

BDA - Assignment 7

Anonymous

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Exercise 1

Exercise 1 A

```
In [1]: #read the bioassay data
import numpy as np
import cv2
import pystan

#reading the data
file = open("c:\\downing.txt", "r")
temp = file.read().splitlines()

years = []
drowned = []
for line in temp:
    values = line.split(" ")
    years.append(float(values[0]))
    drowned.append(float(values[1]))
years = np.array(years)
drowned = np.array(drowned)
```

The mistakes were that: 1) `real<lower=0> sigma` should have been `real<lower=0> sigma`

2) There is no computation of `mu_pred` required for finding out ypred.

3) `ypred = normal_rng(mu, sigma)` should have been `ypred = normal_rng(mu_pred, sigma)`

```
In [8]: fixed_model = '''
data {
    int<lower=0> N; // number of data points
    vector<N> y; // observation year
    vector<N> y; // observation number of drowned
    real xpred; // prediction year
}
parameters {
    real alpha;
    real beta;
    real<lower=0> sigma;
}
transformed parameters {
    vector<N> mu;
    mu = alpha + beta * x;
    pred_mu = alpha + beta * xpred;
}
model {
    y ~ normal(mu, sigma);
}
generated quantities {
    real ypred;
    ypred = normal_rng(pred_mu, sigma);
}
'''
data = {
    "N": len(years),
    "x": years,
    "y": drowned,
    "xpred": 2020,
}
```

```
In [17]: posterior = pystan.StanModel(model_code=fixed_model)
fit = posterior.sampling(data=data)
print(fit)
```

INFO:pystan:COMPILING THE C++ CODE FOR MODEL anon_model_5d2a30d49c6b11b7b660b34acbd17f6 NOW.
WARNING:pystan:941 of 4000 iterations saturated the maximum tree depth of 10 (23.5 %)
WARNING:pystan:Run again with max_treedepth larger than 10 to avoid saturation
Inference for Stan model: anon_model_5d2a30d49c6b11b7b660b34acbd17f6:
4 chains, each with iter=2000; warmup=1000; thin=1;
post-warmup draws per chain=1000, total post-warmup draws=4000.

	mean	se_mean	sd	2.5%	25%	50%	75%	97.5%	n_eff	Rhat
alpha	2464.2	21.98	728.57	1079.6	1984.2	2472.3	2925.8	3936.1	1099	1.0
beta	-1.17	0.01	0.36	-1.9	-1.4	-1.17	-0.93	-0.47	1099	1.0
sigma	26.35	0.08	3.08	21.15	24.2	26.11	28.14	33.24	1393	1.0
mu[1]	157.0	0.22	8.25	140.5	151.62	156.97	162.45	173.44	1371	1.0
mu[2]	155.84	0.21	7.94	139.97	150.7	155.83	161.05	171.66	1398	1.0
mu[3]	154.67	0.2	7.63	139.43	149.76	154.62	159.69	169.91	1429	1.0
mu[4]	153.51	0.19	7.33	138.93	148.79	153.47	158.34	168.12	1465	1.0
mu[5]	152.14	0.18	7.03	138.52	147.82	152.31	157.12	166.32	1509	1.0
mu[6]	151.18	0.17	6.74	138.08	146.8	151.08	155.64	164.56	1561	1.0
mu[7]	150.01	0.16	6.46	137.38	145.79	149.54	154.3	162.88	1624	1.0
mu[8]	148.85	0.15	6.19	136.78	144.78	148.72	152.96	161.15	1698	1.0
mu[9]	147.68	0.14	5.92	136.18	143.78	147.53	151.61	159.55	1791	1.0
mu[10]	146.52	0.13	5.67	135.46	142.78	146.41	150.23	157.87	1892	1.0
mu[11]	145.35	0.12	5.43	134.65	141.81	145.24	148.95	156.14	2010	1.0
mu[12]	144.19	0.11	5.2	133.96	140.82	144.09	147.63	154.45	2154	1.0
mu[13]	143.02	0.1	4.99	133.25	139.73	142.86	146.32	152.88	2329	1.0
mu[14]	141.86	0.1	4.8	132.42	138.72	141.82	145.03	151.48	2542	1.0
mu[15]	140.69	0.1	4.63	131.57	137.63	140.61	143.76	149.88	2797	1.0
mu[16]	139.52	0.08	4.49	130.76	136.6	139.44	142.49	148.46	3095	1.0
mu[17]	138.36	0.07	4.37	129.84	135.5	138.31	141.22	147.14	3415	1.0
mu[18]	137.19	0.07	4.27	128.77	134.2	137.14	140.0	145.87	3696	1.0
mu[19]	136.03	0.07	4.21	127.71	133.24	136.0	138.79	144.53	3932	1.0
mu[20]	134.86	0.07	4.18	126.67	132.1	134.81	137.63	143.18	4025	1.0
mu[21]	133.7	0.07	4.17	125.61	131.01	133.64	136.48	141.95	4031	1.0
mu[22]	132.53	0.07	4.2	124.53	129.77	132.48	135.32	140.9	3970	1.0
mu[23]	131.37	0.07	4.27	123.32	128.6	131.3	134.18	139.9	3756	1.0
mu[24]	130.2	0.07	4.36	122.13	127.34	129.13	133.07	139.9	3484	1.0
mu[25]	129.04	0.08	4.48	120.9	126.5	127.0	131.9	137.9	3190	1.0
mu[26]	127.87	0.09	4.62	118.74	124.83	127.81	130.97	137.03	2856	1.0
mu[27]	126.71	0.09	4.79	117.19	123.57	126.68	129.92	136.22	2594	1.0
mu[28]	125.54	0.1	4.98	115.53	122.27	125.56	128.8	135.31	2392	1.0
mu[29]	124.38	0.11	5.19	113.98	121.0	124.39	127.82	134.48	2193	1.0
mu[30]	123.21	0.12	5.41	112.39	119.67	122.69	126.79	133.86	2044	1.0
mu[31]	122.05	0.13	5.65	110.78	118.57	122.05	125.73	133.17	1907	1.0
mu[32]	120.88	0.14	5.9	109.09	117.04	120.93	124.77	132.45	1787	1.0
mu[33]	119.72	0.15	6.17	107.33	115.71	119.73	123.8	131.77	1690	1.0
mu[34]	118.55	0.16	6.44	105.6	114.3	118.55	122.83	131.19	1610	1.0
mu[35]	117.38	0.17	6.72	103.83	112.86	117.4	121.6	130.43	1594	1.0
mu[36]	116.22	0.18	7.01	102.12	111.77	116.22	120.84	129.86	1488	1.0
mu[37]	115.05	0.19	7.31	100.42	110.44	115.04	119.88	129.32	1443	1.0
mu[38]	113.89	0.2	7.62	98.69	109.16	113.89	118.96	128.7	1406	1.0
mu[39]	112.72	0.21	7.92	96.94	107.7	112.72	117.93	128.17	1375	1.0
mu[40]	111.56	0.22	8.23	95.2	106.36	111.56	116.95	127.67	1347	1.0
pred_mu[10]	139.39	0.23	8.54	93.2	105.02	110.38	116.01	127.12	1163	1.0
pred_mu[11]	138.17	0.5	28.15	55.4	91.86	110.51	129.34	165.24	3187	1.0
lp__	-146.6	0.04	1.25	-149.8	-147.2	-146.3	-145.7	-145.2	1320	1.0

Samples were drawn using NUTS at Thu Oct 20 02:15:55 2022.
For each parameter, `n_eff` is a crude measure of effective sample size, and `Rhat` is the potential scale reduction factor on split chains (at convergence, `Rhat=1`).

Exercise 1 B

We know that beta follows a normal distribution $N(0, \sigma)$.

We also know that the 0.05 percentile is -69, and that the 0.95 percentile is 69.

Ideally, what we would try to do in this case is to find out σ would be to solve for the integral between values -69 and 69 and equate that to 69. But given that the Normal Distribution's form does not have a compact integral, we need to find other ways to solve this.

That's why I will be using the inverse error function (computed I assume with the Taylor Expansion under the hood) to approximate the correct sigma value.

We know that the formula of the CDF is $\frac{1}{2} * [1 + \text{erf}(\frac{x-\mu}{\sigma\sqrt{2}})]$. Starting from this, we can prove that the formula for a quantile is $\mu + \sigma\sqrt{2} * \text{erf}^{-1}(2p - 1)$.

(We also know that μ is 0)

Therefore,

$$69 = 0 + \sigma\sqrt{2} * \text{erf}^{-1}(2 * 0.95 - 1)$$

$$69 = 0 + \sigma\sqrt{2} * \text{erf}^{-1}(0.99)$$

$$69 = 0 + \sigma\sqrt{2} * 1.8213...$$

$$\sigma = 26.79, \text{ using SciPy's erfinv function}$$

```
In [9]: from scipy.special import erfinv
beta_sigma = (69 / (erfinv(0.99) * np.sqrt(2))).round(2)
print(f"Value of beta's sigma is {beta_sigma}")
```

Value of beta's sigma is 26.79

Exercise 1 C

```
In [10]: fixed_model_beta_prior = '''
data {
    int<lower=0> N; // number of data points
    vector<N> x; // observation year
    vector<N> y; // observation number of drowned
    real xpred; // prediction year
}
parameters {
    real alpha;
    real beta;
    real<lower=0> sigma;
}
transformed parameters {
    vector<N> mu;
    mu = alpha + beta * x;
    pred_mu = alpha + beta * xpred;
}
model {
    y ~ normal(0, 26.79); //added this line to account for beta's prior
}
generated quantities {
    real ypred;
    ypred = normal_rng(pred_mu, sigma);
}
'''
data = {
    "N": len(years),
    "x": years,
    "y": drowned,
    "xpred": 2020,
}
```

Solved this by adding line `beta ~ normal(0, 26.79);` in block `model`

Exercise 1 D

My thought process for this exercise is that the intercept should be somewhat compensating for the values of $\beta * x$. Taking this into consideration, given that beta's mean is 0, and the assignment tells us the mean of drownings is 138, this means we'll set the mean of our intercept's prior to 138.

The question is now how to find the variance. We'll apply the same logic we did when computing the variance for the β hyperparameter.

Given that the mean of our x's value is 2000, we'll ask that $P(-2000 + 0 < \alpha < 2000 + 69) = 0.99$.

Using the same function as before, we get that:

```
In [11]: beta_sigma = ((2000+69 - 138) / (erfinv(0.99) * np.sqrt(2))).round(2)
print(f"Value of beta's sigma is {beta_sigma}")
```

Value of beta's sigma is 53521.4

Therefore the prior of alpha will be $N(138, 53521.4)$

```
In [19]: fixed_model_beta_alpha_prior = '''
data {
    int<lower=0> N; // number of data points
    vector<N> x; // observation year
    vector<N> y; // observation number of drowned
    real xpred; // prediction year
}
parameters {
    real alpha;
    real beta;
    real<lower=0> sigma;
}
transformed parameters {
    vector<N> mu;
    mu = alpha + beta * x;
    pred_mu = alpha + beta * xpred;
    give_in = alpha + beta * xpred;
}
model {
    alpha ~ normal(138, 53521.4); // added this line to account for alpha's prior
    beta ~ normal(0, 26.79); // added this line to account for beta's prior
    y ~ normal(mu, sigma);
}
generated quantities {
    real ypred;
    ypred = normal_rng(pred_mu, sigma);
}
'''
data = {
    "N": len(years),
    "x": years,
    "y": drowned,
    "xpred": 2020,
}
```

This is the line added for alpha's prior: `alpha ~ normal(138, 53521.4).`

```
In [20]: posterior = pystan.StanModel(model_code=fixed_model_beta_alpha_priors)
fit = posterior.sampling(data=data)
print(fit)
```

INFO:pystan:COMPILING THE C++ CODE FOR MODEL anon_model_7405d454be49856d06f53ff9737e8f6 NOW.
WARNING:pystan:947 of 4000 iterations saturated the maximum tree depth of 10 (21.2 %)
WARNING:pystan:Run again with max_treedepth larger than 10 to avoid saturation
Inference for Stan model: anon_model_7405d454be49856d06f53ff9737e8f6:
4 chains, each with iter=2000; warmup=1000; thin=1;
post-warmup draws per chain=1000, total post-warmup draws=4000.

```

quality[q1].append(int(values[i]))
    qi += 1

```

First of all, I will explain why I think the model shown in the assignment gives strange results:

The reason for that is the low variance on both μ and σ which impose bias. If the number of samples is low and the prior distribution is narrow, then its influence on the posterior will be high and the data will be neglected. This is why in all of the examples below I made sure to choose a weakly informative prior with high variance for the normal distribution for μ and a wider distribution for σ .

More on this topic could be found here: <https://discourse.mc-stan.org/t/standardizing-predictors-and-outputs-in-a-hierarchical-model/2974/9>

Additionally, there is also a mismatch between the unit scale of the data and the unit scale of the prior. This explanation was taken from https://mc-stan.org/users/documentation/case-studies/weakly_informative_shapes.html.

The differences between the 3 types of models: 1) The separate model builds itself on the assumption that all the machines are independent of each other. That's why each of the values for the priors (we have 6 priors) are independent draws from a pre-defined distribution. And this is why we cannot use this kind of model to infer the behaviour of a new 7th machine.

2) The pooled model builds itself on the assumption that all the machines are the same, thus all of their samples can be treated as belonging to a single machine, a pooled one. That is why there is a single pair of parameters μ and σ which result in a single prior.

3) The hierarchical model is the middleground solution between the separate model and the pooled model. It builds on the assumption that all the machines have some common ground, but they can all behave differently. The way it achieves this is by allowing each of the machines to have its own prior, but all of the priors are "sampled" from a distribution defined tunable hyperparameters. This way, we achieve both the independence of the separate model and the similarity of the pooled model.

Separate Model

Exercise 2 A

$$y_{ij} \sim N(\mu_j, \sigma_j)$$

$$\mu_j \sim N(0, 50)$$

$$\sigma_j \sim \text{Inv} - \chi^2(1)$$

The reason for choosing a variance of 50 for the μ_j term is due to the low number of samples, which would have been completely neglected if we were to choose a low variance, and therefore the value of the mean would be biased (See explanations in the beginning of the exercise). I tried with 1, and the mean was

