

Figure 11.11 Posterior summaries for mortality rates for cardiae surgery data. Posterior means and 0.95 equitailed credible intervals for separate analyses for each hospitaire shown by hollow circles and dotted lines, while blobs and solid lines show the corresponding quantities for a hierarchical model. Nose the shrinkage of the estimates for the hierarchical model towards the overall posterior mean rate, shown as the solid vertical intervals are slightly shorter than those for the simpler model.

Example 11.27 (Spring barley data) Table 10.21 contains data on a field trial intended to compare the yields of 75 varieties of spring barley allocated randomly to plots in three long narrow blocks. The data were analysed in Example 10.35 using a generalized additive model to accommodate the strong fertility trends over the blocks. In the absence of detailed knowledge about the varieties it seems natural to treat them as exchangeable, and we outline a Bayesian hierarchical approach. We also show how the fertility patterns may be modelled using a simple Markov random field.

Let $y = (y_1, \dots, y_n)^T$ denote the yields in the n = 225 plots and let ψ_j denote the unknown fertility of plot j. Let X denote the $n \times p$ design matrix that shows which of the p = 75 variety parameters $\beta = (\beta_1, \dots, \beta_p)^T$ have been allocated to the plots. Then a normal linear model for the yields is

$$y \mid \beta, \psi, \lambda_y \sim N_n(\psi + X\beta, I_n/\lambda_y),$$
 (11.51)

where ψ is the $n \times 1$ vector containing the fertilities and λ_y is the unknown precision of the ys.

We take the prior density of λ_y to be gamma with shape and scale parameters a and b, G(a,b), so that its prior mean and variance are a/b and a/b^2 , where a and b are specified. As there is no special treatment structure, we take for the β_r the exchangeable prior $\beta \sim N_p(0, I_p/\lambda_\beta^{-1})$, with $\lambda_\beta \sim G(c,d)$ and c, d specified. For the fertilities we take the normal Markov chain of Example 6.13, for which

$$\pi(\psi \mid \lambda_{\psi}) \propto \lambda_{\psi}^{n/2} \exp\left\{-\frac{1}{2}\lambda_{\psi} \sum_{i \sim j} (\psi_i - \psi_j)^2\right\}, \quad \lambda_{\psi} > 0, \quad (11.52)$$

the summation being over pairs of neighbouring plots and λ_{ψ}^{-1} being the variance of differences between fertilities. Each ψ_{j} occurs in n_{j} terms, where $n_{j} = 1$ or 2 is the

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Figure 11.11 Posterio summaries for mortalin rates for cardiac surgar data. Posterior mo 0.95 equitailed credition intervals for separat analyses for each hou are shown by bollow circles and dotted lin while blobs and solid! show the corresponding quantities for a hierarchical model. 8 the shrinkage of the estimates for the hierarchical model towards the overall posterior mean can shown as the solid w line; the hierarchical intervals are slightly shorter than those for the simpler model.

number of plots adjacent to plot j. The sum in (11.52) equals $\psi^{T}W\psi$, where W is the $n \times n$ tridiagonal matrix with elements

$$w_{ij} = \begin{cases} n_i, & i = j, \\ -1, & i \sim j, \\ 0, & \text{otherwise} \end{cases}$$

Thus W is block diagonal, with three blocks like the matrix V in Example 6.13 with $\tau = 0$, corresponding to the three physical blocks of the experiment. We take $\lambda_{\psi} \sim G(g,h)$, with g and h specified.

With these conjugate prior densities, the joint posterior density is

$$\pi(\beta, \psi, \lambda) \propto \lambda_y^{n/2} \exp\left\{-\frac{1}{2}\lambda_y (y - \psi - X\beta)^{\mathrm{T}} (y - \psi - X\beta)\right\}$$
$$\times \lambda_\beta^{p/2} \exp\left(-\frac{1}{2}\lambda_\beta \beta^{\mathrm{T}} \beta\right) \times \lambda_\psi^{p/2} \exp\left(-\frac{1}{2}\lambda_\psi \psi^{\mathrm{T}} W \psi\right)$$
$$\times \lambda_y^{n-1} \exp(-b\lambda_y) \times \lambda_\beta^{n-1} \exp(-c\lambda_\beta) \times \lambda_\psi^{n-1} \exp(-h\lambda_\psi),$$

where $\lambda=(\lambda_y,\lambda_{jl},\lambda_{\psi})^{\gamma}.$ The full conditional densities turn out to be

$$\beta \mid \psi, \lambda, y \sim N\{\lambda_y Q_{\beta}^{-1} X^{\tau}(y - \psi), Q_{\beta}^{-1}\},$$
 (11.53)

$$\psi \mid \beta, \lambda, y \sim N\{\lambda, Q_{\psi}^{-1}(y - X\beta), Q_{\psi}^{-1}\},$$
 (11.54)

$$\lambda_y \mid \psi, \beta, y \sim G(a + n/2, b + (y - X\beta - \psi)^{\mathsf{T}}(y - X\beta - \psi)/2).$$
 (11.55)

$$\lambda_{\beta} \mid \psi, \beta, y \sim G(c + p/2, d + \beta^{T}\beta/2),$$
 (11.56)

$$\lambda_{\psi} \mid \psi, \beta, y \sim G(g + n/2, h + \psi^{T} W \psi/2),$$
 (11.57)

where

$$Q_{\beta} = \lambda_{y} X^{\dagger} X + \lambda_{\beta} I_{\rho}, \quad Q_{\psi} = \lambda_{y} I_{n} + \lambda_{\psi} W.$$

The elements of λ are independent conditional on the remaining variables. The relatively simple form of the densities in (11.53)–(11.57) suggests using a time-reversible Gibbs sampler, in which β , ψ , and λ are updated in a random order at each iteration. The most direct approach to simulation in (11.53) and (11.54) is through Cholesky decomposition of Q_{β} and Q_{ψ} : in (11.53), for example, we find the lower triangular matrix L such that $LL^{\tau} = Q_{\beta}^{-1}$, generate $\varepsilon \sim N_{\rho}(0, I_{\rho})$, and let $\beta = \lambda_{\gamma} Q_{\beta}^{-1} X^{\tau}(y - \psi) + L\varepsilon$. The block diagonal structure of W means that the ψ s for different blocks can be updated separately, so the largest Cholesky decomposition needed is that of a 75 × 75 matrix. An alternative is to update individual ψ_{β} s in a random order, but although the computational burden is smaller, the algorithm then converges more slowly than with direct use of (11.54).

Note the strong resemblance of (11.53) and (11.54) to the steps of the backfitting algorithm for the corresponding generalized additive model.

The missing response in block 3 is simply a further unknown whose value may be simulated using the relevant marginal density of (11.51). This adds a fourth component to the simulation in random order of β , ψ , and λ at each iteration; there are no other changes to the algorithm.

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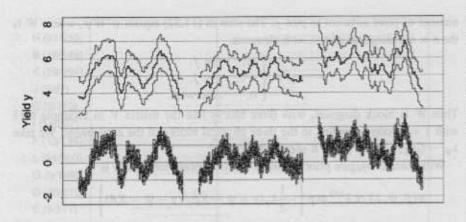
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gure 11.12 Possessor immaries for fertility and ψ for the three ocks of spring barley ta, shown from left to tht. Above: median and (heavy) and overall ψ posterior credible ands. Below: 20 mulated trends from bbs sampler output.

Table 11.12 Posterior probabilities that a variety is ranked among the best r varieties, estimated from 10,000 iterations of Gibbs sampler.

Location

If the matrix X^TX is diagonal, then the full conditional density for the rth variety effect has form

$$\beta_r \mid \psi, \lambda, y \sim N\left(\frac{\lambda_y m_r \overline{\xi}_r}{\lambda_{\beta} + \lambda_y m_r}, \frac{1}{\lambda_{\beta} + \lambda_y m_r}\right),$$

where \bar{z}_r is the current average of $y_j - \psi_j$ for the m_r plots receiving variety r. Thus the β_r are shrunk towards zero by an amount that depends on the ratio $\lambda_\beta/\lambda_\gamma$; with $\lambda_\beta = 0$ the mean for β in (11.53) is the least squares estimate computed by regressing $y - \psi$ on the columns of X. Unlike in Example 11.25, however, the normal distributions of the β_r are here averaged over the posterior densities of ψ , λ_γ and λ_β .

The algorithm described above was run with random initial values for 10,500 iterations. Time series plots of the parameters and log likelihood suggested that it had converged after 500 iterations, and inferences below are based on the final 10,000 iterations. The variance inflation factors $\hat{\tau}$ were less than 4 for ψ and β , about 44, 6 and 30 for λ_y , λ_r and λ_{ψ} , and about 6 for y_{187} . Thus estimation for λ_y is least reliable, being based on a sample equivalent to about 220 independent observations. A longer run of the algorithm would seem wise in practice. Based on this run, the posterior 0.9 credible intervals for λ_y , λ_{ψ} and λ_{β} were (5.2, 12.4), (5.0, 11.5) and (2.7, 5.7) respectively, and differences of two variety effects have posterior densities very close to normal with typical standard deviation of 0.35. The corresponding standard error for the generalized additive model was 0.41, so use of a hierarchical model and injection of prior information has increased the precision of these comparisons.

Figure 11.12 shows some simulated values of ψ and pointwise 0.90 credible envelopes for the true ψ . These envelopes are constructed by joining the 0.05 quantiles of the fertilities simulated from the posterior density, for each location, and likewise with the 0.95 quantiles. By contrast with the analysis in Example 10.35, the effective degrees of freedom for ψ , controlled by λ_{ψ} , are here equal for each block, leading to apparent overfitting of the fertilities for block 2 compared to the generalized additive model, A difference between the models is that the current model corresponds

11.12 Posterior	Liui	dia.	Mirko	3.007	enle	Va	riety	1370	- (t)	W.P	(0)94
mics, estimated from the iterations of Gibb	P	56	35	72	31	55	47	54	18	38	40
	1 2 5 10	0.327 0.518 0.814 0.959	0.182 0.357 0.690 0.908	0.149 0.299 0.643 0.887	0.129 0.270 0.621 0.871	0.075 0.174 0.486 0.795	0.055 0.136 0.416 0.743	0.019 0.050 0.234 0.560	0.015 0.042 0.183 0.497	0.012 0.035 0.153 0.429	0.006 0.020 0.106 0.344

to first differences of ψ being a normal random sample, while in the earlier model the second differences are a normal random sample, giving a smoother fit.

The posterior probabilities that certain varieties rank among the r best are given in Table 11.12. The ordering is somewhat different from that in Example 10.35, perhaps due to the slightly different treatment of fertility effects. As mentioned previously, no single variety strongly outperforms the rest, and future field experiments would have to include several of those included in this trial. This type of information is difficult to obtain using frequentist procedures, but is readily found by manipulating the output of the simulation algorithm described above.

This analysis is relatively easily modified when elements of the model are changed. Indeed the priors and other components chosen largely for convenience should be varied in order to assess the sensitivity of the conclusions to them; see Exercise 11.3.6. Metropolis-Hastings steps would then typically replace the Gibbs updates in the algorithm.

As mentioned above, more complicated hierarchies involve several layers of nested variation. Such models are widely used in certain applications, but their assessment and comparison can be difficult. For instance, shrinkage makes it unclear just how many parameters a hierarchical model has. Hierarchical modelling is an active area of current research.

Justification of (11.49)

To establish (11.49), suppose that r lies in $0, \ldots, n$ and that m > n. Then exchangeability of U_1, \ldots, U_m implies that the conditional probability

$$Pr(U_1 + \cdots + U_n = r \mid U_1 + \cdots + U_m = s)$$

equals the probability of seeing r 1's in n draws without replacement from an urn containing s 1's and m-s 0's, which is $\binom{m}{n}^{-1}\binom{s}{r}\binom{m-s}{n-r}$ for $s=r,\ldots,m-(n-r)$ and zero otherwise. Hence

$$\Pr(U_1 + \dots + U_n = r) = \sum_{s=r}^{m-(n-r)} {m \choose r}^{-1} {s \choose r} {m-s \choose n-r} \Pr(U_1 + \dots + U_m = s)$$

$$= {n \choose r} \sum_{s=r}^{m-(n-r)} \frac{s^{(r)}(m-s)^{(n-r)}}{m^{(n)}} \Pr(U_1 + \dots + U_m = s),$$

	_	lock I	Н.	llock 2	Block 3		
Location t	Variety	Yield y	Variety	Yield y	Variety	Yield	
1	57	9.29	49	7.99	63	11.0	
2	39	8.16	18	9.56		11.7	
3	3	8.97	8	9.02	38	12.03	
4	48	8.33	69	8.91	14	12.2	
5	75	8.66	29	9.17	71	10.96	
6	21	9.05	59		22	9.94	
7	66	9.01	19	9.49	46	9.27	
8	12	9.40	39	9.73 9.38	6	11.05	
9	30	10.16	67		30	11.40	
10	32	10.30	57	8.80	16	10.78	
-11	59	10.73	37	9.72	24	10.30	
12	50	9.69	26	10.24	40	11.27	
13	5	11.49		10.85	64	11.13	
14	23	10.73	16	9,67	8	10.55	
15	14	10.71	6	10.17	56	12.82	
16	68		47	11.46	32	10.95	
17	41	10.21	36	10.05	48	10.92	
18	1	10.52	64	11.47	54	10.77	
19	64	11.09	6.3	10.63	37	11.08	
20	28	11.39	33	11.03	21	10.22	
21	46	11.24	74	10.85	29	10.59	
22	73	10.65	13	11.35	62	11.35	
23	37	10.77	43	10.25	5	11.39	
24	55	10.92	3	10.08	70	10.59	
25	19	12.07	53	10.25	13	11.26	
26	10	11.03	23	9.57	- 11	11.79	
27		11.64	62	11.34	44	12.25	
28	35	11.37	52	10.19	36	12.23	
29	26	10.34	12	10.80	52	10.84	
30	17	9.52	2	10.04	60	10.92	
31	71	8.99	32	9.69	68	10.41	
32	8	8.34	22	9.36	3	10.96	
	62	9.25	42	9.43	19	9.94	
33	44	9.86	72	11.46	67	11.27	
34	53	9.90	73	9.29	59	11.79	
35	74	11.04	25	10.10	2	11.51	
36	20	10.30	45	9.53	75	11.64	
37	56	11.56	15	10.55	27		
38	29	9.69	35	11.34	43	9.78	
39	2	10.68	66	11.36	51	8.86	
40	47	10.91	5	10.88	10	10.28	
41	11	10.05	56	11.61	35	12.15	
42	38	10.80	46	10.33	74	10.36	
43	65	10.06	71	10.53	66	9.59	
44	13	10.04	51	8.67	34	10.53	
45	31	10.50	21	9.56	18	11.26	
46	40	9.51	1	9.95	50	10.37	
47	4	9.20	31	11.10	42	10.10	
48	67	9.74	11	10.11	1	9.95	
49	22	8.84	41	9.36	58	9.80	
50	49	9.33	61	10.23	26	10.58	
51	58	9.51	55	11.38	41	9.31	
52	43	9.35	14	11.30	25		
					447	9.29	

Table 10.21 Spring barley data (Bessa of 1995). Spatial layou applot yield at harvest (standardized to be crude variance) in a sassessment trial of 75 varieties of speins harley. The varieties sown in three blocks each variety replicate thrice in the design yield for variety 27 missing in the third to

Table 10.21 Spring barley data (Besag et al. 1995). Spatial layout plot yield at harvest y (standardized to have crude variance) in a fassessment trial of 75 varieties of apring barley. The varieties asown in three blocks, each variety replication to the design. The yield for variety 27 is missing in the third is

Location t	Blo	ck 1	Blo	ck 2	Block 3		
	Variety	Yield y	Variety	Yield y	Variety	Yield y	
53	7	9.01	44	10.90	33	10.03	
54	25	10.58	34	10.97	9	9.49	
55	61	11.03	54	12.22	17	11.52	
56	16	9.89	24	10.10	57	12.24	
57	52	11.39	4	11.22	65	11.64	
58	70	11.24	65	10.01	49	10.74	
59	34	12.18	75	10.29	73	10.29	
60	42	10.21	38	10.95	7	10.25	
61	24	11.08	17	9.66	23	11.39	
62	33	11.05	68	9.31	72	13.34	
63	51	10.29	7	8.84	55	12.73	
64	60	10.57	27	10.64	31	12.62	
65	69	10.42	58	9.45	39	10.19	
66	1.5	10.49	48	9.66	47	11.61	
67	6	10.00	28	9.85	.15	10.52	
68	63	9.23	60	9.24	20	9.07	
69	54	10.57	30	10.11	61	10.76	
70	18	10.27	70	9.63	28	9.91	
71	45	8.86	20	9.04	53	10.17	
72	72	9.45	9	8.43	69	8.68	
73	9	8.03	40	10.97	45	8.74	
74	36	9.22	50	8.98	12	9.15	
75	27	8.70	10	9.88	4	9.39	

10.7.3 More general models

We now consider how the discussion above should be modified when there are explanatory variables as well as a smooth variable, treating certain covariates nonparametrically and others not, and allowing the response to have a density other than the normal.

Let the data consist of independent triples $(x_1, t_1, y_1), \ldots, (x_n, t_n, y_n)$, with jth log likelihood contribution $\ell_j(\eta_j, \kappa)$, where $\eta_j = x_j^{\mathrm{T}} \beta + g(t_j)$; for now we suppress dependence on κ . Then the analogue of (10.47) is the penalized log likelihood

$$\ell_{\lambda}(\beta, g) = \sum_{j=1}^{n} \ell_{j}(\eta_{j}) - \frac{1}{2}\lambda \int_{a}^{b} \{g''(t)\}^{2} dt, \quad \lambda > 0,$$
 (10.49)

where a and b are chosen so that $a < t_1, \ldots, t_n < b$. If all the t_j are distinct and $\lambda = 0$, the maximum is obtained by choosing $g_j = g(t_j)$ to maximise the jth log likelihood contribution, but this is not useful because the resulting model has n parameters and is too rough. The integral in (10.49) penalizes roughness of g(t), so λ has the same interpretation as before.

If the ordered distinct values of t_1, \ldots, t_n are $s_1 < \cdots < s_q$ and if g(t) is a natural cubic spline with knots at the s_i , then the integral in (10.49) may be written $g^T K g$, where the $q \times 1$ vector g has ith element $g_i = g(s_i)$. Given a value of λ , our aim