

Basic CAR Model

Alvin Sheng

6/30/2021

```
library(here)

## here() starts at /Users/Alvin/Documents/NCSU_Fall_2021/NIH_SIP/flood-risk-health-effects
library(coda)
library(CARBayes)

## Loading required package: MASS
## Loading required package: Rcpp
## Registered S3 method overwritten by 'GGally':
##   method from
##   +.gg      ggplot2
```

CAR model results

Inference is based on 3 markov chains, each of which has been run for 100000 samples, the first 10000 of which has been removed for burn-in. The remaining 90000 samples are thinned by 5, resulting in $18000 \times 3 = 54000$ samples for inference across the 3 Markov chains.

```
load(here("modeling_files/model_3chains_var_exclude.RData"))
```

Output for the first chain is shown below.

```
chain1

##
## #####
## #### Model fitted
## #####
## Likelihood model - Gaussian (identity link function)
## Random effects model - Leroux CAR
## Regression equation - Y ~ X
## <environment: 0x7fd4b8b61700>
## Number of missing observations - 0
##
## #####
## #### Results
## #####
## Posterior quantities and DIC
##
##               Median    2.5%   97.5% n.effective Geweke.diag
## (Intercept)      77.7453 77.7253 77.7651      18000.0         1.9
```

```

## Xpct_fs_risk_2020_5      -0.1179 -0.2087 -0.0283      10734.0      1.1
## Xpct_floodfactor2        0.0171 -0.0283  0.0625      12960.6      0.8
## Xpct_floodfactor3       -0.0154 -0.0617  0.0308      12684.5     -0.5
## Xpct_floodfactor4        0.0479 -0.0024  0.0992       9935.2     -0.4
## Xpct_floodfactor5       -0.0196 -0.0827  0.0441      11001.7     -1.0
## Xpct_floodfactor6        0.0262 -0.0392  0.0918      12597.8      1.1
## Xpct_floodfactor7       -0.0120 -0.0623  0.0395      10856.9     -0.9
## Xpct_floodfactor8       -0.0144 -0.0583  0.0289      14894.6      0.1
## Xpct_floodfactor9        0.0877  0.0139  0.1609      11986.6     -1.1
## Xavg_risk_fsf_2020_100   0.1202  0.0388  0.2000       8785.6      0.2
## Xavg_risk_score_sfha     0.0083 -0.0430  0.0588      12590.9      1.9
## Xavg_risk_score_no_sfha -0.0063 -0.0954  0.0819      10668.5      0.3
## KEP_POV                  -0.1779 -0.2448 -0.1106      11811.2      0.4
## KEP_UNEMP                -0.0596 -0.1079 -0.0106      12352.9      1.1
## KEP_PCI                   0.2465  0.1733  0.3203      10025.0      0.5
## KEP_NOHSDP               -0.0541 -0.1368  0.0286       8373.5      0.0
## KEP_DISABL               -0.1211 -0.1789 -0.0630       9023.2     -1.2
## KEP_SNGPNT               -0.2182 -0.2656 -0.1727      14142.0      2.8
## KEP_MINRTY               -0.3378 -0.4233 -0.2514       5446.1     -0.7
## KEP_LIMENG                0.3492  0.2820  0.4150       8736.9     -0.3
## KEP_MUNIT                 0.1100  0.0494  0.1697       8593.8     -0.1
## KEP_MOBILE               -0.0883 -0.1493 -0.0271       7809.0      0.8
## KEP_CROWD                -0.0735 -0.1215 -0.0249       9235.4     -1.1
## KEP_NOVEH                -0.1260 -0.1793 -0.0722      10100.2     -0.1
## KEP_GROUPQ               0.1105  0.0758  0.1454      14218.3      0.7
## KEP_UNINSUR              -0.0314 -0.0895  0.0286       7662.7     -0.4
## Xco                       -0.0954 -0.1659 -0.0265       4192.4      1.0
## Xno2                     -0.0193 -0.1204  0.0829       3562.1     -0.7
## Xo3                      -0.0394 -0.1733  0.0929        943.6     -0.3
## Xpm10                     0.1123  0.0308  0.1929       3248.1     -0.5
## Xpm25                    -0.2881 -0.4125 -0.1668       1405.1      0.6
## Xso2                      -0.0758 -0.1279 -0.0248       6256.1      1.3
## Xtotal_mean              -0.9107 -0.9826 -0.8377       6328.3      0.3
## nu2                      0.3197  0.2576  0.3815       1883.7     -0.3
## tau2                     1.8848  1.5878  2.2258       1910.0      0.1
## rho                      0.9924  0.9753  0.9992       7881.1      0.8
##
## DIC = 6926.558      p.d = 1660.539      LMPL = -3825.94

```

The smallest effective sample size is 935.8, for ozone (o3).

```
chain1$accept
```

```

##      beta      phi      nu2      tau2      rho
## 100.0000 100.0000 100.0000 100.0000 45.2697

```

It appears that beta, phi, nu2, and tau2 probably have Gibbs steps, whereas rho has a Metropolis-Hastings step. In any case, the acceptance probabilities are acceptable.

Model Diagnostics

Beta samples

```

beta_samples <- mcmc.list(chain1$samples$beta, chain2$samples$beta,
                          chain3$samples$beta)

saveRDS(beta_samples, file = here("modeling_files/model_3chains_var_exclude_beta_samples.rds"))

plot(beta_samples)

gelman.diag(beta_samples)

## Potential scale reduction factors:
##
##      Point est. Upper C.I.
## [1,]          1          1
## [2,]          1          1
## [3,]          1          1
## [4,]          1          1
## [5,]          1          1
## [6,]          1          1
## [7,]          1          1
## [8,]          1          1
## [9,]          1          1
## [10,]         1          1
## [11,]         1          1
## [12,]         1          1
## [13,]         1          1
## [14,]         1          1
## [15,]         1          1
## [16,]         1          1
## [17,]         1          1
## [18,]         1          1
## [19,]         1          1
## [20,]         1          1
## [21,]         1          1
## [22,]         1          1
## [23,]         1          1
## [24,]         1          1
## [25,]         1          1
## [26,]         1          1
## [27,]         1          1
## [28,]         1          1
## [29,]         1          1
## [30,]         1          1
## [31,]         1          1
## [32,]         1          1
## [33,]         1          1
## [34,]         1          1
##
## Multivariate psrf
##
## 1

```

Examining tau2, nu2, rho

```
tau2_samples <- mcmc.list(chain1$samples$tau2, chain2$samples$tau2,  
                          chain3$samples$tau2)  
  
nu2_samples <- mcmc.list(chain1$samples$nu2, chain2$samples$nu2,  
                        chain3$samples$nu2)  
  
rho_samples <- mcmc.list(chain1$samples$rho, chain2$samples$rho,  
                        chain3$samples$rho)
```

```
plot(tau2_samples)
```

```
plot(nu2_samples)
```

```
plot(rho_samples)
```

```
gelman.diag(tau2_samples)
```

```
## Potential scale reduction factors:  
##  
##      Point est. Upper C.I.  
## [1,]          1          1
```

```
gelman.diag(nu2_samples)
```

```
## Potential scale reduction factors:  
##  
##      Point est. Upper C.I.  
## [1,]          1          1
```

```
gelman.diag(rho_samples)
```

```
## Potential scale reduction factors:  
##  
##      Point est. Upper C.I.  
## [1,]          1          1
```

Examining a sample of the 3108 phi parameters

```
phi_samples <- mcmc.list(chain1$samples$phi, chain2$samples$phi, chain3$samples$phi)  
  
set.seed(1157, kind = "Mersenne-Twister", normal.kind = "Inversion", sample.kind = "Rejection")  
  
phi_subset_idx <- sample(1:3108, size = 10)  
  
phi_samples_subset <- phi_samples[, phi_subset_idx]
```

```
plot(phi_samples_subset)
```

```
gelman.diag(phi_samples_subset)
```

```
## Potential scale reduction factors:  
##  
##      Point est. Upper C.I.
```

```
## [1,]      1      1
## [2,]      1      1
## [3,]      1      1
## [4,]      1      1
## [5,]      1      1
## [6,]      1      1
## [7,]      1      1
## [8,]      1      1
## [9,]      1      1
## [10,]     1      1
##
## Multivariate psrf
##
## 1
```

Inference

```
beta_samples_matrix <- rbind(chain1$samples$beta, chain2$samples$beta, chain3$samples$beta)
colnames(beta_samples_matrix) <- colnames(chain1$X)

(beta_inference <- round(t(apply(beta_samples_matrix, 2, quantile, c(0.5, 0.025, 0.975))),5))
```

```
##              50%      2.5%     97.5%
## (Intercept)    77.74541 77.72526 77.76547
## Xpct_fs_risk_2020_5 -0.11846 -0.20947 -0.02771
## Xpct_floodfactor2   0.01733 -0.02848  0.06292
## Xpct_floodfactor3  -0.01574 -0.06166  0.02993
## Xpct_floodfactor4   0.04814 -0.00219  0.09916
## Xpct_floodfactor5  -0.02001 -0.08280  0.04319
## Xpct_floodfactor6   0.02603 -0.03848  0.09156
## Xpct_floodfactor7  -0.01147 -0.06218  0.03928
## Xpct_floodfactor8  -0.01455 -0.05858  0.02901
## Xpct_floodfactor9   0.08748  0.01451  0.16079
## Xavg_risk_fsf_2020_100 0.12019  0.03839  0.20060
## Xavg_risk_score_sfha  0.00802 -0.04303  0.05908
## Xavg_risk_score_no_sfha -0.00614 -0.09451  0.08119
## XEP_POV          -0.17780 -0.24495 -0.11046
## XEP_UNEMP         -0.05955 -0.10768 -0.01091
## XEP_PCI           0.24681  0.17371  0.32026
## XEP_NOHSDP        -0.05413 -0.13624  0.02840
## XEP_DISABL        -0.12113 -0.17901 -0.06317
## XEP_SNGPNT        -0.21845 -0.26570 -0.17201
## XEP_MINRTY        -0.33722 -0.42218 -0.25146
## XEP_LIMENG         0.34887  0.28217  0.41493
## XEP_MUNIT         0.11002  0.04950  0.17033
## XEP_MOBILE        -0.08845 -0.14917 -0.02743
## XEP_CROWD         -0.07362 -0.12170 -0.02504
## XEP_NOVEH         -0.12598 -0.17966 -0.07259
## XEP_GROUPQ        0.11039  0.07611  0.14525
## XEP_UNINSUR       -0.03139 -0.09039  0.02834
## Xco              -0.09513 -0.16489 -0.02588
## Xno2             -0.01956 -0.12031  0.08103
```

```
## Xo3                -0.04186 -0.17364  0.09052
## Xpm10              0.11263  0.03151  0.19444
## Xpm25             -0.28796 -0.41248 -0.16528
## Xso2              -0.07616 -0.12793 -0.02453
## Xtotal_mean       -0.91055 -0.98245 -0.83768
```

List of significant beta coefficients:

```
colnames(beta_samples_matrix)[sign(beta_inference[, 2]) == sign(beta_inference[, 3])]
```

```
## [1] "(Intercept)"      "Xpct_fs_risk_2020_5"  "Xpct_floodfactor9"
## [4] "Xavg_risk_fsf_2020_100" "XEP_POV"             "XEP_UNEMP"
## [7] "XEP_PCI"           "XEP_DISABL"          "XEP_SNGPNT"
## [10] "XEP_MINRTY"        "XEP_LIMENG"          "XEP_MUNIT"
## [13] "XEP_MOBILE"        "XEP_CROWD"           "XEP_NOVEH"
## [16] "XEP_GROUPQ"        "Xco"                 "Xpm10"
## [19] "Xpm25"             "Xso2"                "Xtotal_mean"
```

My sparse implementation

```
load(here("modeling_files/model_1chain_var_exclude_sparse.RData"))
```

```
chain1$modelfit
```

```
##          DIC          p.d          WAIC          p.w          LMPL
##    6913.325    1649.640    6876.002    1227.458    -3806.863
## loglikelihood
##    -1807.022
```

```
mcmc_samps <- chain1$samples
```

```
effectiveSize(mcmc_samps$beta)
```

```
##      var1      var2      var3      var4      var5      var6      var7
## 17824.8740 10588.2173 13177.1579 13446.7282 10459.9458 10412.9187 11606.2789
##      var8      var9      var10      var11      var12      var13      var14
## 11633.0534 14869.8877 11052.7277  9542.8497 11781.1272 10761.8732 12304.7678
##      var15      var16      var17      var18      var19      var20      var21
## 12082.5510  9529.2160  9427.1320  9654.0071 13385.0412  4968.8319 10662.4156
##      var22      var23      var24      var25      var26      var27      var28
##  9006.2581  8618.2333 10854.5943 10996.9025 14468.5769  8008.8227  4197.2541
##      var29      var30      var31      var32      var33      var34
##  3307.6520   873.2518  3393.8121  1465.3655  6072.1736  6791.8248
```

It's easier to achieve a high sample size. I can have 10x fewer iterations.

```
effectiveSize(mcmc_samps$sigma2)
```

```
##      var1
## 1982.105
```

```
effectiveSize(mcmc_samps$nu2)
```

```
##      var1
## 2003.7
```

```
effectiveSize(mcmc_samps$rho)
```

```
##      var1  
## 8737.944
```

```
effectiveSize(mcmc_samps$Y)
```

```
##      var1      var2      var3      var4      var5      var6      var7      var8  
## 14958.76 16219.32 16425.60 15512.36 15399.18 15819.92 15661.60 16051.90  
##      var9      var10  
## 16502.47 16298.51
```

```
t(apply(mcmc_samps$beta, 2, quantile, c(0.5, 0.025, 0.975)))
```

```
##           50%           2.5%           97.5%  
## var1  77.746506148 77.726411457 77.76645330  
## var2  -0.119540299 -0.210452554 -0.02909804  
## var3   0.017900183 -0.027865030  0.06325522  
## var4  -0.016757355 -0.062849320  0.02888706  
## var5   0.050129231 -0.001709434  0.10149961  
## var6  -0.020836773 -0.084061452  0.04178001  
## var7   0.027238928 -0.037526315  0.09284415  
## var8  -0.010973621 -0.060695936  0.04078919  
## var9  -0.015118646 -0.059089625  0.02768246  
## var10  0.086282980  0.013777734  0.16009979  
## var11  0.121360779  0.042087761  0.20293893  
## var12  0.008158981 -0.043258548  0.05890849  
## var13 -0.006309570 -0.095211432  0.08202875  
## var14 -0.178166872 -0.244198785 -0.11073784  
## var15 -0.061460637 -0.110337992 -0.01330997  
## var16  0.244703557  0.170855815  0.31917766  
## var17 -0.053992619 -0.137481557  0.02888784  
## var18 -0.118966450 -0.177010691 -0.06076987  
## var19 -0.217929506 -0.264566749 -0.17075672  
## var20 -0.337053185 -0.422677017 -0.25329809  
## var21  0.347558929  0.282064515  0.41340951  
## var22  0.111408456  0.050665553  0.17062336  
## var23 -0.087001928 -0.146387155 -0.02668316  
## var24 -0.074577134 -0.123766678 -0.02567313  
## var25 -0.127042726 -0.180567305 -0.07353508  
## var26  0.109783964  0.074951764  0.14426383  
## var27 -0.028954755 -0.088080431  0.03103675  
## var28 -0.093294506 -0.163674562 -0.02383003  
## var29 -0.021817409 -0.119213178  0.07528710  
## var30 -0.050301272 -0.186201008  0.08593165  
## var31  0.115483751  0.034014079  0.19704008  
## var32 -0.287547411 -0.407863043 -0.16612464  
## var33 -0.078473225 -0.130344934 -0.02708925  
## var34 -0.913616225 -0.986850856 -0.84071714
```

```
quantile(mcmc_samps$nu2, c(0.5, 0.025, 0.975))
```

```
##           50%           2.5%           97.5%  
## 0.3216937 0.2609731 0.3824641
```

```
quantile(mcmc_samps$sigma2, c(0.5, 0.025, 0.975))
```

```
##      50%      2.5%      97.5%  
## 1.875368 1.576425 2.213394
```

```
quantile(mcmc_samps$rho, c(0.5, 0.025, 0.975))
```

```
##      50%      2.5%      97.5%  
## 0.9927457 0.9762306 0.9992126
```

Imputed Y values

```
t(apply(mcmc_samps$Y, 2, quantile, c(0.5, 0.025, 0.975)))
```

```
##      50%      2.5%      97.5%  
## var1  79.53963 77.91262 81.15431  
## var2  77.62478 75.98932 79.26445  
## var3  75.73865 74.12137 77.34398  
## var4  80.92168 79.21596 82.62262  
## var5  82.47714 80.79385 84.17298  
## var6  76.85689 75.17219 78.55423  
## var7  75.78624 74.16714 77.41043  
## var8  81.49117 79.95209 83.05497  
## var9  77.56431 75.99440 79.11463  
## var10 76.68195 75.07241 78.32694
```