## Bayesian Overview

* Bayes’ Theorem:
* **Evidence –** P(B) likelihood, averaged over the prior.
* **Prior –** P(A) probability distribution of the parameter before the evidence is taken into account.
  + Is updated based on the data.
* **Likelihood –** P(B|A) relative number of ways that the data can occur within the bounds of the prior.
* **Posterior distribution –** P(A|B) probability distribution of the parameter, given the prior and the data.
* **Parameter –** unknown. Estimated from the data.

## Priors

* **Informative –** specific, definite information about a variable.
* **Regularizing –** keeps inferences within a reasonable range, but he data does most of the informing.
  + Use this kind for myctophids.
  + We have some idea about what the parameters might be from previous studies, so can keep within a range, but we want the data do most of the informing.
* **Diffuse prior –** equal probabilities to all outcomes.

## Multilevel Models

1. Adjust for repeated sampling.
   * Many individuals per species.
2. Adjust for imbalance in sampling.
   * Different number of individuals per species.
3. Avoids averaging.
   * Individuals per species rather than species averages.

## MCMC

* **Monte Carlo simulations** - approximates a parameter by repeatedly generating random numbers.
* **Markov Chains** – sequence of event, where the next event is predicted by the current event.
* Basic method:

1. Pick a random potential parameter value.
2. Test how likely the parameter value is to explain the data, given the prior.
3. The parameter then informs the next parameter to be picked.

* Samples directly from posterior without assuming a shape.
  + I.E. doesn’t assume normality.
* **Gibbs Sampling –** adjusts the distribution of a proposed parameter value, depending on the current parameter values.
  + Used in BUGS and JAGS (and therefore MixSIAR).
  + Uses **conjugate priors** (prior and posterior distributions are from the same family, e.g. both Gaussian).
* **Hamiltonian Monte Carlo –** sends a frictionless particle to glide over the posterior, turning when the gradient of the posterior specifies.
  + Used in Stan.
  + Doesn’t require conjugate priors.
  + More efficient than Gibbs sampling.

## MixSIAR

* Uses JAGS (Just Another Gibbs Sampler).
* Mixing models that estimate proportion of sources to an isotopic mixture.
* Advantages of Bayesian:
  + Takes uncertainty into account for source values and discrimination.
  + Incorporates prior information.

### Priors

* Use the uninformative priors.

### MCMC Parameters

* **“Normal”**
  + Chain length = 100,000
  + Burn in = 50,000
    - Proportion of the chain that may not converge on the target distribution yet (discarded).
  + Thinning parameter = 50
  + Number of chains = 3

### Error Structure

* Error structure is “**process only**” because we only have one mixture data point.
  + Error comes from consumers sampling in different areas of the diet and DIC distribution.

### Diagnostics

* **Gelman-Rubin diagnostic** is the same as R-hat.
  + Needs to be a close to 1 to ensure convergence.
* **Geweke diagnostic** is a z-score.
  + MixSIAR tells you how many variables were outside +/- 1.96 in each chain.
* From PDF:
  + Deviance – Deviance information criterion.
    - Can ignore for our purposes.
  + ILR – Isometric log ratio transformation.
    - Used in the analysis of compositional data.
  + Loglik – log likelihood.
    - P(B|A)
  + p.fac and p.global – probability densities of each source.
    - Are the same for M because we are only looking at one individual at a time.
    - [1,1] is DIC and [1,2] is diet.
    - P(A|B)

### Outputs

* Posterior plot of contributions of each source.

## Body Mass Temperature Model

* M ~ a + a\_Var[Species] + b\_W\*Weight + b\_T\*Temp\_est
* Priors:
  + A, b\_W & b\_T ~ dnorm(0, 1)
  + Sigma & sigma\_Species ~ dexp(1)
* Adapt delta 🡪 size of steps that stan takes when sampling.
  + 0.999 to avoid divergent iterations.

### Test Run

* 3 chains, 1000 iter, 500 warmup.
* Errors:
  + 4 transitions after warmup that exceeded maximum treedepth.
  + 2 chains where the estimated Bayesian Fraction of Missing information was low.
  + ---
  + 809 transitions after warmup that exceeded maximum treedepth.
  + 2 chains where estimated Bayesian Fraction of missing information was low.
* Treedepth.
  + “When the maximum allowed tree depth is reached it indicates that NUTS is terminating prematurely to avoid excessively long execution time.”
  + NUTS 🡪 No U Turn Sampler (a variant of HMC used by Stan).
  + Try increasing max\_treedepth to 15.
    - Test 1 – worked!
    - Test 2 – worked!
* Bayesian Fraction of Missing information.
  + “This implies that the adaptation phase of the Markov Chains did not turn out well and those chains likely did not explore the posterior distribution efficiently.”
  + Try with longer chain (2000, 1000 iter).
    - Test 1 – 1 chain where BFMI was low.
    - Test 2 – 3 chains where BFMI was low.
  + E-BFMI should be above 0.2.
    - Chain 2 = 0.136.
    - Chain 3 = 0.056.
  + Try changing sigma to Cauchy prior.
    - Seems to have helped?
      * BFMI good.
      * Treedepth and iterations good.
    - Still some “banana” shape between b\_T and sigma.
* Tried long model.
  + 4000 + divergent iterations.
  + BFMI = 0.82 🡪 low.
* Try with longer warmup (5e5) and adapt delta = 0.9999.
  + Still getting errors, even worse than above.
* Try with JAGS number of iterations.
  + 100,000 iter 50,000 warmup.
  + N\_eff good.
  + BFMI low.
  + 60 divergent iterations.
  + 19 iterations exceeded max treedepth.
* Reparameterise model.
  + <https://mc-stan.org/docs/2_18/stan-users-guide/reparameterization-section.html>
  + Test with 3 chains. Iter = 1000, warmup = 500.

|  |  |  |
| --- | --- | --- |
| **Feature** | **Regular** | **Reparameterized – with sigma** |
| Error messages | None | “Stan model 'M\_est ~ dnorm(mu, sigma)' does not contain samples.” |
| N\_eff | Small for b\_T (71) and sigma(31). | Small for b\_T (52) and sigma (40). |
| R\_hat | High for sigma (1.08). | All below 1.01. |
| BFMI | 2 chains where BFMI was between 0.2 and 0.15. | Ok |
| Treedepth | OK | Ok |
| Divergent iterations | 0 | 0 |
| Pairs plot | All parameters fairly unimodal.  Correlation between b\_T and sigma. | All parameters fairly unimodal.  Correlation between b\_T and sigma. |
| Parameters | * A = 0.1835 * B\_W = 0.0016 * B\_T = -0.0034 * Sigma\_species = 0.338 * Sigma = 0.109 | * A = 0.1819 * B\_W = 0.0017 * B\_T = -0.0032 * Sigma\_species = 0.0341 * Sigma = 0.0114 |

* Try with default iter and warmup – regular.
  + Should have twice the number of effective samples.
    - Still low n\_eff for sigma (22) and b\_T (17).
    - Correspondingly high Rhat (1.16 and 1.13).
    - Correlation of 0.78.
* Email Andrew Jackson.
  + Give outline of the problem first rather than full on.

### Andrew Suggestions

* Try Z-scoring body mass and temperature.
  + It worked!

# Among Species

## Results – Body Mass and Temperature Model

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90
* Diagnostics:
  + Pairs plot, looking happy.
  + E-BFMI good.
  + Treedepth all ok.
  + No divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1825 | 0.0161 | 0.1528 | 0.2168 | 1213 | 0.9998 |
| b\_W | 0.0019 | 0.0045 | -0.0069 | 0.0109 | 5000 | 0.9998 |
| b\_T | -0.0036 | 0.0028 | -0.0089 | 0.0019 | 5000 | 0.9999 |
| a\_Var\_ELN | 0.0299 | 0.0165 | -0.0044 | 0.0615 | 1225 | 0.9999 |
| a\_Var\_ELC | -0.0097 | 0.0167 | -0.0420 | 0.0236 | 1263 | 0.9998 |
| a\_Var\_GYR | 0.0185 | 0.0169 | -0.0133 | 0.0526 | 1291 | 0.9998 |
| a\_Var\_GYN | -0.0381 | 0.0186 | -0.0780 | -0.0048 | 1472 | 0.9998 |
| a\_Var\_KRA | 0.0108 | 0.0174 | -0.0223 | 0.0455 | 1377 | 0.9999 |
| a\_Var\_PRM | -0.0113 | 0.0171 | -0.0440 | 0.0236 | 1418 | 0.9998 |
| sigma\_Species | 0.0352 | 0.0174 | 0.0124 | 0.0691 | 1901 | 0.9998 |
| sigma | 0.0170 | 0.0012 | 0.0147 | 0.0193 | 5000 | 0.9998 |

## Results – Temperature Model

* 3000 iter, 1500 warmup.
* Diagnostics:
  + Pairs plot, looking happy.
  + E-BFMI good.
  + Treedepth all ok.
  + No divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1819 | 0.1746 | 0.1547 | 0.2090 | 257 | 1.0060 |
| b\_T | -0.0041 | 0.0023 | -0.0083 | 0.0008 | 181 | 0.9993 |
| a\_Var\_ELN | 0.0296 | 0.0138 | -0.0001 | 0.0579 | 277 | 1.0045 |
| a\_Var\_ELC | -0.0084 | 0.0139 | -0.0363 | 0.0201 | 265 | 1.0048 |
| a\_Var\_GYR | 0.0191 | 0.0139 | -0.0073 | 0.0487 | 286 | 1.0048 |
| a\_Var\_GYN | -0.0335 | 0.0141 | -0.0604 | -0.0026 | 271 | 1.0029 |
| a\_Var\_KRA | 0.0097 | 0.0136 | -0.0161 | 0.0411 | 267 | 1.0059 |
| a\_Var\_PRM | -0.0135 | 0.0137 | -0.0413 | 0.0135 | 271 | 1.0033 |
| sigma\_Species | 0.0319 | 0.0140 | 0.0131 | 0.0608 | 585 | 1.0019 |
| sigma | 0.0124 | 0.0017 | 0.0089 | 0.0158 | 436 | 1.0004 |

## Results – Weight Model

* 3000 iter, 1500 warmup.
* Diagnostics:
  + Pairs plot, looking happy.
  + E-BFMI good.
  + Treedepth all ok.
  + No divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1808 | 0.0163 | 0.1474 | 0.2128 | 356 | 0.9993 |
| b\_W | 0.0036 | 0.0043 | -0.0045 | 0.0122 | 1500 | 1.0002 |
| a\_Var\_ELN | 0.0324 | 0.0167 | -0.0020 | 0.0644 | 361 | 0.9993 |
| a\_Var\_ELC | -0.0125 | 0.0167 | -0.0481 | 0.0196 | 364 | 0.9993 |
| a\_Var\_GYR | 0.0232 | 0.0168 | -0.0136 | 0.0531 | 375 | 0.9993 |
| a\_Var\_GYN | -0.0431 | 0.0181 | -0.0779 | -0.0065 | 419 | 0.9993 |
| a\_Var\_KRA | 0.0138 | 0.0174 | -0.0196 | 0.0500 | 374 | 0.9994 |
| a\_Var\_PRM | -0.0093 | 0.0174 | -0.0436 | 0.0254 | 389 | 0.9993 |
| sigma\_Species | 0.0384 | 0.0186 | 0.0165 | 0.0716 | 666 | 0.9996 |
| sigma | 0.0132 | 0.0015 | 0.0104 | 0.0163 | 648 | 1.0035 |

# Within Species

## Results – Body Mass Temp Model – Within Species

### ELN

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90
* Diagnostics:
  + Fairly strong multicollinearity between b\_W and b\_T (0.29) and b\_T and sigma (-0.18).
  + E-BFMI good.
  + Treedepth all ok.
  + 37 divergent iterations.

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| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2146 | 0.0051 | 0.2046 | 0.2248 | 5800 | 0.9999 |
| b\_W | 0.0015 | 0.0054 | -0.0095 | 0.0125 | 2666 | 1.0013 |
| b\_T | 0.0016 | 0.0056 | -0.0097 | 0.0120 | 396 | 1.0011 |
| sigma | 0.0155 | 0.0058 | 0.0042 | 0.0279 | 441 | 0.9998 |

### ELC

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 68 divergent iterations.
  + E-BFMI and treedepth ok.
  + Some autocorrelation between b\_T and b\_W (0.14).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1687 | 0.0032 | 0.1626 | 0.1751 | 3377 | 0.9998 |
| b\_W | -0.0034 | 0.0034 | -0.0103 | 0.0031 | 2211 | 0.9999 |
| b\_T | 0.0001 | 0.0027 | -0.0053 | 0.0056 | 834 | 1.0001 |
| sigma | 0.0055 | 0.0037 | 0.0007 | 0.0142 | 269 | 1.0061 |

### GYR

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 294 divergent iterations.
  + E-BFMI and treedepth ok.
  + Some multicolliniarity between b\_W and b\_T (0.55)

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2073 | 0.0032 | 0.2012 | 0.2136 | 1912 | 1.0000 |
| b\_W | -0.0051 | 0.0039 | -0.0129 | 0.0027 | 791 | 0.9998 |
| b\_T | -0.0007 | 0.0032 | -0.0072 | 0.0055 | 314 | 1.0001 |
| sigma | 0.0066 | 0.0037 | 0.0015 | 0.0152 | 223 | 1.0030 |

### **GYN**

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 145 divergent iterations.
  + E-BFMI and treedepth ok.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1429 | 0.0051 | 0.1327 | 0.1533 | 3366 | 0.9998 |
| b\_W | -0.0029 | 0.0054 | -0.0133 | 0.0081 | 3049 | 1.0005 |
| b\_T | -0.0005 | 0.0049 | -0.0099 | 0.0091 | 599 | 0.9999 |
| sigma | 0.0105 | 0.0061 | 0.0017 | 0.0246 | 465 | 1.0042 |

### KRA

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + E-BFMI and treedepth OK.
  + 86 divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1908 | 0.0035 | 0.1839 | 0.1977 | 2922 | 0.9998 |
| b\_W | 0.0015 | 0.0037 | -0.0059 | 0.0090 | 1963 | 1.0000 |
| b\_T | -0.0032 | 0.0031 | -0.0090 | 0.0035 | 401 | 1.0042 |
| sigma | 0.0078 | 0.0041 | 0.0018 | 0.0171 | 318 | 1.0022 |

### PRM

* 10000 iter, 5000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + E-BFMI and treedepth OK.
  + 46 divergent iterations.
  + Strong autocorrelation between b\_T and sigma.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1678 | 0.0051 | 0.1577 | 0.1779 | 4096 | 1.0007 |
| b\_W | 0.0115 | 0.0052 | 0.0012 | 0.0216 | 4816 | 0.9999 |
| b\_T | -0.0064 | 0.0041 | -0.0137 | 0.0027 | 618 | 1.0040 |
| sigma | 0.0140 | 0.0063 | 0.0038 | 0.0270 | 397 | 1.0003 |

## Results – Weight Model – Within Species

### ELN

* 3000 iter, 1500 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 0 divergent transitions.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2148 | 0.0046 | 0.2056 | 0.2237 | 1500 | 0.9993 |
| b\_W | 0.0011 | 0.0048 | -0.0084 | 0.0103 | 1500 | 0.9995 |
| sigma | 0.0176 | 0.0046 | 0.0095 | 0.0269 | 891 | 0.9993 |

### ELC

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 44 divergent transitions.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1688 | 0.0030 | 0.1630 | 0.1749 | 845 | 0.9997 |
| b\_W | -0.0034 | 0.0031 | -0.0096 | 0.0021 | 816 | 1.0002 |
| sigma | 0.0058 | 0.0033 | 0.0011 | 0.0122 | 162 | 1.0069 |

### GYR

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 1 divergent transition.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2072 | 0.0030 | 0.2015 | 0.2131 | 1194 | 1.0009 |
| b\_W | -0.0046 | 0.0031 | -0.0103 | 0.0019 | 917 | 0.9997 |
| sigma | 0.0070 | 0.0037 | 0.0011 | 0.0139 | 182 | 0.9997 |

### GYN

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 4 divergent transitions.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1430 | 0.0047 | 0.1338 | 0.1528 | 1967 | 1.000 |
| b\_W | -0.0032 | 0.0049 | -0.0132 | 0.0062 | 1764 | 0.9998 |
| sigma | 0.0117 | 0.0057 | 0.0017 | 0.0222 | 486 | 1.0002 |

### KRA

* 3000 iter, 1500 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 1 divergent transitions.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1907 | 0.0034 | 0.1838 | 0.1974 | 2109 | 0.9997 |
| b\_W | 0.0020 | 0.0035 | -0.0047 | 0.0094 | 1696 | 0.9999 |
| sigma | 0.0099 | 0.0041 | 0.0019 | 0.0172 | 287 | 0.9997 |

### PRM

* 3000 iter, 1500 warmup.
  + Adapt\_delta = 0.80.
* Diagnostics:
  + 1 divergent transitions.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1678 | 0.0052 | 0.1580 | 0.1789 | 1500 | 0.9994 |
| b\_W | 0.0125 | 0.0050 | 0.0034 | 0.0226 | 1500 | 0.9997 |
| sigma | 0.0198 | 0.0044 | 0.0116 | 0.0286 | 1064 | 0.9994 |

## Results – Temp Model – Within Species

### ELN

* 3000 iter, 1500 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 30 divergent iterations.
  + E-BFMI and treedepth ok.
  + Some autocorrelation between b\_T and sigma (-0.15).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2147 | 0.0048 | 0.2053 | 0.2243 | 2183 | 1.0002 |
| b\_T | 0.0011 | 0.0051 | -0.0090 | 0.0102 | 289 | 1.0055 |
| sigma | 0.0152 | 0.0039 | 0.0039 | 0.0239 | 355 | 1.0098 |

### ELC

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 219 divergent iterations.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1689 | 0.0030 | 0.1630 | 0.1753 | 1334 | 1.000 |
| b\_T | 0.0005 | 0.0029 | -0.0043 | 0.0068 | 74 | 0.9997 |
| sigma | 0.0049 | 0.0034 | 0.0006 | 0.0114 | 107 | 1.0355 |

### GYR

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 86 divergent iterations.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2068 | 0.0033 | 0.1998 | 0.2128 | 1990 | 1.0000 |
| b\_T | 0.0016 | 0.0029 | -0.0043 | 0.0070 | 188 | 1.0054 |
| sigma | 0.0074 | 0.0036 | 0.0019 | 0.0140 | 301 | 1.0045 |

### GYN

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 78 divergent iterations.
  + E-BFMI and treedepth OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1427 | 0.0052 | 0.1317 | 0.1521 | 1614 | 0.9988 |
| b\_T | -0.0005 | 0.0045 | -0.0096 | 0.0084 | 385 | 1.0077 |
| sigma | 0.0105 | 0.0053 | 0.0020 | 0.0206 | 418 | 1.0010 |

### KRA

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 91 divergent iterations.
  + E-BFMI and treedepth OK.
  + Some autocorrelation between b\_T and sigma (0.23).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1907 | 0.0032 | 0.1847 | 0.1969 | 1642 | 1.0011 |
| b\_T | -0.0029 | 0.0029 | -0.0080 | 0.0035 | 277 | 1.0023 |
| sigma | 0.0075 | 0.0037 | 0.0021 | 0.0142 | 255 | 1.0006 |

### PRM

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + 41 divergent iterations.
  + E-BMFI and treedepth OK.
  + Some autocorrelation between b\_T and sigma (0.53).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1678 | 0.0055 | 0.1580 | 0.1796 | 3000 | 1.0001 |
| b\_T | -0.0084 | 0.0045 | -0.0163 | 0.0013 | 204 | 1.0073 |
| sigma | 0.0141 | 0.0075 | 0.0014 | 0.0266 | 133 | 1.0066 |

## Results - Belcher

* 10000 iter, 5000 warmup.
* Diagnostics:
  + E-BFMI and treedepth ok.
  + 0 divergent iterations.
  + Autocorrelation OK.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1861 | 0.0030 | 0.1801 | 0.1919 | 5000 | 0.9998 |
| b | -0.0047 | 0.0029 | -0.0106 | 0.0010 | 5000 | 0.9999 |
| sigma | 0.0295 | 0.0022 | 0.0251 | 0.0336 | 5000 | 1.0007 |

### ELN

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics.
  + E-BFMI and treedepth OK.
  + 0 divergent transitions.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2156 | 0.0049 | 0.2063 | 0.2255 | 3000 | 0.9998 |
| b | -0.0030 | 0.0054 | -0.0129 | 0.0083 | 2493 | 0.9997 |
| sigma | 0.0169 | 0.0050 | 0.0077 | 0.0269 | 1622 | 0.9998 |

### ELC

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics.
  + E-BFMI and treedepth OK.
  + 121 divergent iterations.
  + Some autocorrelation between b\_M and sigma.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1664 | 0.0040 | 0.1584 | 0.1742 | 1743 | 0.9997 |
| b | 0.0033 | 0.0042 | -0.0053 | 0.0120 | 1265 | 0.9998 |
| sigma | 0.0063 | 0.0044 | 0.0008 | 0.0148 | 112 | 1.0073 |

### GYR

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics.
  + E-BFMI and treedepth OK.
  + 23 divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.2069 | 0.0030 | 0.2006 | 0.2123 | 1615 | 1.0019 |
| b | 0.0051 | 0.0030 | -0.0008 | 0.0114 | 1076 | 1.0000 |
| sigma | 0.0068 | 0.0036 | 0.0009 | 0.0134 | 190 | 1.0042 |

### GYN

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics.
  + E-BFMI and treedepth OK.
  + 47 divergent iterations.
  + Autocorrelation between a and b (0.90).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1427 | 0.0051 | 0.1327 | 0.1527 | 1709 | 0.9998 |
| b | 0.0016 | 0.0052 | -0.0089 | 0.0119 | 1128 | 0.9997 |
| sigma | 0.0121 | 0.0059 | 0.0014 | 0.0225 | 359 | 0.9999 |

### PRM

* 6000 iter, 3000 warmup.
  + Adapt\_delta = 0.90.
* Diagnostics.
  + E-BFMI and treedepth OK.
  + 10 divergent iterations.
  + Autocorrelation between b\_M and sigma (0.43).

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1678 | 0.0046 | 0.1587 | 0.1768 | 3000 | 0.9997 |
| b | -0.0145 | 0.0047 | -0.0239 | -0.0055 | 715 | 1.0006 |
| sigma | 0.0123 | 0.0058 | 0.0024 | 0.0227 | 283 | 1.0010 |

### KRA

* Iter = 6000, warmup = 3000.
  + Adapt\_delta = 0.90.
* Diagnostics:
  + E-BFMI and treedepth OK.
  + 41 divergent iterations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Parameter** | **Mean** | **SD** | **Lower 95%** | **Upper 95%** | **N\_eff** | **R\_hat** |
| a | 0.1906 | 0.0033 | 0.1845 | 0.1973 | 2024 | 1.0008 |
| b | -0.0028 | 0.0037 | -0.0105 | 0.0041 | 416 | 0.9997 |
| sigma | 0.0090 | 0.0042 | 0.0015 | 0.0161 | 132 | 1.0082 |