into a single joint model
 - Must implicitly handle two different priors for same quantity
 - Must handle non-invertible deterministic transformations
 - 2. Fit the submodels one at a time
 - Minimize burden on practitioners
 - 3. Understanding of reverse operation to joining splitting
- Models m = 1, ..., M each with joint density $p_m(\phi, \psi_m, Y_m)$ where:
 - $-\phi$ is the common parameter linking the models
 - $-\psi_m$ are model specific unobserved parameters
 - $-Y_m$ are model specific observed quantities
- Join together to create $p(\phi, \psi_1, \dots, \psi_M, Y_1, \dots, Y_M)$

2.2. Markov combination

To-do: add references to Dawid, Massa

Suppose that the marginals are consistent i.e. $p_m(\phi) = p(\phi)$ for all m then one can define the Markov combination p_{comb} of submodels p_1, \ldots, p_m as

$$\begin{aligned} p_{\text{comb}}\left(\phi,\psi_{1},\ldots,\psi_{M},Y_{1},\ldots,Y_{M}\right) &= p(\phi) \prod_{m=1}^{M} p_{m}\left(\psi_{m},Y_{m}|\phi\right) \\ &= p(\phi) \prod_{m=1}^{M} \frac{p_{m}\left(\phi,\psi_{m},Y_{m}\right)}{p(\phi)} \\ &= \frac{\prod_{m=1}^{M} p_{m}\left(\phi,\psi_{m},Y_{m}\right)}{p(\phi)^{M-1}} \end{aligned}$$

2.3. Pooling marginal distributions

If the marginals are inconsistent then instead a pooled density $p_{\text{pool}}(\phi) = g(p_1(\phi), \dots, p_M(\phi))$ can be used instead. Similar idea to combining expert opinions. This suggests the joint model

$$p_{\text{meld}}(\phi, \psi_1, \dots, \psi_M, Y_1, \dots, Y_M) = p_{\text{pool}}(\phi) \prod_{m=1}^M p_m(\psi_m, Y_m | \phi)$$
$$= p_{\text{pool}}(\phi) \prod_{m=1}^M \frac{p_m(\phi, \psi_m, Y_m)}{p_m(\phi)}$$

Goudie calls this Markov melding.

2.4. Inference and computation

Joint posterior, given data $Y_m = y_m$ for $m = 1, \dots, M$, under Melded model is

$$p_{\text{meld}}(\phi, \psi_1, \dots, \psi_M | y_1, \dots, y_M) \propto p_{\text{pool}}(\phi) \prod_{m=1}^M \frac{p_m(\phi, \psi_m, y_m)}{p_m(\phi)}$$

Metropolis-Hastings candidate values $(\phi^*, \psi_1^*, \dots, \psi_M^*)$ drawn from a proposal $q(\phi^*, \psi_1^*, \dots, \psi_M^* | \phi, \psi_1, \dots, \psi_M)$ and accepted with probability min(1, r) where

$$r = \frac{R\left(\phi^{\star}, \psi_{1}^{\star}, \dots, \psi_{M}^{\star}, \phi, \psi_{1}, \dots, \psi_{M}\right)}{R\left(\phi, \psi_{1}, \dots, \psi_{M}, \phi^{\star}, \psi_{1}^{\star}, \dots, \psi_{M}^{\star}\right)}$$

where $R(\phi^{\star}, \psi_1^{\star}, \dots, \psi_M^{\star}, \phi, \psi_1, \dots, \psi_M)$ is the target-to-proposal density ratio

$$R\left(\phi^{\star}, \psi_{1}^{\star}, \dots, \psi_{M}^{\star}, \phi, \psi_{1}, \dots, \psi_{M}\right) = p_{\text{pool}}\left(\phi^{\star}\right) \prod_{m=1}^{M} \frac{p_{m}\left(\phi^{\star}, \psi_{m}^{\star}, y_{m}\right)}{p_{m}\left(\phi^{\star}\right)} \times \frac{1}{q\left(\phi^{\star}, \psi_{1}^{\star}, \dots, \psi_{M}^{\star} \middle| \phi, \psi_{1}, \dots, \psi_{M}\right)}$$

2.4.1. Metropolis-within-Gibbs

Sample from the full conditionals using Metropolis-Hastings.

For each of the latent parameter updates $(\psi_m \text{ for } m = 1, \dots, M)$ we have

$$R\left(\phi,\psi_{1},\ldots,\psi_{m}^{\star},\ldots,\psi_{M},\phi,\psi_{1},\ldots,\psi_{M}\right)=p_{\text{pool}}\left(\phi\right)\prod_{j\neq m}\frac{p_{j}\left(\phi,\psi_{j},y_{j}\right)}{p_{j}\left(\phi\right)}\times\frac{p_{m}\left(\phi,\psi_{m}^{\star},y_{m}\right)}{p_{m}\left(\phi\right)}\frac{1}{q\left(\psi_{m}^{\star}|\psi_{m}\right)}$$

so that

$$r = \frac{p_m \left(\phi, \psi_m^{\star}, y_m\right) \times \frac{1}{q(\psi_m^{\star} | \psi_m)}}{p_m \left(\phi, \psi_m, y_m\right) \times \frac{1}{q(\psi_m | \psi_m^{\star})}}$$

and for the link parameter update

$$R\left(\phi, \psi_{1}, \dots, \psi_{m}^{\star}, \dots, \psi_{M}, \phi, \psi_{1}, \dots, \psi_{M}\right) = p_{\text{pool}}\left(\phi^{\star}\right) \prod_{m=1}^{M} \frac{p_{m}\left(\phi^{\star}, \psi_{m}, y_{m}\right)}{p_{m}\left(\phi^{\star}\right)} \times \frac{1}{q\left(\phi^{\star} | \phi\right)}$$

2.4.2. Multi-stage Metropolis-within-Gibbs

Factorise the pooled prior (can be done in many ways)

$$p_{\text{pool}}(\phi) = \prod_{m=1}^{M} p_{\text{pool},m}(\phi)$$

Define lth stage posterior as

$$p_{\text{meld},l}\left(\phi,\psi_1,\ldots,\psi_\ell|y_1,\ldots,y_\ell\right) \propto \prod_{m=1}^{\ell} \left(\frac{p_m\left(\phi,\psi_m,y_m\right)}{p_m(\phi)}p_{\text{pool},m}(\phi)\right)$$

Basis obtain samples $\left(\phi^{(h,1)}, \psi_1^{(h,1)}\right)$ for $h = 1, \dots, H_1$ from $p_{\text{meld},1}\left(\phi, \psi_1 | y_1\right)$ (by MCMC typically)

Inductive construct a Metropolis-within-Gibbs sampler for $(\phi, \psi_1, \dots, \psi_\ell)$ given the data (y_1, \dots, y_ℓ)

3. (Lindsten, 2017) D&C-SMC

To-do

4. Example: Gambia Malaria Data

The gambia dataset from the R package geoR contains observations of n=2035 Gambian children. The eight variables measured are:

- \bullet x the x-coordinate of the village (Universal Transverse Mercator similar to latitude and longitude)
- y the y-coordinate of the village (UTM)
- pos presence (1) or absence (0) of malaria in a blood sample taken from the child
- age age of the child, in days
- netuse indicator variable denoting whether (1) or not (0) the child regularly sleeps under a bed-net
- treated indicator variable denoting whether (1) or not (0) the bed-net is treated (coded 0 if netuse = 0)
- green satellite-derived measure of the green-ness of vegetation in the immediate vicinity of the village (arbitrary units)
- phc indicator variable denoting the presence (1) or absence (0) of a health center in the village

4.1 Logistic regression

Consider response $Y \in \{0,1\}$ modelled as $Y \sim \text{Bern}(q)$ and covariates $x \in \mathbb{R}^p$ with

$$\log\left(\frac{q(x)}{1-q(x)}\right) = \beta_0 + \beta^T x,$$

where $\beta \in \mathbb{R}^p$. Then

$$q(x) = \frac{\exp(\beta_0 + \beta^T x)}{1 + \exp(\beta_0 + \beta^T x)}$$

```
# Classify to 1 with probability
q <- function(x, b) {
  exp(b %*% x) / (1 + exp(b %*% x))
}</pre>
```

Observe labelled data $\mathcal{D} = \{(x_1, y_1), \dots, (x_n, y_n)\}$. The likelihood function is

$$\mathcal{L}(\beta_0, \beta) = \prod_{i=1}^{n} q(x_i)^{y_i} (1 - q(x_i))^{1 - y_i}$$

Place Gaussian priors on β and β_0 such that

$$\beta_0 \sim \mathcal{N}(\mu_0, \sigma_0^2), \ \beta \sim \mathcal{N}_p(\mu, \operatorname{diag}(\sigma_1^2, \dots, \sigma_p^2))$$

Then the posterior is proportional to

$$p(\beta_0, \beta | y_1, \dots, y_n) \propto \prod_{i=0}^{p} \exp\left(\frac{1}{2\sigma_j^2}(\beta_j - \mu_j)^2\right) \prod_{i=1}^{n} q(x_i)^{y_i} (1 - q(x_i))^{1-y_i}.$$

Taking the logarithm gives

$$\log p(\beta_0, \beta | y_1, \dots, y_n) \propto \sum_{j=0}^{p} \frac{1}{2\sigma_j^2} (\beta_j - \mu_j)^2 + \sum_{i=1}^{n} \{ y_i \log q(x_i) + (1 - y_i) \log (1 - q(x_i)) \}.$$

The log-likelihood can be rewritten as

$$\sum_{i=1}^{n} \{y_i \log q(x_i) + (1 - y_i) \log (1 - q(x_i))\} = \sum_{i=1}^{n} \{y_i \log \left(\frac{q(x_i)}{1 - q(x_i)}\right) + \log (1 - q(x_i))\}
= \sum_{i=1}^{n} \{y_i \log \left(\frac{q(x_i)}{1 - q(x_i)}\right) + \log (1 - q(x_i))\}
= \sum_{i=1}^{n} \{y_i \left(\beta_0 + \beta^T x\right) - \log \left(1 + \exp(\beta_0 + \beta^T x)\right)\},$$

so that the log-posterior is

$$\log p(\beta_0, \beta | y_1, \dots, y_n) \propto \sum_{j=0}^{p} \frac{1}{2\sigma_j^2} (\beta_j - \mu_j)^2 + \sum_{j=1}^{n} \{ y_i \left(\beta_0 + \beta^T x \right) - \log \left(1 + \exp(\beta_0 + \beta^T x) \right) \}$$

```
# (proportional to) log posterior in the indep normals prior case
logpost <- function(b, X, mu, sigma) {
  logprior <- sum((b - mu)^2 / 2*sigma)
  nu <- apply(X, 1, function(x) b %*% x) # Vector of linear predictors
  loglike <- sum(nu[Y == 1]) + sum(-log(1 + exp(nu)))
  logprior + loglike
}</pre>
```

4.2. Full and submodels

Firstly, the full model \mathcal{M} is the logistic regression of response pos on the other variables including an intercept term but excluding the co-ordinates x and y.

$$\log\left(\frac{q(x)}{1 - q(x)}\right) = \eta$$

$$\eta = \beta_0 + \beta_1 \cdot \mathtt{age} + \beta_2 \cdot \mathtt{netuse} + \beta_3 \cdot \mathtt{treated} + \beta_4 \cdot \mathtt{green} + \beta_5 \cdot \mathtt{phc}$$

Define submodels \mathcal{M}_1 and \mathcal{M}_2 with linear predictors

$$\eta_1 = \beta_{0,1} + \beta_1 \cdot \text{age} + \beta_4 \cdot \text{green} + \beta_5 \cdot \text{phc},$$

and

$$\eta_2 = \beta_{0,2} + \beta_2 \cdot \mathtt{netuse} + \beta_3 \cdot \mathtt{treated} + \beta_5 \cdot \mathtt{phc}.$$

	Intercept	β_1	β_2	β_3	β_4	β_5
Submodel 1	√	√			√	√
Submodel 2	\checkmark		\checkmark	\checkmark		\checkmark

The link parameter is $\phi = \beta_5$ and model specific parameters are $\psi_1 = (\beta_{0,1}, \beta_1, \beta_4)$ and $\psi_2 = (\beta_{0,2}, \beta_2, \beta_3)$. Both submodels have the same observable random variables $Y_1 = Y_2 = Y$, the response variable pos.

Define q_k for k = 1, 2 by

$$q_k(x) = \frac{\exp(\eta_k)}{1 + \exp(\eta_k)}$$

Take a normal prior as in Section 1 for $\beta_{1:5}$, and similarly take a normal prior for the intercepts

$$\beta_{0,k} \sim \mathcal{N}(\mu_{0,k}, \sigma_{0,k}^2).$$

Then the submodels have consistent prior marginals in the link parameter and Markov combination can be applied.

The joint distribution corresponding to submodel \mathcal{M}_1 , as a function of the parameters, is proportional to the posterior, which itself is proportional to the prior times the likelihood

$$p_{1}(\phi, \psi_{1}, \mathbf{y}_{1}) \propto p_{1}(\phi, \psi_{1}|\mathbf{y}_{1})$$

$$= p_{1}(\beta_{0,1}, \beta_{1}, \beta_{4}, \beta_{5}|\mathbf{y})$$

$$\propto \exp\left(\frac{1}{2\sigma_{0,1}^{2}}(\beta_{0,1} - \mu_{0,1})^{2}\right) \prod_{j=1,4,5} \exp\left(\frac{1}{2\sigma_{j}^{2}}(\beta_{j} - \mu_{j})^{2}\right) \times \underbrace{\prod_{i=1}^{2035} q_{1}(x_{i})^{y_{i}}(1 - q_{1}(x_{i}))^{1-y_{i}}}_{\text{Likelihood}}.$$

Similarly for \mathcal{M}_2

$$p_{2}(\phi, \psi_{2}, \mathbf{y}_{2}) \propto p_{2}(\phi, \psi_{2}|\mathbf{y}_{2})$$

$$= p_{2}(\beta_{0,2}, \beta_{2}, \beta_{3}, \beta_{5}|\mathbf{y})$$

$$\propto \exp\left(\frac{1}{2\sigma_{0,2}^{2}}(\beta_{0,2} - \mu_{0,2})^{2}\right) \prod_{j=2,3,5} \exp\left(\frac{1}{2\sigma_{j}^{2}}(\beta_{j} - \mu_{j})^{2}\right) \times \underbrace{\prod_{i=1}^{2035} q_{2}(x_{i})^{y_{i}} (1 - q_{2}(x_{i}))^{1-y_{i}}}_{\text{Likelihood}}.$$

Therefore the Markov combination in this case is

$$p_{\text{comb}}(\phi, \psi_{1}, \psi_{2}, \mathbf{y}_{1}, \mathbf{y}_{2}) = \frac{p_{1}(\phi, \psi_{1}, \mathbf{y}_{1}) p_{2}(\phi, \psi_{2}, \mathbf{y}_{2})}{p(\phi)}$$

$$\propto \frac{p_{1}(\beta_{0,1}, \beta_{1}, \beta_{4}, \beta_{5}|\mathbf{y}) p_{2}(\beta_{0,2}, \beta_{2}, \beta_{3}, \beta_{5}|\mathbf{y})}{p(\beta_{5})}$$

$$\propto \prod_{k=1}^{2} \exp\left(\frac{1}{2\sigma_{0,k}^{2}} (\beta_{0,k} - \mu_{0,k})^{2}\right) \prod_{j=1}^{5} \exp\left(\frac{1}{2\sigma_{j}^{2}} (\beta_{j} - \mu_{j})^{2}\right)$$

$$\times \prod_{i=1}^{2035} q_{1}(x_{i})^{y_{i}} (1 - q_{1}(x_{i}))^{1-y_{i}} q_{2}(x_{i})^{y_{i}} (1 - q_{2}(x_{i}))^{1-y_{i}}$$
(*)

Informally p_{comb} contains the product of the priors on the full set of parameters (ϕ, ψ_1, ψ_2) with both likelihoods from \mathcal{M}_1 and \mathcal{M}_2 . So this is almost like Bayesian inference for full model but with a different likelihood. It seems as if the information contained by the data \mathcal{D} is being used more than once.

4.3. Monte Carlo schemes

Want to sample from the target (*) using the methods described in (Goudie 2018). Continue to use (symmetric) normal proposals throughout.

4.3.1 Metropolis-within-Gibbs

- Update first latent parameter ψ_1 by systematic scan Metropolis-within-Gibbs
- Update second latent parameter ψ_2 by systematic scan Metropolis-within-Gibbs
- Update link parameter ϕ by Metropolis-within-Gibbs

4.3.2 Multi-stage Metropolis-within-Gibbs

To-do

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