

# BDA - Project

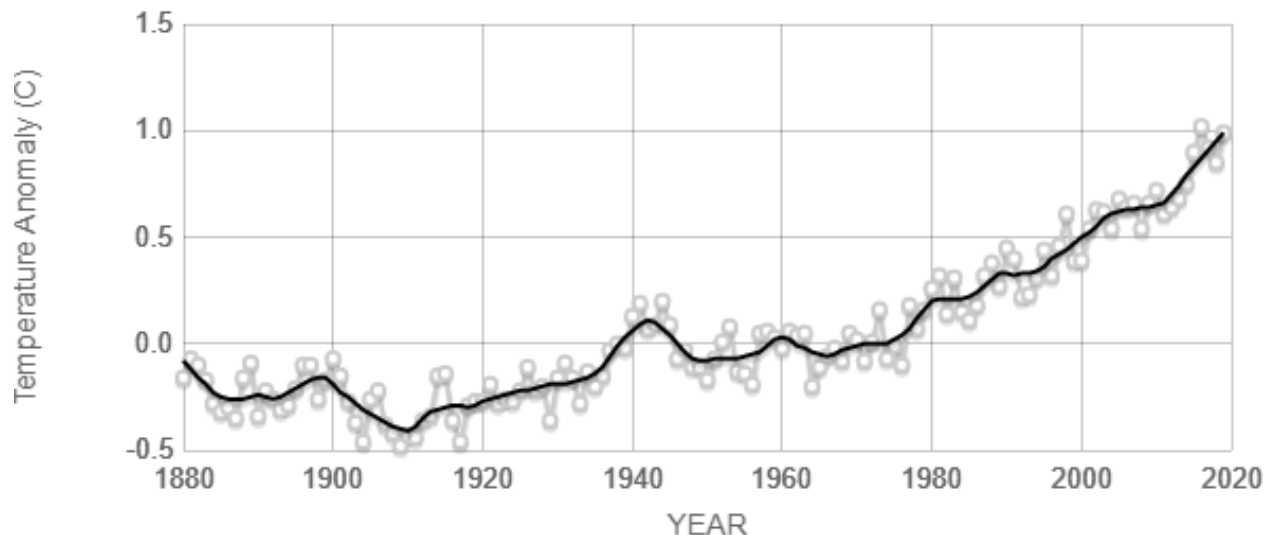
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# 1. Introduction

One of the biggest challenges of humankind in the 2020s is figuring out ways to slow down the growth of greenhouse gas emissions and stop global warming (due to human activities) under 2 °C. The increasing trend of global temperature is easily seen in Figure 1 [cite NASA] in which the global surface temperature is illustrated relative to 1951-1980 average temperatures. Warming can also be seen with one's own eyes by observing the winters that are warming year by year, by noticing that the number of devastating hurricanes is increased, and by finding out the increased rate of ice melting in glaciers during summer.



Source: [climate.nasa.gov](https://climate.nasa.gov)

Figure 1: Global Land-Ocean Temperature Index

In response to that warming, many countries have declared a climate emergency to emphasize the criticality of the situation. In addition, young people have organized climate demonstrations around the world, politicians are talking more and more about climate change, and presidents and prime ministers are negotiating agreements and commitments to solve this, one of humanity's greatest, problem. But what if, despite attempts of negotiation, the necessary CO<sub>2</sub> reduction decisions are not achieved?

In this project, our goal is to model the historical emission trends of selected countries as well as attempts to model their future emissions. We are examining a scenario in which emissions continue to develop at a historical rate, and the necessary reductions are not achieved. In our modeling, the other parameters e.g. population growth and technical conditions, are similar to historical data in our modeling.

## 2. Data description

Our CO<sub>2</sub> data was obtained from *Our World in Data* (OWID) web page [cite OWID\_net] and the actual CSV file from OWID GitHub page [cite OWID\_git]. As mentioned earlier, climate change is a hot topic in the daily news, and there is a lot of studies and research concerning how CO<sub>2</sub> emissions are influencing global warming. The data set was also used, for example, when researchers studied the climate impact of the different policy recommendation which targeted to reduce greenhouse gases from the atmosphere.

### 2.1 Choosing the sample and estimating it's resemblance

In our modeling, we selected 19 different countries from the OWID data set and examined CO<sub>2</sub> data between the years 1950-2018. We decided to not take all countries into the modeling as there are holes and missing

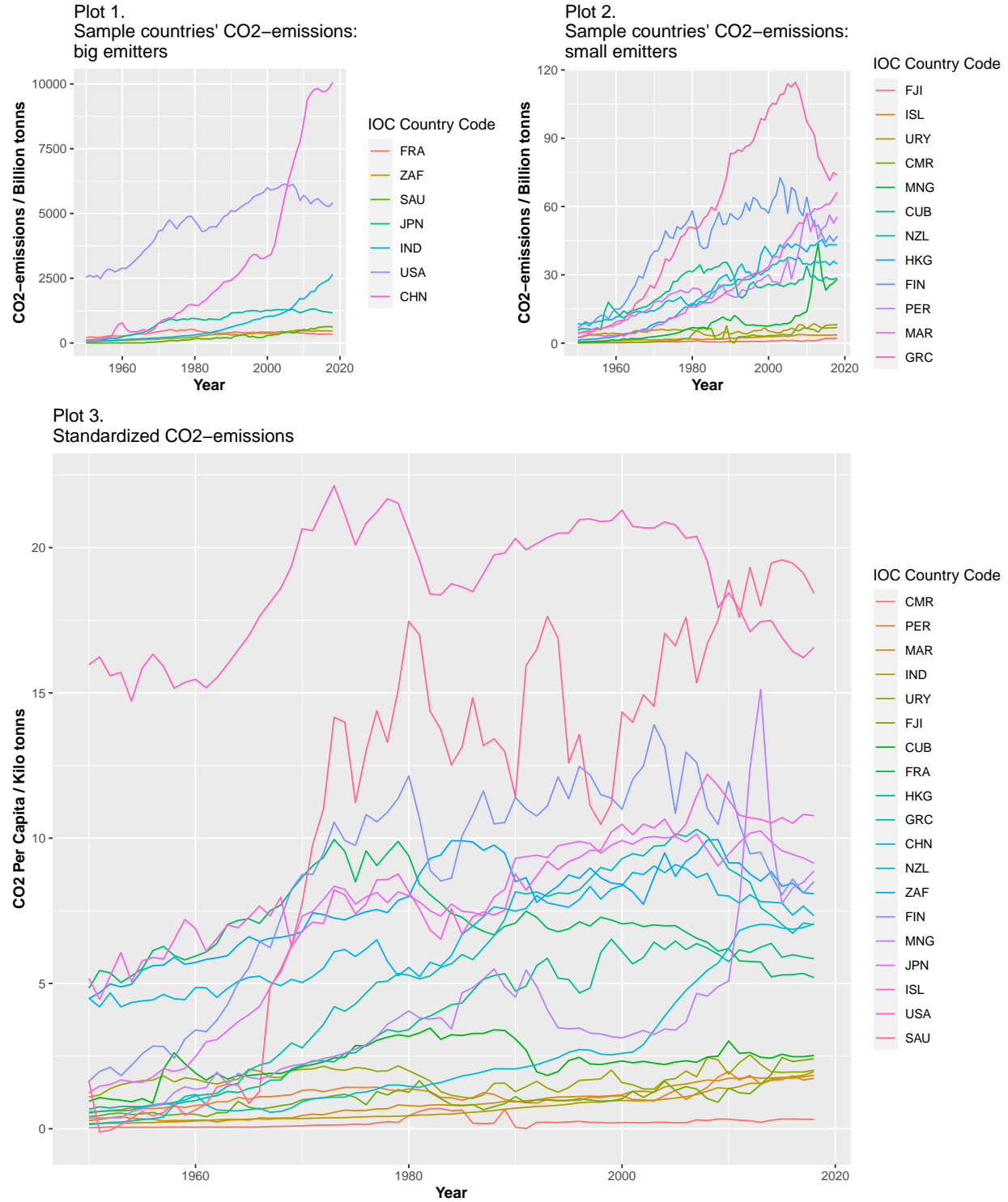
information in the dataset. The countries we chose cover the whole globe and are roughly evenly distributed across continents. However, we estimated that the data is probably more reliable in the western countries and thus were more open-minded in selecting them. Even that said, we think that the geographical distribution covers the whole world pretty well. Another important aspect of division is the division between large and small emitters. Even though it is quite difficult to perform such division, we tried to take countries from both ends pretty evenly. However, it is worth noting that this division was performed intuitively and it does not rely on any actual metrics. Lastly, we thought that the division between developing and western countries is extremely important to consider too. Therefore, this aspect was taken into account when considering the sample countries, too. We estimated that the number of developing countries in the world exceeds the number of western countries and thus tried to choose developing countries a bit more into the sample set.

For the reasons presented above, we believe that the sample we use in this project, resembles the situation in the world quite well. However, we estimated that it is possible that the sample is slightly biased towards western countries. It is important to note this since we examine results where the CO<sub>2</sub>-emissions data is standardized with the countries' population. As the CO<sub>2</sub>-emissions are standardized, the importance of correct ratio (number) of countries between different division-aspects increases. As the sample may be a bit biased, the results may propose higher numbers of CO<sub>2</sub>-emissions per capita in the world than what they actually are.

## 2.2 Plotting the sample

Below is plotted three graphs. On the first row, we investigate our sample countries' CO<sub>2</sub>-emissions by country. Please note the y-axis difference between large and small emitters in the graphs. It is worth noting that the CO<sub>2</sub>-emissions development of China is very concerning as it has almost doubled its CO<sub>2</sub>-emissions during the last 15 years. In addition, India, Greece, Morocco, Peru and Mongolia has been showing a bit concerning trend during last decades.

On the second row, we plotted the sample countries' emission standardized with the population of the country. Thus, we obtained a "CO<sub>2</sub> per capita" -estimate for each country. This is the data that we used later in our models. Especially between 1950s and 70s, western countries play significant role as the big emitters. However, during 2000s, the situation has changed as western countries have systematically been able to lower their emissions per capita. At the same time, developing countries have been increasing their emissions and thus the situation has tied.



### 3 Model description

In this chapter, we will present our model structures and stan codes of our implementation of a non-hierarchical pooled model and a hierarchical model. Before this, we briefly introduce the mathematical structure behind the models.

### 3.1 Pooled model

A pooled model is one of the most straightforward model structure to understand. In the pooled model, all data points are used as one “pool” without considering groups or particular features different pieces of data could have. The whole dataset is used as a one, and modeling is done based on that collection of data. If we assume that priors of the mean and standard deviation follow standard normal distributions, we can present the mathematical structure of the pooled model in the form

$$\mu \sim N(0, 1) \tag{1}$$

$$\sigma \sim N(0, 1) \tag{2}$$

$$y_i \sim N(\mu, \sigma) \tag{3}$$

We used these standardized normal priors just for illustration purposes, and the correct choice of priors we utilized when modeling is presented in chapter 4. Respectively, the pooled model’s implementation with probabilistic programming language *Stan* is presented in chapter 5.

### 3.2 Hierarchical normal model

Unlike the previous model, the hierarchical model takes into account the possibility that some of the subgroups of the whole dataset have similar properties. Due to this observation, the hierarchical model presents a “hyper-prior” that is common to each group. For each group, its posterior distribution of mean is calculated using that hyper-prior, taking into account only all samples belonging to that group. This property can be illustrated in Figure 2. In Figure 2,  $\tau$  is a hyper-parameter and  $\theta_i$ s presents modeled parameter of each group.

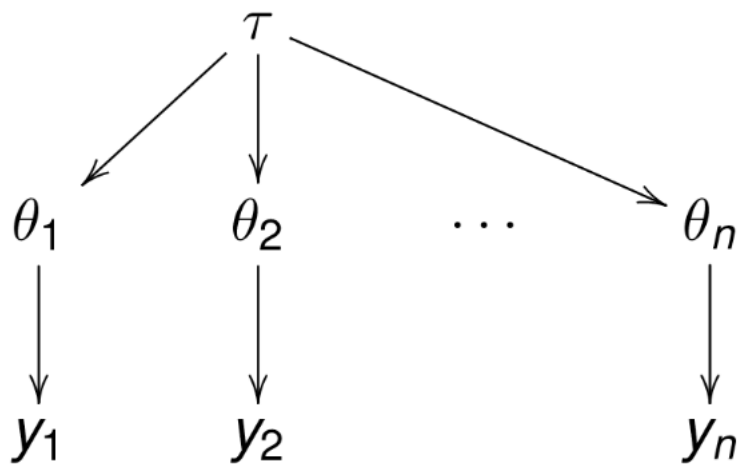


Figure 2: A hierarchical model –

We can summarize the hierarchical model mathematically as

$$\mu_0 \sim N(0, 1) \tag{4}$$

$$\sigma_0 \sim N(0, 1) \tag{5}$$

$$\mu_i \sim N(\mu_0, \sigma_0) \tag{6}$$

$$\sigma \sim N(0, 1) \tag{7}$$

$$y_{ji} \sim N(\mu_i, \sigma) \tag{8}$$

where  $\mu_0$  and  $\sigma_0$  are hyper-priors for mean and standard deviation. Again, we used these normal distributions just for illustration purposes. In addition, we assumed through the project that all the groups have a common variance ( $\sigma$  in [4]).

## 4. Priors

For our modeling, we needed to define hyper priors  $\mu_0$  and  $\sigma_0$ . In addition, common  $\sigma$  was defined for the hierarchical model's standard deviation between data points.

Our first goal was to define the hyper prior  $\mu_0$ . To aid this problem, we searched information in the internet about country-wise CO2-emissions per capita. The figure below illustrates the results that we found. The figure is taken from <https://www.economicshelp.org/blog/10296/economics/top-co2-polluters-highest-per-capita/> on the 1st of December, 2020.

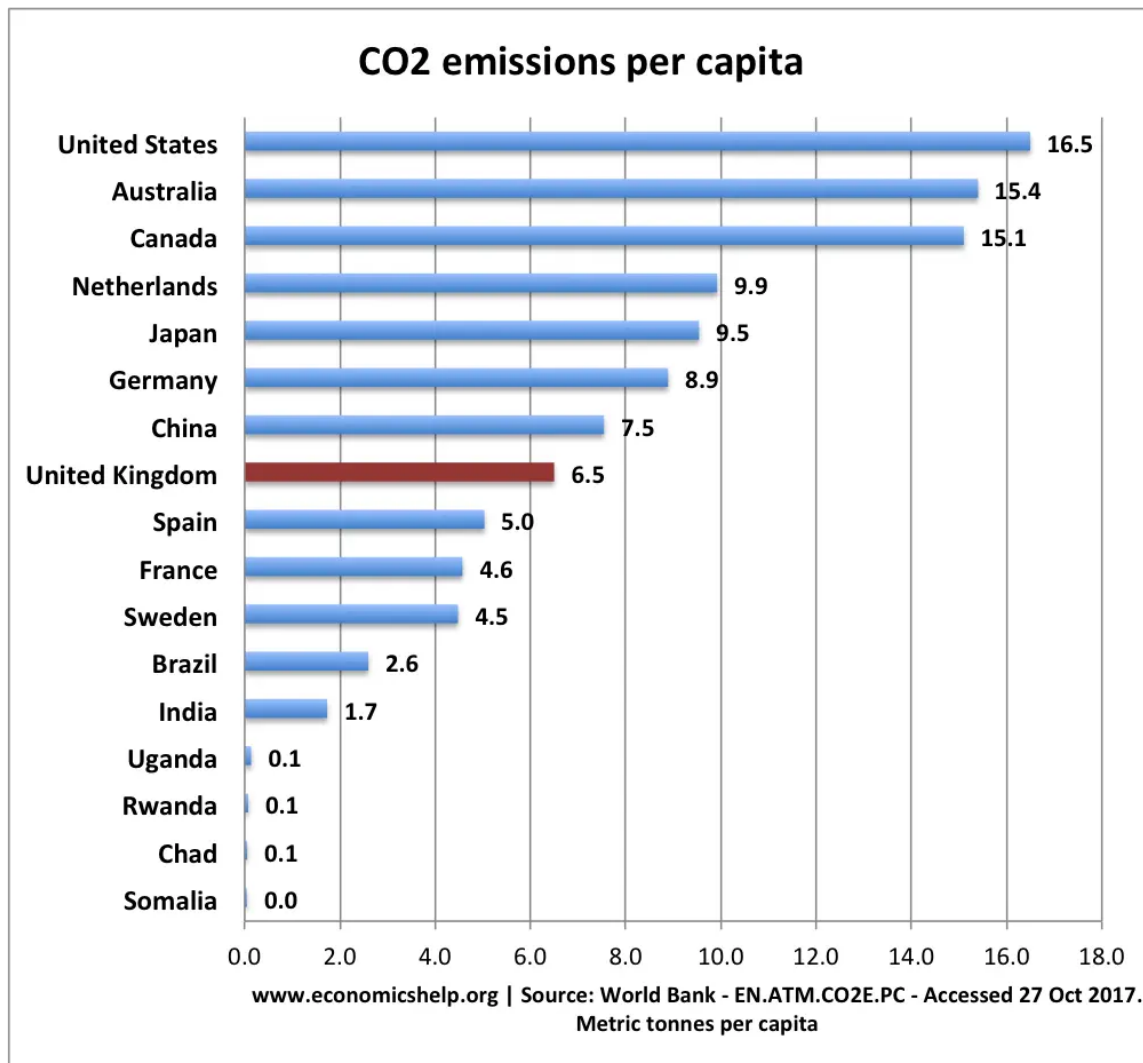


Figure 3: Selected countries CO2 emissions per capita

Source: <https://www.economicshelp.org/blog/10296/economics/top-co2-polluters-highest-per-capita/>  
Accessed December 1, 2020.

## 5. Stan

```
data { int <lower=0> N; // number of observations vector[N] y; // observations }
parameters { real mu; real<lower=0> sigma; }
model { mu ~ lognormal(0, 10); // priors from last week sigma ~ inv_chi_square(1); // priors from last week
// pooled model likelihood, common mu and sigma for all observations y ~ normal(mu, sigma); }
generated quantities { real ypred; vector[N] log_lik;
//predictive distribution for any machine ypred = normal_rng(mu, sigma);
for (i in 1:N){
  log_lik[i] = normal_lpdf(y[i] | mu, sigma);
}
}
```

### 5.2 Hierarchical model

Example code for hierarchical model

```
data { int<lower=0> N; // Number of observations int<lower=0> N_c; // Number of countries vector[N_c]
y[N]; // Observations }
parameters { vector[N_c] mu; // group means real hyper_mu; // prior mean real<lower=0> hyper_sigma;
// prior std constrained to be positive real<lower=0> sigma; // COMMON std constrained to be positive
}
model { hyper_mu ~ normal(0, 100); // weakly informative hyper-prior hyper_sigma ~ inv_chi_square(1);
// weakly informative hyper-prior
mu ~ normal(hyper_mu, hyper_sigma); // population prior with unknown parameters
sigma ~ inv_chi_square(1); // weakly informative prior for group (common) std

for (j in 1:N_c) {
  y[,j] ~ normal(mu[j], sigma); // likelihood
}
}

generated quantities { real y_pred; vector[N_c] log_lik[N];
y_pred = normal_rng(hyper_mu, sigma);

for (j in 1:N_c) {
  for (i in 1:N) {
    log_lik[i, j] = normal_lpdf(y[i,j] | mu[j], sigma);
  }
}
}

df_data <- data.frame(years=seq(1950,2018), data_co2_population)
df_plot <- melt(data = df_data, id.vars = "years", variable.name = "country")
vectedored_data_pop <- data.frame(df_plot[, 'value'])
N <- nrow(vectedored_data_pop)

num_of_iter <- 2000
num_of_warmup <- 1000
```



```

pool_data <- list(N = N,
                 y = vectored_data_pop[,1])

pool_model <- rstan::stan_model(file = "pooled_model_stan_without_loglik.stan")

pool_fit <- rstan::sampling(object = pool_model,
                           data = pool_data,
                           iter = num_of_iter,
                           warmup = num_of_warmup)

hier_data <- list(N = nrow(data_co2_population),
                 N_c = ncol(data_co2_population),
                 y = data_co2_population)

hier_model <- rstan::stan_model(file = "hier_model_stan_without_loglik.stan")

hier_fit <- rstan::sampling(object = hier_model,
                           data = hier_data,
                           iter = num_of_iter,
                           warmup = num_of_warmup,
                           refresh = 0,
                           seed=12345)

hier_fit
monitor(pool_fit)
monitor(hier_fit)

check_hmc_diagnostics(hier_fit)

pooled_plotters <- function() {
  pooled_df =data.frame(rstan::extract(pool_fit, permuted=T))

  #Histogram
  hist(pooled_df$mu,
       breaks = 100,
       xlim=c(0,22),
       xlab = "Mean of the quality measurements",
       col = "lightyellow",
       main="Posterior distribution of the mean of the sixth machine")
}

hierarchical_plotters <- function() {
  hierarchical_df =data.frame(rstan::extract(hier_fit, permuted=T))

  #MCMC Areas
  mcmc_areas_df <- hierarchical_df %>% select(starts_with('mu')) %>%
    setNames(colnames(data_co2_population))
  mcmc_areas(mcmc_areas_df) + xlab("Testttttttt")

  #Histograms of countries together
  m <- 19
  {
    plot(0,0,type="n",

```

```

    xlim=c(0,20), ylim=c(0,1100),
    xlab="x",ylab="freq",
    main="Histograms of each country separately, plotted together")
for(n in 1:m) {
  var_name <- paste("mu.",n, sep="")
  #hier_matrix[n,] <- unlist(hierarchical_df[var_name])
  plot(
    hist(unlist(hierarchical_df[var_name]), breaks = 12, plot=FALSE),
    col=alpha('blue', 0.25),
    add=T # Add to main plot
  )
}
}

#One histogram for whole data
one_hist_data <- unlist(hierarchical_df %>% select(starts_with('mu')))
plot(hist(one_hist_data, breaks = 100, xlim = c(0,22), ylim = c(0, 5000)),
      col = 'lightblue')
}

```

## References