

# Bayesian Synthesis Models

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Data Confidentiality

# Outline

- 1 Introduction
- 2 Preserving relationships and Bayesian models
- 3 Bayesian synthesis models estimation
- 4 Generating synthetic values for sensitive variables
- 5 Miscellany

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# Adding random noise

- Add noise to
  - ▶ provide privacy protection: noise.
  - ▶ preserve important relationships: signal.
- For a sensitive continuous variable, e.g. family income  $Y_i$ , we can add noise  $Y_i^*$  from a normal centered at 0 (why 0?):

$$Y_i^* \sim \text{Normal}(0, \sigma)$$

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- This approach preserves distributional characteristics of family income, but not its relationships with other variable(s).

# The CE sample

- The Consumer Expenditure Surveys (CE)
  - ▶ conducted by the U.S. Census Bureau for the U.S. Bureau of Labor Statistics.
  - ▶ contains data on expenditures, income, and tax statistics about consumer units (CU) across the country.
  - ▶ provides information on the buying habits of U.S. consumers.

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```
CEdata <- read.csv(file = "CEdata.csv")
head(CEdata)
```

##	UrbanRural	Income	Race	Expenditure
## 1	1	98600	1	5972.167
## 2	1	24360	1	5854.500
## 3	1	80200	1	5506.667
## 4	1	150500	1	8968.891
## 5	1	130000	1	10092.833
## 6	1	32836	1	5520.267

# The CE sample cont'd

Variable	Information
UrbanRural	Binary; the urban / rural status of CU: 1 = Urban, 2 = Rural.
Income	Continuous; the amount of CU income before taxes in past 12 months.
Race	Categorical; the race category of the reference person: 1 = White, 2 = Black, 3 = Native American, 4 = Asian, 5 = Pacific Islander, 6 = Multi-race.
Expenditure	Continuous; CU's total expenditures in last quarter.



# The synthesis goals and questions

- Suppose Income is deemed the most sensitive among the four variables, and we want to add noise to it for privacy protection.
- To complete this, we can consider the following questions:
  - ① What kind of relationships do you think are important to preserve in the confidential data?

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  - ④ And finally, how do you evaluate whether your methods have achieved your goals?
- This lecture focuses on Questions 1-3.

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- What kind of relationships between `Income` and the other three (`UrbanRural`, `Race`, `Expenditure`) do you think are important to preserve? Why?
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  - ▶ Inferences done on synthetic data are “close” to those done on confidential data.



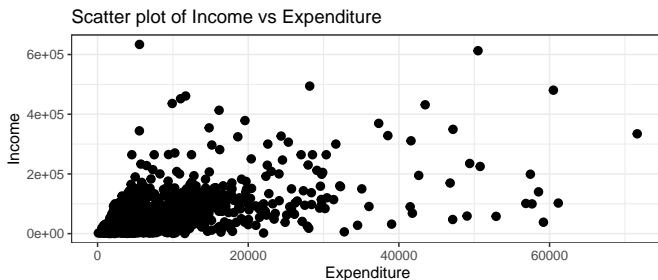
# Important relationships?

- What kind of relationships between `Income` and the other three (`UrbanRural`, `Race`, `Expenditure`) do you think are important to preserve? Why?
- When we say “preserve”, what do we mean?
  - ▶ Inferences done on synthetic data are “close” to those done on confidential data.
  - ▶ However, too “close” could mean less privacy protection.
  - ▶ We need to strike the balance between utility and disclosure risks.

## Example: Relationship between Income and Expenditure

- Suppose we want to preserve the relationship between Income and Expenditure.
- Visualizing this relationship on the raw scale of these two variables shows fanning trend.

```
ggplot(CEdata, aes(x = Expenditure, y = Income)) +  
  geom_point(size = 1) +  
  labs(title = "Scatter plot of Income vs Expenditure") +  
  theme_bw(base_size = 6, base_family = "")
```



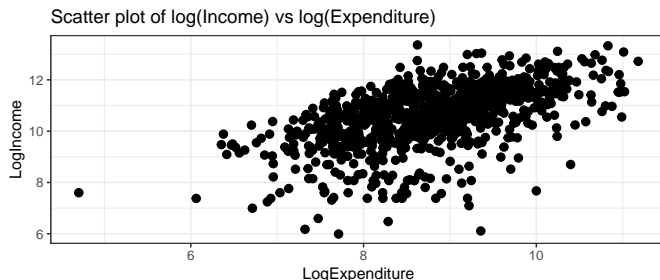
# Income and Expenditure: log scale

- On the log scale, the relationship between  $\log(\text{Income})$  and  $\log(\text{Expenditure})$  appears to be linear.

```
CEdata$LogIncome <- log(CEdata$Income)
CEdata$LogExpenditure <- log(CEdata$Expenditure)
```

```
ggplot(CEdata, aes(x = LogExpenditure, y = LogIncome)) +
  geom_point(size = 1) +
  labs(title = "Scatter plot of log(Income) vs log(Expenditure)") +
  theme_bw(base_size = 6, base_family = "")
```

# Income and Expenditure: log scale cont'd



- To capture and preserve this linear relationship, a Bayesian linear regression model of  $\log(\text{Expenditure})$  on  $\log(\text{Income})$  seems a good choice.

# A Bayesian simple linear regression model

- Let  $Y_i$  be the  $\log(\text{Income})$  and  $X_i$  be the  $\log(\text{Expenditure})$  for CU  $i$ , a Bayesian simple linear regression model can be expressed as:

$$Y_i \mid \mu_i, \sigma \stackrel{\text{ind}}{\sim} \text{Normal}(\mu_i, \sigma) \quad (1)$$

$$\mu_i = \beta_0 + \beta_1 X_i \quad (2)$$

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- The expected  $\log(\text{Income})$  is  $\mu_i$ , which is a linear function of  $\log(\text{Expenditure})$   $X_i$  through the intercept parameter  $\beta_0$  and the slope parameter  $\beta_1$ .
- The intercept  $\beta_0$ : the expected  $\log(\text{Income})$   $\mu_i$  for CU  $i$  that has zero  $\log(\text{Expenditures})$  (i.e.  $X_i = 0$ ).
- The slope  $\beta_1$ : the change in the expected  $\log(\text{Income})$   $\mu_i$  when the  $\log(\text{Expenditures})$  of CU  $i$  increases by 1 unit (i.e.  $X_i$  increases by  $\$ \log(1)$ ).

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## A weakly informative prior

- To estimate the proposed Bayesian linear regression model, we need to assign appropriate prior distributions for all parameters in the model:  $\{\beta_0, \beta_1, \sigma\}$ .



## A weakly informative prior

- To estimate the proposed Bayesian linear regression model, we need to assign appropriate prior distributions for all parameters in the model:  $\{\beta_0, \beta_1, \sigma\}$ .
- If we have limited prior information about these parameters, we could use a weakly informative prior distribution. Assuming independence of the three parameters:

$$\pi(\beta_0, \beta_1, \sigma) = \pi(\beta_0)\pi(\beta_1)\pi(\sigma). \quad (3)$$

# A weakly informative prior cont'd

- We can then give individual weakly informative prior for each parameter:

$$\beta_0 \sim \text{Normal}(\mu_0, s_0) \quad (4)$$

$$\beta_1 \sim \text{Normal}(\mu_1, s_1) \quad (5)$$

$$1/\sigma^2 \sim \text{Gamma}(a, b), \quad (6)$$

where we can use  $\mu_0 = \mu_1 = 0$ ,  $s_0 = s_1 = 100$ , and  $a = b = 1$ .

# MCMC simulation by JAGS

- Let's use JAGS (Just Another Gibbs Sampler) to estimate our chosen Bayesian simple linear regression model.

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# MCMC simulation by JAGS

- Let's use JAGS (Just Another Gibbs Sampler) to estimate our chosen Bayesian simple linear regression model.
- We will obtain pre-specified number of posterior parameter draws from the JAGS output, which will be used for synthetic data generation through the posterior predictive distribution.
- Make sure that we require the `runjags` and `coda` libraries.

```
require(runjags)  
require(coda)
```

# Using JAGS: part 1

- Describe the model by a script.

```
modelString <-"
model {
  ## sampling
  for (i in 1:N){
    y[i] ~ dnorm(beta0 + beta1*x[i], invsigma2)
  }

  ## priors
  beta0 ~ dnorm(mu0, g0)
  beta1 ~ dnorm(mu1, g1)
  invsigma2 ~ dgamma(a, b)
  sigma <- sqrt(pow(invsigma2, -1))
}
"
```

## Using JAGS: part 2

- Define the data and prior parameters.

```

y <- as.vector(CEdata$LogIncome)
x <- as.vector(CEdata$LogExpenditure)
N <- length(y)
the_data <- list("y" = y, "x" = x, "N" = N,
                 "mu0" = 0, "g0" = 0.0001,
                 "mu1" = 0, "g1" = 0.0001,
                 "a" = 1, "b" = 1)

initsfunction <- function(chain){
  .RNG.seed <- c(1,2)[chain]
  .RNG.name <- c("base::Super-Duper",
                 "base::Wichmann-Hill")[chain]
  return(list(.RNG.seed=.RNG.seed,
              .RNG.name=.RNG.name))
}

```

## Using JAGS: part 3

- Generate samples from the posterior distribution.

```
posterior <- run.jags(modelString,  
                      n.chains = 1,  
                      data = the_data,  
                      monitor = c("beta0", "beta1", "sigma"),  
                      adapt = 1000,  
                      burnin = 5000,  
                      sample = 5000,  
                      thin = 50,  
                      inits = initsfunction)
```

- The value of thin is set given MCMC diagnostics.

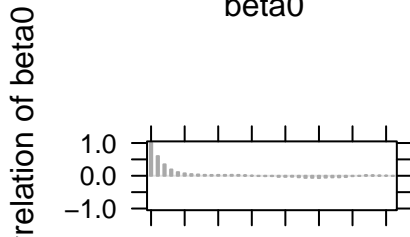
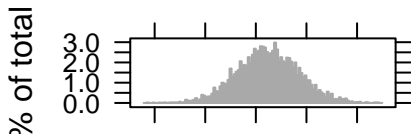
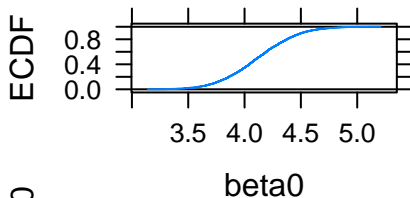
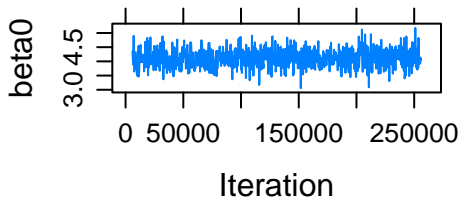


## Using JAGS: part 4

- MCMC diagnostics (check  $\beta_1$  and  $\sigma$  as well).

```
plot(posterior, vars = "beta0")
```

```
## Generating plots...
```



## Using JAGS: part 5

- Saving posterior parameter draws.

```
post <- as.mcmc(posterior)
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- `post` contains 5000 rows and 3 columns:
  - ▶ each column corresponds to a parameter: `beta0`, `beta1`, and `sigma`.
  - ▶ each row corresponds to one of the 5000 MCMC iterations.

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- `post` contains 5000 rows and 3 columns:
  - ▶ each column corresponds to a parameter: `beta0`, `beta1`, and `sigma`.
  - ▶ each row corresponds to one of the 5000 MCMC iterations.
- Next, we will use the posterior parameter draws saved in `post` to generate synthetic values for `log(Income)`.

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# Generating one set of synthetic data

- The synthetic data generation process is no different from making prediction of future values. To predict future  $\log(\text{Income})$ ,  $\tilde{Y}_i$  for a CU given its  $\log(\text{Expenditure})$ ,  $X_i$ :

$$\tilde{Y}_i \mid \beta_0, \beta_1, \sigma \stackrel{\text{ind}}{\sim} \text{Normal}(\beta_0 + \beta_1 X_i, \sigma). \quad (7)$$

# Generating one set of synthetic data cont'd

- Given  $X_i$  and one set of posterior draws of the parameters  $\{\beta_0, \beta_1, \sigma\}$ , we could simulate  $\tilde{Y}_i$  for each all  $n$  CUs:

simulate  $E[Y_1] = \beta_0 + \beta_1 X_1 \rightarrow$  sample  $\tilde{Y}_1 \sim \text{Normal}(E[Y_1], \sigma)$

simulate  $E[Y_2] = \beta_0 + \beta_1 X_2 \rightarrow$  sample  $\tilde{Y}_2 \sim \text{Normal}(E[Y_2], \sigma)$

$\vdots$

simulate  $E[Y_n] = \beta_0 + \beta_1 X_n \rightarrow$  sample  $\tilde{Y}_n \sim \text{Normal}(E[Y_n], \sigma)$



## Generating one set of synthetic data cont'd

- Suppose we use **one set of posterior draws**, the function below returns a synthetic dataset with synthesized  $\log(\text{Income})$  and un-synthesized  $\log(\text{Expenditure})$ .

```
synthesize <- function(X, index, n){  
  mean_Y <- post[index, "beta0"] + X * post[index, "beta1"]  
  synthetic_Y <- rnorm(n, mean_Y, post[index, "sigma"])  
  data.frame(X, synthetic_Y)  
}
```

- The input  $X$  is a vector of the un-synthesized variable, i.e.  $\log(\text{Expenditure})$ .
- $\text{index}$  indicates which set of posterior draws to be used.

## Generating one set of synthetic data cont'd

- For example, `index = 1` if we use the first of 5000 sets. `n` is the number of observations.

```
synthesize <- function(X, index, n){
  mean_Y <- post[index, "beta0"] + X * post[index, "beta1"]
  synthetic_Y <- rnorm(n, mean_Y, post[index, "sigma"])
  data.frame(X, synthetic_Y)
}
```

```
n <- dim(CEdata)[1]
synthetic_one <- synthesize(CEdata$LogExpenditure, 1, n)
names(synthetic_one) <- c("logExpenditure", "logIncome_syn")
```

# Generating multiple sets of synthetic data

- Typical practice generates **multiple sets of synthetic data**, for example,  $m = 20$ .
- Later, we will explore why multiple synthetic datasets are needed for data utility evaluation.

## Generating multiple sets of synthetic data cont'd

- To use the last  $m = 20$  sets of the obtained 5000 MCMC iterations for generating  $m = 20$  synthetic datasets:

```
n <- dim(CEdata)[1]
m <- 20
synthetic_m <- vector("list", m)
for (l in 1:m){
  synthetic_one <- synthesize(CEdata$LogExpenditure, 4980+1, n)
  names(synthetic_one) <- c("logExpenditure", "logIncome_syn")
  synthetic_m[[l]] <- synthetic_one
}
```

## Generating multiple sets of synthetic data cont'd

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  names(synthetic_one) <- c("logExpenditure", "logIncome_syn")
  synthetic_m[[l]] <- synthetic_one
}
```

- Use a list `synthetic_m` to save all  $m = 20$  synthetic datasets.
- Each synthetic dataset contains **the un-synthesized** `logExpenditure`, and **the synthesized** `logIncome_syn` from the estimated Bayesian simple linear regression model.

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# Preserve important relationships

- If `Expenditure` is deemed sensitive, what relationships do you want to preserve, why, and how?
- If `UrbanRural` is deemed sensitive, what relationships do you want to preserve, why, and how?
- If `Race` is deemed sensitive, what relationships do you want to preserve, why, and how?

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- If `Race` is deemed sensitive, what relationships do you want to preserve, why, and how?
- If `Income` and `UrbanRural` are deemed sensitive, what relationships do you want to preserve, why, and how?



# Partial synthesis vs full synthesis

- Partial synthesis: only a subset of variables / attributes are deemed sensitive and to be synthesized.
- Full synthesis: all variables / attributes are deemed sensitive and to be synthesized.

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- Partial synthesis: only a subset of variables / attributes are deemed sensitive and to be synthesized.
- Full synthesis: all variables / attributes are deemed sensitive and to be synthesized.
- Implications for Bayesian synthesis models?
- Implications for disclosure risks evaluation?