Implementing a BYM Model in Stan to Fit Boston Housing Price Data

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1 Data on Boston Housing Prices in 1978

For this study we'll be using a dataset originally published in 1978 on housing prices in Boston, Massachusetts (David and Rubinfeld (1978)). This data was made accessible through the spData package in R (Bivand (2022)). Specifically, it contains median house prices and potential covariates for 506 tracts in the Greater Boston Area. Fig. 1 shows the median price per tract of land. It should be understood that this data is censored and median values over \$50,000 are capped. A full list of covariates can be found in the documentation for the boston dataset in spData. What we wish to understand is which of these covariates are indeed related to housing prices after being added to a spatial bayesian model.

Starting with some exploratory data analysis we identified four covariates of particular interest. Crime rates per capita, average number of rooms per dwelling, weighted distance to employment centers, and the nitric oxide concentrations per town.

First, as the distribution of values is right skewed (and following in the footsteps of (Moraga (2023))) we took as our target variable the logarithm of the median house value per tract instead of the median value itself. Fig. 2 shows the distribution of our target.

Fig. 3 shows the relationship between per capita crime rate and our target. There appears to be a clear negative relationship between then two and a rather wide spread of values in and around 0.

Fig. 4 shows how our target varies with the average number of rooms per dwelling in each tract. Here we can see a strong positive relationship between the two which makes a great deal of sense.

In Fig. 5 we see what looks to be a similar positive relationship between the logarithm of the average weighted distance to employment centers and our target. However this relationship is not as clear or strong as the one between our target and the number of rooms per dwelling.

Finally Fig. 6 shows us the relationship between nitric oxide concentrations in parts per million and our target with a somewhat spurious negative relationship.

2 Defining the Model

We will be using a variant of the Besag-York-Mollié (BYM) model (Moraga (2023))(Mitzi Morris (2019)). In this model we assume that we have an observations of our target variable Y_i (log median housing value in our case) for each tract i. Furthermore we assume our Y_i can be modeled as a normal distribution:

$$Y_i \sim Normal(\mu_i, \sigma^2)$$

where it is the μ_i that will be a function by our covariates. Specifically:

$$\mu_i = \beta_0 + \vec{\beta}\vec{x_i} + \sigma_r \left(\sqrt{\rho}\phi_i + \sqrt{1-\rho}\theta_i\right)$$

where β_0 is our intercept, $\vec{\beta}$ are the coefficients for our models effects from each covariate, x_i are our covariates corresponding to tract i, ϕ_i and θ_i are spatial and random effects respectively, and $sigma_r$ and ρ allow us to control the effect of the random variables as well as the degree to which our model has spatial and/or unstructured noise (if $\rho = 1$ we have only spatial structure whereas if $\rho = 0$ it is totally unstructured).

For our more straightforward priors we will have:

$$\beta_0 \sim Normal(0, 1)$$

$$\beta_i \sim Normal(0, 1)$$

$$\theta_i \sim Normal(0, 1)$$

$$\sigma_r \sim Uniform(0, 1)$$

$$\rho = \frac{e^r}{1 + e^r}, r \sim Normal(0, 1)$$

In the initial experimentation with this model we also had the prior:

$$\sigma \sim Normal(0,1)$$

but consistently found $\sigma \approx 0$. Therefore going forward we simply assume $\sigma = 0.01$ in order to not over complicate the sampling.

Our spatial terms ϕ_i are a little more complicated.

Each spatial interaction term ϕ_i is modeled as conditional on the other terms:

$$\phi_i | \phi_j \sim N\left(\sum_j w_{ij}\phi_j, \sigma^2\right), i \neq j$$

This gives us a conditional autoregressive model (CAR). A key result that Besag proved (Julian (1974)) is that the joint distribution ϕ ends up being multivariate normal random variable centered at 0

$$\vec{\phi} \sim N(0, Q^{-1})$$

where $Q = D(I - \alpha A)$. D is a diagonal "neighbors" matrix (each element on the diagonal is the number of neighbors unit i has), A is an adjacency matrix where if i, j are neighbors then the i, j element is 1, and α lets us control spatial dependence. This results in a log probability density of $\vec{\phi}$ which is proportional to (Mitzi Morris (2019)):

$$\frac{n}{2}\log\left(\det Q\right) - \frac{1}{2}\vec{\phi}^T Q\vec{\phi}$$

Given $\det Q$ is a constant and MCMC samplers compute the log probability up to a proportionality constant (Mitzi Morris (2019)) the first term drops out of the computation thereby reducing the computational intensity of this evaluation.

In our case, as we will be following the stan implementation from the paper (Mitzi Morris (2019)) we will be setting $\alpha=1$ and thereby get an intrinsic conditional autoregressive model (ICAR) in which Q reduces to D-A. With an ICAR model each ϕ_i is distributed with a mean equal to the average of its neighbors. If we additionally assume that $\vec{\phi}$ is centered at zero with common variance 1, then the joint probability of $\vec{\phi}$ becomes (Mitzi Morris (2019)):

$$p(\vec{\phi}) \propto \exp\left(-\frac{1}{2}\sum_{i\sim j}(\phi_i - \phi_j)^2\right)$$

where $i \sim j$ indicates that i and j are neighbors. This then is our prior for the ϕ_i - an ICAR model centered at 0 with common variance 1.

3 Fitting the Model

After computing the appropriate features (covariates and neighbors arrays) we fit our model using rstan (Stan Development Team (2024)) in the R programming language (R Core Team (2024)). Each model was fit with 4 chains and 7,500 iterations per chain.

Two models were fit, one with the crime rates feature and one without (see section 4). The trace plots for the former can be found in figures 7-10 and the trace plots for the latter are in figures 15-18. Densities are in figures 11-14 and 19-22 respectively. The order of features was crime, rooms, distance, and nox for the full model and rooms, distance, and nox for the limited model.

See section 5 for the $\,$ stan $\,$ model itself and section 6 for the $\,$ R $\,$ code used to build inputs.

4 Results

Parameter	95% CI	Mean	SD	SE	\hat{R}	n_{EFF}
β_0	2.130, 2.397	2.264	0.198	0.004	1.001	2495
β_{CRIM}	-0.009, -0.007	-0.008	0.001	≈ 0	1.000	14346
eta_{RM}	0.237, 0.255	0.246	0.014	≈ 0	1.000	8457
$\beta_{\log(DIS)}$	-0.171, -0.054	-0.112	0.085	0.004	1.008	546
β_{NOX}	-1.234, -0.972	-1.105	0.197	0.003	1.002	3557
σ_r	0.341, 0.363	0.352	0.016	≈ 0	1.001	2889
ho	0.941, 0.966	0.952	0.019	≈ 0	1.001	1461

Given the very small value on β_{CRIM} we also trained a model without that feature present.

Parameter	95% CI	Mean	SD	SE	\hat{R}	n_{EFF}
eta_0	2.041, 2.312	2.177	0.203	0.004	1.001	2552
eta_{RM}	0.237, 0.257	0.247	0.015	≈ 0	1.000	7170
$\beta_{\log(DIS)}$	-0.119, 0.003	-0.061	0.085	0.004	1.009	395
β_{NOX}	-1.260, -0.983	-1.122	0.205	0.004	1.003	3205
σ_r	0.361, 0.381	0.371	0.015	≈ 0	1.002	2623
ho	0.952, 0.973	0.961	0.016	0.001	1.005	947

Here we can see that the standard errors are small and the \hat{R} values are all below 1.05. In addition the traceplots all look grassy (Figures 15-18).

Therefore it does not seem there is much evidence to suggest a lack of convergence.

The results themselves are quite interesting. We can see we have the positive relationship between number of rooms per dwelling and our median housing price with β_{RM} being positive. It is also interesting to note just how inconsequential β_{CRIM} was when we took into account the other features and the spatial correlations among tracts (through the ϕ_i). We can also see that contrary to our original view, there is in fact a negative relationship between the distance to centers of employment and median housing prices per tract ($\beta_{\log DIS}$). However the relationship is also quite small. Finally there is a strong negative relationship between nitric oxide levels and housing prices.

It is also quite interesting to note that ρ ended up being very nearly one meaning our noise was almost entirely spatially structured.

Given the relatively small value of $\beta_{\log DIS}$ and the fact that its 95% credible interval crosses 0, we also looked at the bayes factor for the hypothesis that $\beta_{\log DIS} < 0$. The posterior odds were found to be ≈ 3.18 and given the prior was a normal distribution centered at 0 our prior odds are simply 1 giving us a bayes factor of 3.18. Thus there is some weak evidence that the relationship between distance to center of employment and our median housing prices is non-zero. Strong evidence, however, is lacking.

All in all then, it seems like our two strongest determinants of housing price are the number of rooms per dwelling and the levels of pollution as indicated by nitric oxide levels. Beyond that we also found that there was significant spatial structure in the median housing prices. All of these conclusions seem reasonable given the context.

5 The Stan Model

```
functions {
                   real icar_normal_lpdf(vector phi, int N, int[] node1, int[] node2) {
                                     return -0.5 * dot_self(phi[node1] - phi[node2])
                                                       + normal_lpdf(sum(phi) | 0, 0.001 * N);
                  }
}
data {
                  int<lower=0> N; // number of tracts
                  \mathbf{int} \hspace{-0.05cm} < \hspace{-0.05cm} \mathtt{lower} \hspace{-0.05cm} = \hspace{-0.05cm} \mathtt{0S} \hspace{-0.05cm} \hspace{-0.05cm} \hspace{-0.05cm} \hspace{-0.05cm} \mathtt{N\_edges} \hspace{-0.05cm} \hspace{-0.05cm} ; \hspace{0.2cm} \hspace{-0.05cm} / \hspace{-0.05cm} \hspace{-0.05cm} \textit{number of unique edges} \hspace{-0.05cm} \hspace{-0.05cm}
                  int<lower=1, upper=N> node1[N_edges]; // start of edge
                  int<lower=1, upper=N> node2[N_edges]; // end of edge
                  \begin{array}{lll} \textbf{int} < \textbf{lower=1} > \; K; \; \; / / \; \; \textit{number of covariates} \\ \text{matrix} \left[ N, \; K \right] \; \; x; \; \; / / \; \; \textit{design matrix} \end{array}
                   real y[N]; // target
}
parameters {
                  real beta0;
                   vector [K] betas;
                   real logit_rho;
                   vector [N] phi;
                   vector [N] theta;
                  real<lower=0> sigma_r;
transformed parameters {
                   real < lower = 0, upper = 1> rho = inv_logit (logit_rho);
                  vector[N] convolved_re = sqrt(rho) * phi
                                                                                                                                                     + \operatorname{sqrt}(1 - \operatorname{rho}) * \operatorname{theta};
}
model {
                  y ~ normal(beta0 + x * betas + convolved_re * sigma_r, 0.01);
                  target += icar_normal_lpdf(phi | N, node1, node2);
                  \begin{array}{ccc} \text{beta0} & \tilde{} & \text{normal(0, 1);} \\ \text{betas} & \tilde{} & \text{normal(0, 1);} \end{array}
                  logit_rho ~ normal(0, 1);
                  theta \tilde{} normal(0, 1);
                  sigma_r ~ uniform(0, 1);
}
            Note the term normal_lpdf(sum(phi)|0,0.001*N) used to center \vec{\phi} at
```

6 Building Inputs

```
library (sf)
library (spData)
library (spdep)
map \leftarrow st_read(
    system.file("shapes/boston_tracts.shp", package = "spData"),
    quiet = TRUE
# build neighbors arrays
nb \leftarrow poly2nb (map)
N = length (map$MEDV)
node1 = c()
node2 = c()
for (i in 1:N) {
    for (j in nb[[i]]) {
        if (j > i) 
            node1 = c(node1, i)
            node2 = c(node2, j)
        }
    }
N_{-}edges = length(node1)
\# build target and covariates
y = map log_median_value
x = cbind(map$CRIM, map$RM, log(map$DIS), map$NOX)
K = dim(x)[2]
   Packages used - spData (Bivand (2022)), sf (Pebesma and Bivand
(2023a) Pebesma (2018)), spdep (Bivand and Wong (2018) Roger Bivand
(2022) Bivand et al. (2013) Pebesma and Bivand (2023b))
```

7 Bibliography

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8 Figures

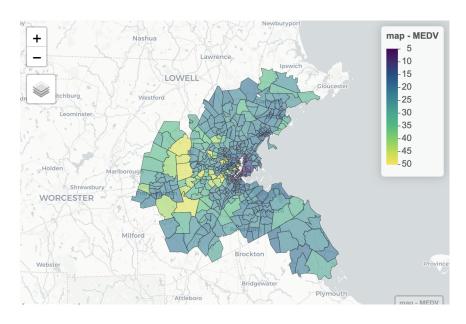


Figure 1: Median Housing Price

The median house value (in \$1000 USD) by census tract in the Greater Boston Area.

Frequency 50 100 150

2.5

0

2.0

Distribution of Log Median Values

Figure 2: Log Median Housing Price Distribution

Log Median Value

3.0

3.5

4.0

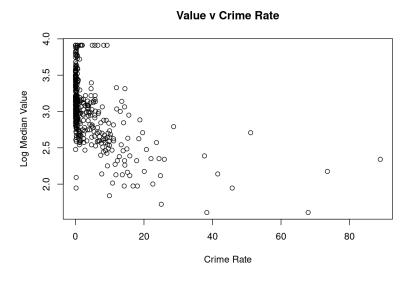


Figure 3: Log Median Price vs Crime Rate

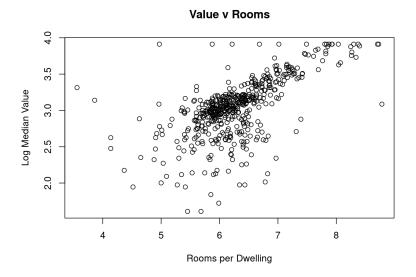


Figure 4: Log Median Price vs Rooms per Dwelling



Figure 5: Log Median Price vs Log Weighted Distance to Employment Centers

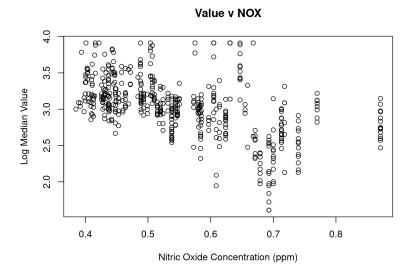


Figure 6: Log Median Price vs NOX

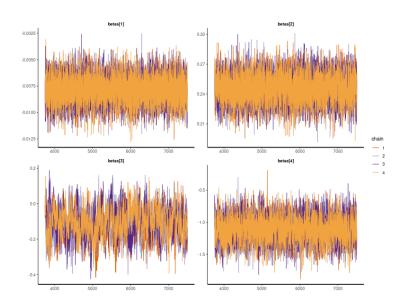


Figure 7: $\vec{\beta}$ Traceplots for 4 Feature Model

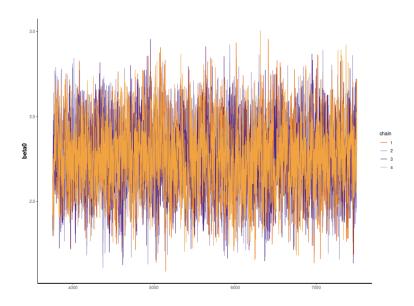


Figure 8: β_0 Traceplot for 4 Feature Model

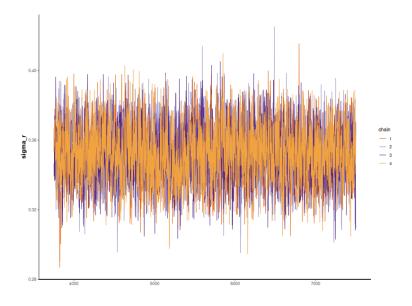


Figure 9: σ_r Traceplot for 4 Feature Model

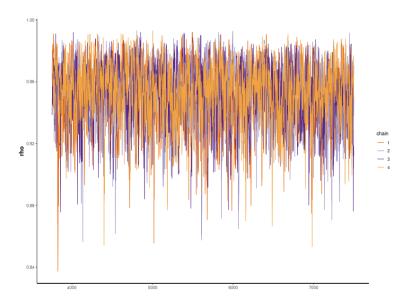


Figure 10: ρ Traceplot for 4 Feature Model

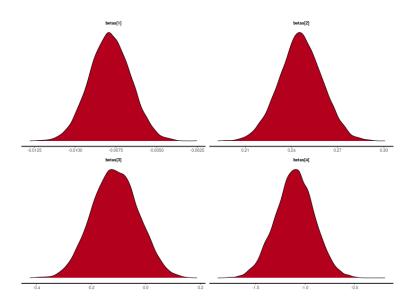


Figure 11: $\vec{\beta}$ Densities for 4 Feature Model

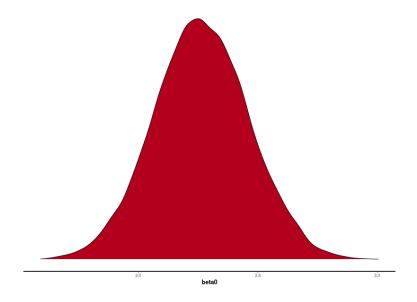


Figure 12: β_0 Density for 4 Feature Model

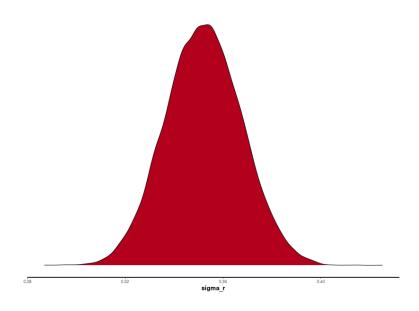


Figure 13: σ_r Density for 4 Feature Model

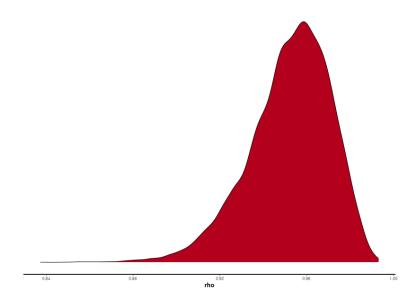


Figure 14: ρ Density for 4 Feature Model

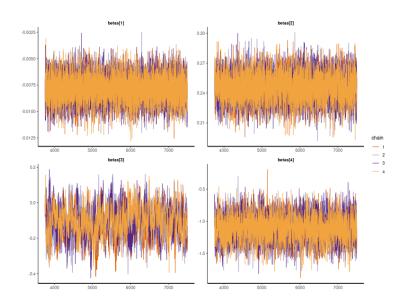


Figure 15: $\vec{\beta}$ Trace plots for 3 Feature Model

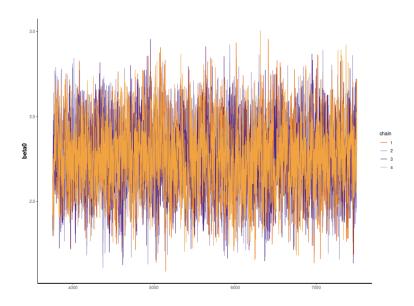


Figure 16: β_0 Traceplot for 3 Feature Model

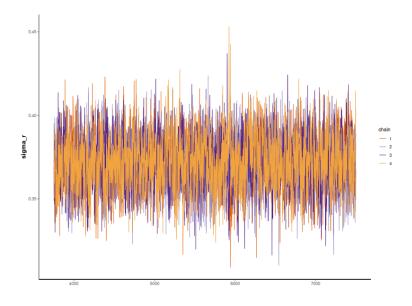


Figure 17: σ_r Traceplot for 3 Feature Model

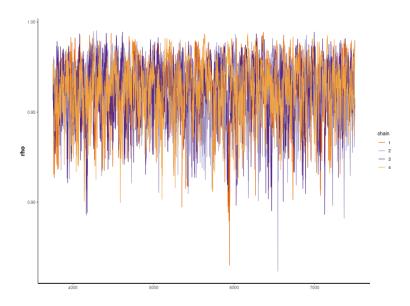


Figure 18: ρ Traceplot for 3 Feature Model

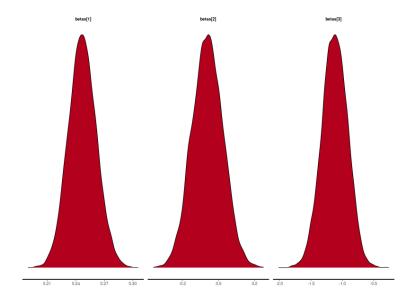


Figure 19: $\vec{\beta}$ Densities for 3 Feature Model

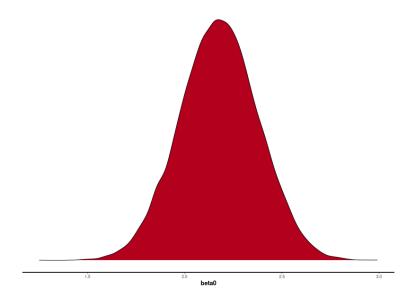


Figure 20: β_0 Density for 3 Feature Model

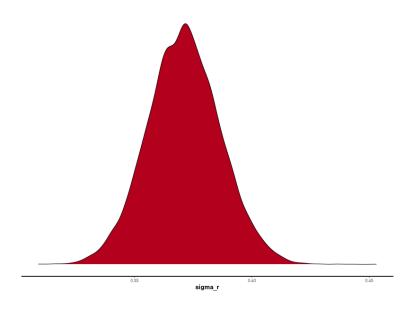


Figure 21: σ_r Density for 3 Feature Model

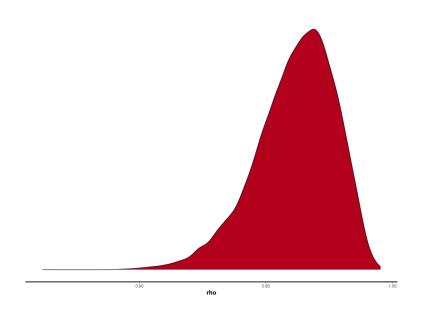


Figure 22: ρ Density for 3 Feature Model