

cs208 HW 2

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Note: I include the code I wrote directly in line for some questions, such as for 2 (a) and (b) which are purely implementation problems. For other questions, my code can be found in the Appendix and on [Github](#). I worked with Bhaven Patel and Lipika Ramaswamy for this assignment.

Question 1

(a) Both (i) and (iv) are $(\epsilon, 0)$ -differentially private. I demonstrate that below using two neighboring datasets, $x = [0, 0, \dots, 0]$ and $x' = [0, 0, \dots, 1]$.

(i)

$$\frac{P([\bar{x} + Z]_0^1 = r)}{P([x' + Z]_0^1 = r)}$$
$$\frac{\frac{n}{4} \exp(-\frac{n}{2}|r - \bar{x}|)}{\frac{n}{4} \exp(-\frac{n}{2}|r - x'|)}$$
$$\exp(\frac{n}{2}|r - \bar{x}'| - |r - \bar{x}|)$$

By the triangle inequality, this expression is less than or equal to

$$\exp(\frac{n}{2}|r - \bar{x}' - r + \bar{x}|) = \exp(\frac{n}{2}|\bar{x} - \bar{x}'|)$$

Since $|\bar{x} - \bar{x}'|$ is just the global sensitivity, we get that

$$\exp(\frac{n}{2}|\bar{x} - \bar{x}'|) = \exp(\frac{n}{2} \frac{1}{n}) = \exp(\frac{1}{2})$$

Thus, $M(x)$ is $(\epsilon, 0)$ -differentially private for $\epsilon \geq 0.5$.

(iv)

$$\frac{\exp(\frac{-n}{10}|y - \bar{x}|)}{\exp(\frac{-n}{10}|y - \bar{x}'|)} * \frac{\int_0^1 \exp(\frac{-n}{10}|z - \bar{x}|) dz}{\int_0^1 \exp(\frac{-n}{10}|z - \bar{x}'|) dz}$$

I'll evaluate this term by term. First, the left term:

$$\exp(\frac{n}{10}(-|y - \bar{x}| + |y - \bar{x}'|))$$

$$\exp(\frac{n}{10}(-|y| + |y - \frac{1}{n}|))$$

By the triangle inequality, this is less than or equal to

$$\exp(\frac{n}{10}(y - \frac{1}{n} - y)) = \exp(\frac{1}{10})$$

Now, for the right term:

$$\begin{aligned} & \frac{\int_0^1 \exp(\frac{-n}{10}|z - \frac{1}{n}|)dz}{\int_0^1 \exp(\frac{-nz}{10})dz} \\ & \frac{\int_0^{\frac{1}{n}} \exp(\frac{-n}{10}(z - \frac{1}{n}))dz + \int_{\frac{1}{n}}^1 \exp(\frac{-n}{10}(z - \frac{1}{n}))dz}{\int_0^{\frac{1}{n}} \exp(\frac{-nz}{10})dz + \int_{\frac{1}{n}}^1 \exp(\frac{-nz}{10})dz} \\ & \frac{\exp(\frac{1}{10}) \int_0^{\frac{1}{n}} \exp(\frac{-nz}{10})dz + \exp(\frac{1}{10}) \int_{\frac{1}{n}}^1 \exp(\frac{-nz}{10})dz}{\int_0^{\frac{1}{n}} \exp(\frac{-nz}{10})dz + \int_{\frac{1}{n}}^1 \exp(\frac{-nz}{10})dz} \end{aligned}$$

This reduces to $\exp(\frac{1}{10})$. Putting the two terms together, we have

$$\exp(\frac{1}{10}) * \exp(\frac{1}{10}) = \exp(\frac{1}{5})$$

Thus, this mechanism is $(\epsilon, 0)$ -differentially private for $\epsilon \geq 0.2$.

(b) Mechanisms (ii) and (iii) are not $(\epsilon, 0)$ -differentially private. Below I'll provide a counterexample that demonstrates this and find a minimum value of δ for which they are (ϵ, δ) -differentially private.

- (ii) Consider $x = [0, 0, \dots, 0]$ and $x' = [0, 0, \dots, 1]$. Now, $P(M(x) = -1) \geq 0$ while $P(M(x') = -1) = 0$. This violates $P(M(x) = -1) \leq \exp(\epsilon)P(M(x') = -1)$, so this mechanism is not $(\epsilon, 0)$ -differentially private. Now let's consider the minimum value of δ for which it is (ϵ, δ) -differentially private.

$$\delta \geq \max_{x, x'} \left[\int_y \max(P(M(x) = y) - \exp(\epsilon)P(M(x') = y), 0) \right]$$

Consider the worst-case scenario I defined above, where $x = [0, \dots, 0]$ and $x' = [0, \dots, 0, 1]$. The expression inside the integral will only be ≥ 0 for $y \in [-\infty, -1 + \frac{1}{n}]$ because $P(M(x') = y) = 0$ here.

$$\begin{aligned} & \max \left[\int_{-\infty}^{\infty} P(M(x) = y) - \exp(\epsilon)P(M(x') = y), 0 \right] \\ & \int_{-\infty}^{-1 + \frac{1}{n}} P(M(x) = y) \\ & \int_{-1}^{-1 + \frac{1}{n}} P(\bar{x} + Z = y) \end{aligned}$$

Since $\bar{x} = 0$,

$$\int_{-\infty}^{-1 + \frac{1}{n}} P(Z = y)$$

$$\int_{-\infty}^{-1+\frac{1}{n}} \frac{n}{4} \exp\left(\frac{-n|y|}{2}\right)$$

$$\int_{-\infty}^{-1+\frac{1}{n}} \frac{n}{4} \exp\left(\frac{ny}{2}\right)$$

$$\frac{n}{4} \int_{-\infty}^{-1+\frac{1}{n}} \exp\left(\frac{ny}{2}\right)$$

After integrating we are left with

$$\frac{1}{2} [\exp\left(\frac{ny}{2}\right)]_{-\infty}^{-1+\frac{1}{n}}$$

$$\frac{1}{2} [\exp\left(\frac{-n+1}{2}\right) - \exp\left(\frac{-n\infty}{2}\right)]$$

Since the second expression in the parenthesis goes to 0, we have

$$\frac{1}{2} \exp\left(\frac{-n+1}{2}\right)$$

Thus, this mechanism is (ϵ, δ) -differentially private for $\delta \geq \frac{1}{2} \exp\left(\frac{-n+1}{2}\right)$.

- (iii) Consider $x = [0, 0, \dots, 1]$ and $x' = [0, 0, \dots, 0]$. Now, $P(M(x) = 1) = \frac{1}{n}$ while $P(M(x') = 1) = 0$. This clearly violates $P(M(x) = 1) \leq \exp(\epsilon) P(M(x') = 1)$, so this mechanism is not $(\epsilon, 0)$ -differentially private. Now let's consider the minimum value of δ for which it is (ϵ, δ) -differentially private.

$$\delta \geq \max_{x, x'} [\Sigma_y \max(P(M(x) = y) - \exp(\epsilon) P(M(x') = y), 0)]$$

$$\begin{aligned} & \Sigma_{y \in [0,1]} \max[(P(M(x) = y) - \exp(\epsilon) P(M(x') = y), 0] \\ & \max[P(M(x) = 1) - \exp(\epsilon) P(M(x') = 1), 0] + \max[P(M(x) = 0) - \exp(\epsilon) P(M(x') = 0), 0] \end{aligned}$$

$$\max\left[\frac{1}{n} - \exp(\epsilon) * 0, 0\right] + \max\left[1 - \frac{1}{n} - \exp(\epsilon) * 1, 0\right] = \frac{1}{n} + 0 = \frac{1}{n}$$

Thus, this mechanism is (ϵ, δ) -differentially-private for $\delta \geq \frac{1}{n}$.

(c)

- (i) For this mechanism, the minimum value of ϵ depends on the bounds. Since the data were bounded on $[0,1]$, the minimum value of ϵ is 0.5, but if $/epsilon$ were on $[a,b]$, then the minimum value is $\frac{b-a}{n}$. From here the value of ϵ used for the mechanism can be tuned, i.e., made larger based on an evaluation of the privacy-utility tradeoff.
- (ii) •
- (iii) •
- (iv) Similarly to (i), this minimum value of ϵ depends on the bounds of the data. Here, the minimum value would be $\frac{b-a}{5}$.

(d) I would consider mechanism (i) to be the best for releasing a mean. For one, -we understand laplace mechanism well -clamping to protect outliers?

Question 2

(a)

```
poissonDGP <- function(n){ return(rpois(n, lambda=10)) }
```

(b) I use the first mechanism from Question 1 to answer this question. From here on out I'll refer to it as the clamped Laplace mean release mechanism.

```
sgn <- function(x) {      # function borrowed from class
  return(ifelse(x < 0, -1, 1))
}
```

```
rlap = function(mu=0, b=1, size=1) {      # function borrowed from class
  p <- runif(size) - 0.5
  draws <- mu - b * sgn(p) * log(1 - 2 * abs(p))
  return(draws)
}
```

```
clip <- function(x, lower, upper){      # function borrowed from class
  x.clipped <- x
  x.clipped[x.clipped<lower] <- lower
  x.clipped[x.clipped>upper] <- upper
  return(x.clipped)
}
```

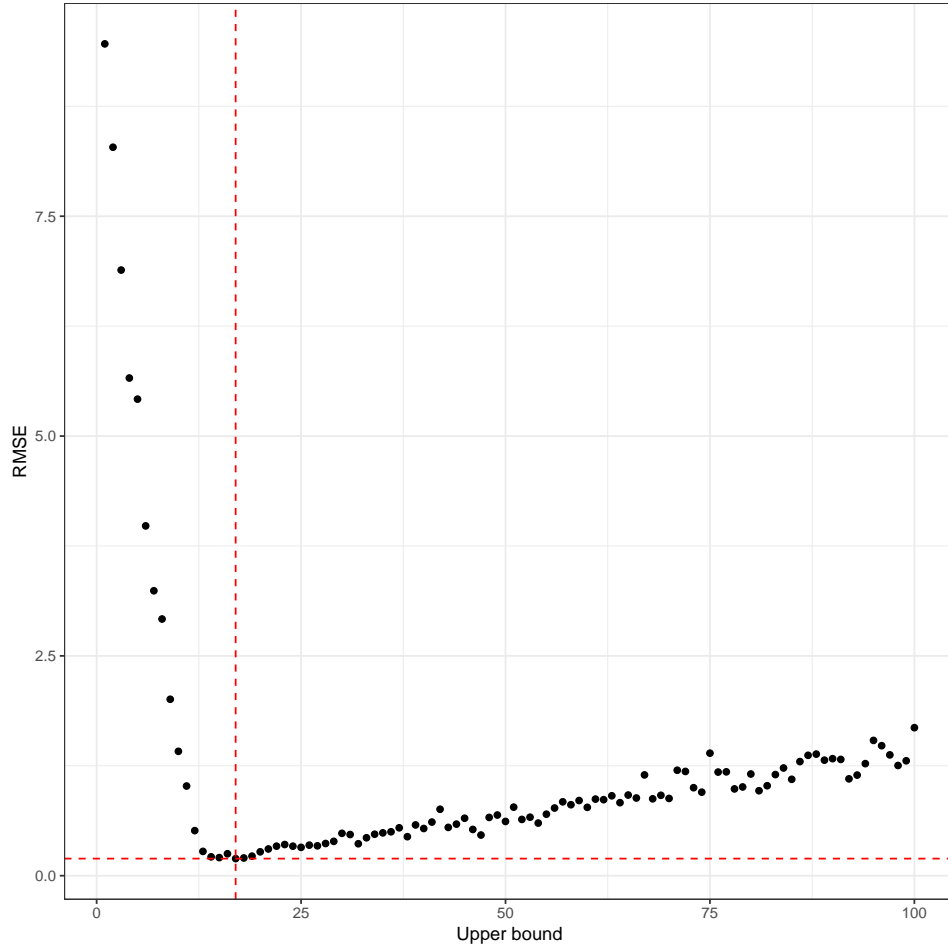
```
laplaceClampMeanRelease <- function(x, epsilon, a=0, b=1){
  n <- length(x)
  sensitivity <- (a - b)/n
  scale <- sensitivity / epsilon

  x.clipped <- clip(x, a, b)
  clipped.mean <- mean(x.clipped)
  noisy.mean <- clipped.mean + rlap(mu=0, b=scale, size=1)
  release.mean <- clip(noisy.mean, a, b)
  true.mean <- mean(x)

  return(list(release=release.mean, true=true.mean))
}
```

(c) I run 100 Monte Carlo simulations per upper bound value. In each simulation I draw 200 samples randomly drawn from a Poisson distribution and compute a noisy mean using the clamped Laplace mean mechanism. Then, I compare the root mean squared error between the true and noisy means for each upper bound value and plot these for every upper bound value. This is shown in the plot below.

Based on my analysis, the optimal value for the upper bound b - i.e., the one that minimizes RMSE - is 19. I will use that value for the remainder of this problem and for Question 3.



(d) This bootstrap approach is not a good mechanism to preserve privacy because we may leak information about outliers. Consider a dataset with a lot of observations between 0 and 10, a few less between 20 and 30, and a handful of observations between 35 and 40. When we bootstrap (resample without replacement) there is a possibility, albeit a small one, that our bootstrapped sample will overrepresent these outliers. If we get one of those bootstrapped samples when we evaluate an upper bound of 20, it will appear that 20 is a poor upper bound to choose from a utility perspective, but if we get one of those bootstrapped samples when the upper bound in question is 40, then this cutoff will look like a good choice. Thus, the upper bound value may actually leak information about outliers in the dataset.

(e) -propose new mechanism -local sensitivity \rightarrow lower sensitivity in general, means that we draw less noisy means -BOOM -or a different statistic, different mechanism -see section notes

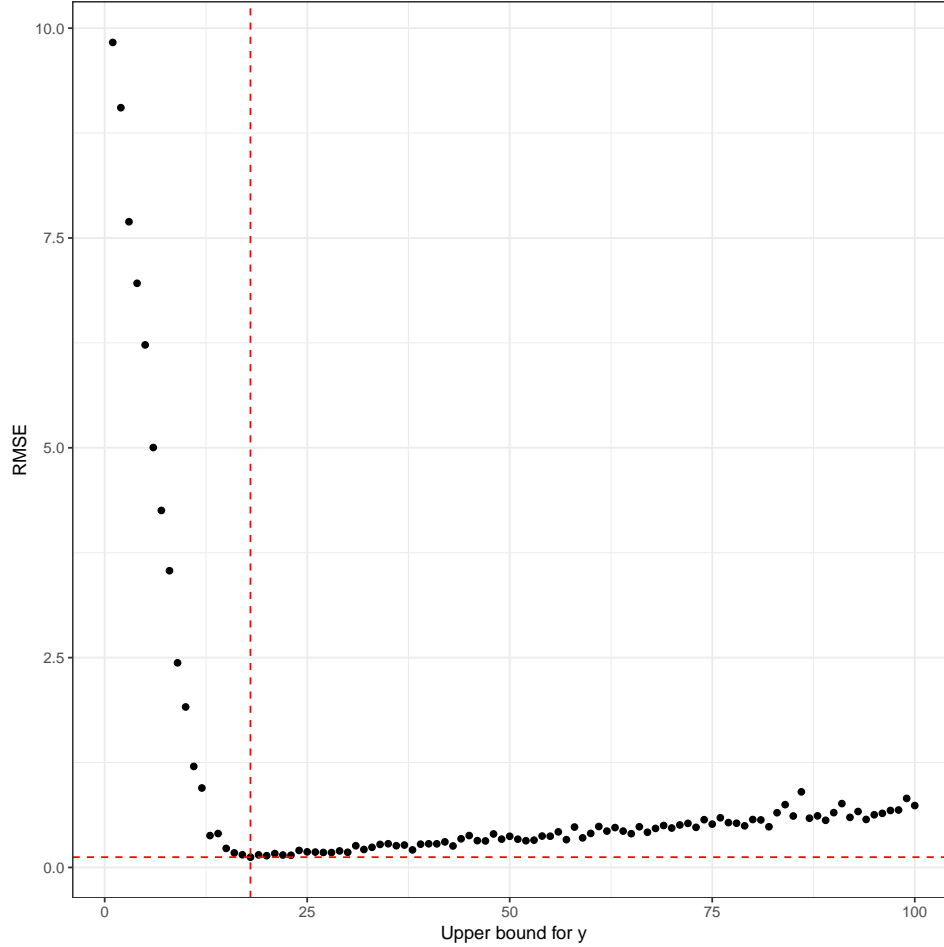
Question 3

(a) I break up the regression coefficient and intercept release mechanism into four differentially private subroutines. We showed in class that simply computing a differentially private version of the coefficient directly would be useless because the global sensitivity approaches ∞ .

First, I know that I can write the coefficient of the regression as $\frac{S_{xy}}{S_{xx}}$. Thus, computing a differentially private version of the coefficient requires two subroutines: one for S_{xy} and another for S_{xx} . Next, to calculate the intercept, I can use the formula $\hat{\alpha} = \bar{y} - \hat{\beta}\bar{x}$. Since our $\hat{\beta}$ is now differentially private, we can use the post-processing property to use it in this calculation with no extra privacy loss. However, we also need to compute a differentially private version of \bar{x} , and \bar{y} . To do this, I'll use the clamped Laplace mean release

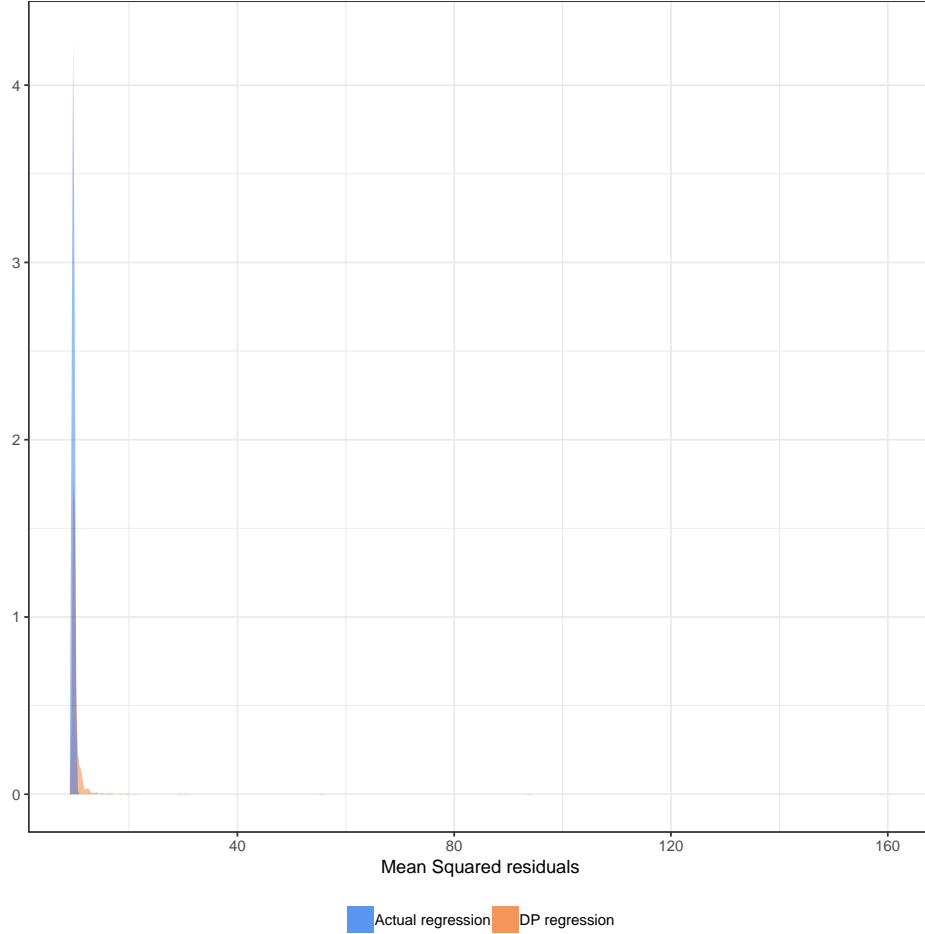
mechanism.

Furthermore, I will clamp both x and y as a part of this regression release mechanism. In Question 2 I found 19 to be a good upper bound value for x , and I'll continue to use that here. To get a good upper bound for y , I repeated the same process that I used for x but will also feed the data randomly drawn from a Poisson distribution into the noisy linear function and find the best upper bound for y that reduces the RMSE between the actual and noisy y values from the clamped Laplace mean release mechanism. As shown in the plot below, I find that 17 is the optimal upper bound for y .



(b) I generate 1,000 different datasets, perform my regression release on each dataset (splitting my budget evenly across the four subroutines), and then compare the performance of my noisy regression parameters to the performance of the true regression parameters by computing the mean-squared residuals for the original data. Here I consider the true regression to be the one computed on the clipped data so that my comparison focuses specifically on the effect of the added noise and not on the clipping procedure.

The plot below shows the distributions of the mean-squared residuals for the private and non-private regressions (the plot shows kernel density estimates). In general, the distribution of the mean-squared residuals for the private regression has much longer tail, implying that this approach has a larger expected error and could potentially lead to an enormous, utility-diminishing amount of error.



(c) Now I run a grid search to attempt to identify a better partitioning of the privacy budget that leads to lower error. To set up my grid, I consider weights for each statistic ranging from 0.1 to 0.9 by steps of 0.1. I then apply a softmax function to ensure that each partition of epsilon adds up to 1 - the total privacy budget for this problem. In total, I consider 6,561 partitions, although some are redundant. For each partition in my grid, I run 10 simulations in which I compute the differentially private regression and take the average of the mean squared residuals across these 10 simulations. I use the same underlying dataset across all the simulations so that the only variability comes from the Laplace noise in my mechanism.

I do not find that any of these partitions substantially improve the utility provided by the equal partition. I find that the minimum mean squared residual is 9.56 and comes when the weights for the weights for S_{xx} , S_{xy} , \bar{x} , and \bar{y} are 0.20, 0.18, 0.28, and 0.34, respectively. In comparison the minimum mean squared residual for the equal partition was 9.20 and the average was 10.46.

Question 4

First we can write out the expected fraction of bits reconstructed as the sum of the probabilities of reconstructing each bit.

$$\frac{P[A(M(X_i, X_{-i})) = X_i]}{n}$$

By a union bound, this is less than or equal to

$$\frac{\sum_{i=1}^n P[A(M(X_i, X_{-i})) = X_i]}{n}$$

By the definition of (ϵ, δ) -differential privacy, we have

$$\frac{\sum_{i=1}^n P[A(M(X_i, X_{-i})) = X_i]}{n} \leq \frac{\sum_{i=1}^n \exp(\epsilon) P[A(M(X_i, X_{-i})) = X_i] + \delta}{n}$$

For now, I'll drop the denominator since it is not affecting our evaluation of the numerator. Considering the post-processing property of a (ϵ, δ) -differentially private mechanism, we can replace each X_i with either 0 or 1. Let $P(X_i = 0) = p$ and $P(X_i = 1) = p - 1$. The left hand term above is now equal to

$$\max(\sum_{i=1}^n \exp(\epsilon) P[A(M(0, X_{-i})) = X_i], \sum_{i=1}^n \exp(\epsilon) P[A(M(1, X_{-i})) = X_i]) + \delta$$

$$\max(\sum_{i=1}^n \exp(\epsilon) p, \sum_{i=1}^n \exp(\epsilon) (1 - p)) + \delta$$

$$\max(p, (1 - p)) * \exp(\epsilon) * n + \delta$$

The n in this expression goes away when we divide by the denominator n that I omitted for these calculations. Further, we can rewrite $\delta^* = \delta/n$. Thus, we have that

$$E \left[\frac{\#i \in [n] : A(M(x))_i = X_i}{n} \right] \leq \max(p, (1 - p)) * \exp(\epsilon) + \delta^*$$

Question 5

I will be working with Bhaven Patel for the final project.

Appendix

I put the code for all of my analyses here. You can also find it on [Github](#).

Question 2

```
# c
rmse <- function(pred, true){ return(sqrt(mean((pred-true)^2))) }

n = 200
epsilon = 0.5
b_vals = seq(from=1, to=100, by=1)
n_sims <- 100

results <- matrix(NA, nrow=(length(b_vals)*n_sims), ncol=4)
i = 1
for (b in b_vals){
  dat <- poissonDGP(n)
  for (j in 1:n_sims){
    DPrelease <- laplaceClampMeanRelease(dat, epsilon, a=0, b=b)
```



```

      results[i,1] <- b
      results[i,2] <- j
      results[i,3] <- DPrelease$release
      results[i,4] <- DPrelease$true
      i = i + 1
    }
  }
results_df <- data.frame(results)
names(results_df) <- c("b", "sim", "release", "true")
avg_results_df <- results_df %>% group_by(b) %>% summarise(rmse = rmse(release, true))

q2_plot <- ggplot(data=avg_results_df, aes(x=b, y=rmse)) +
  geom_point() + geom_hline(yintercept = min(avg_results_df$rmse), col="red", lty=2) +
  geom_vline(xintercept = avg_results_df[which.min(avg_results_df$rmse), ]$b, col="red", lty=2) +
  labs(x="Upper bound", y="RMSE") + theme_bw()

```

Question 3

```

# a
poissonDGP <- function(n){ return(rpois(n, lambda=10)) }
noisyLinearDGP <- function(x, n, alpha, beta, mu=0, sd=1) { return(beta*x + alpha + rnorm(n, mu, sd)) }

sgn <- function(x) {      # function borrowed from class
  return(ifelse(x < 0, -1, 1))
}

rlap = function(mu=0, b=1, size=1) {      # function borrowed from class
  p <- runif(size) - 0.5
  draws <- mu - b * sgn(p) * log(1 - 2 * abs(p))
  return(draws)
}

clip <- function(x, lower, upper){      # function borrowed from class
  x.clipped <- x
  x.clipped[x.clipped<lower] <- lower
  x.clipped[x.clipped>upper] <- upper
  return(x.clipped)
}

rmse <- function(pred, true){ return(sqrt(mean((pred-true)^2))) }

laplaceClampMeanRelease <- function(x, epsilon, a=0, b=1){      # from q2
  n <- length(x)
  sensitivity <- (a - b)/n
  scale <- sensitivity / epsilon

  x.clipped <- clip(x, a, b)
  clipped.mean <- mean(x.clipped)
  noisy.mean <- clipped.mean + rlap(mu=0, b=scale, size=1)
  release.mean <- clip(noisy.mean, a, b)
  true.mean <- mean(x)

  return(list(release=release.mean, true=true.mean))
}

```

```

}

regressionRelease <- function(y, x, ylower=0, yupper=17, xlower=0, xupper=19, eplpsilon, partition){
  x <- clip(x, xlower, xupper)
  y <- clip(y, ylower, yupper)

  n <- length(x)
  sens.Sxy <- ((xupper-xlower)*(yupper-ylower))
  sens.Sxx <- ((xupper-xlower)^2)

  scale.Sxy <- sens.Sxy / (epsilon*partition$Sxy)
  scale.Sxx <- sens.Sxx / (epsilon*partition$Sxx)

  true.beta <- sum((x - mean(x))*(y - mean(y))) / sum((x - mean(x))^2)
  true.alpha <- mean(y) - true.beta*mean(x)

  release.Sxy <- sum((x - mean(x))*(y - mean(y))) + rlap(mu=0, b=scale.Sxy, size=1)
  release.Sxx <- sum((x - mean(x))^2) + rlap(mu=0, b=scale.Sxx, size=1)
  release.beta <- release.Sxy/release.Sxx

  release.x.bar <- laplaceClampMeanRelease(x, epsilon*partition$x.bar, a=xlower, b=xupper)$release
  release.y.bar <- laplaceClampMeanRelease(y, epsilon*partition$y.bar, a=ylower, b=yupper)$release
  release.alpha <- release.y.bar - release.beta*release.x.bar

  release.mean.sq.residuals <- mean((y - release.beta*x - release.alpha)^2)
  true.mean.sq.residuals <- mean((y - true.beta*x - true.alpha)^2)

  return(list(release.beta=release.beta,
             release.alpha=release.alpha,
             true.beta=true.beta,
             true.alpha=true.alpha,
             release.mean.sq.residuals=release.mean.sq.residuals,
             true.mean.sq.residuals=true.mean.sq.residuals))
}

# get optimal upper bound for y
n = 200
epsilon = 1
b_vals = seq(from=1, to=100, by=1)
n_sims <- 100

results_y <- matrix(NA, nrow=(length(b_vals)*n_sims), ncol=4)
i = 1
for (b in b_vals){
  dat <- poissonDGP(n)
  y <- noisyLinearDGP(dat, n, alpha=1, beta=1, mu=0, sd=1)
  for (j in 1:n_sims){
    DPrelease <- laplaceClampMeanRelease(y, epsilon, a=0, b=b)
    results_y[i,1] <- b
    results_y[i,2] <- j
    results_y[i,3] <- DPrelease$release
    results_y[i,4] <- DPrelease$true
    i = i + 1
  }
}

```

```

}
results_y_df <- data.frame(results_y)
names(results_y_df) <- c("b", "sim", "release", "true")
avg_results_y_df <- results_y_df %>% group_by(b) %>% summarise(rmse = rmse(release, true))

q3_plot1 <- ggplot(data=avg_results_y_df, aes(x=b, y=rmse)) +
  geom_point() + geom_hline(yintercept = min(avg_results_y_df$rmse), col="red", lty=2) +
  geom_vline(xintercept = avg_results_y_df[which.min(avg_results_y_df$rmse), ]$b, col="red", lty=2) +
  labs(x="Upper bound for y", y="RMSE") + theme_bw()
pdf("plots/q3_plot1.pdf", width=8, height=8)
q3_plot1
dev.off()

# b
equal_partition <- list(Sxy=0.25, Sxx=0.25, x.bar=0.25, y.bar=0.25)
n = 1000
alpha = beta = eplison = sd = 1
n_sims = 1000

results_reg <- matrix(NA, nrow=n_sims, ncol=6)
for (i in 1:n_sims){
  x <- poissonDGP(n)
  y <- noisyLinearDGP(dat, n, alpha=1, beta=1, mu=0, sd=1)
  DPrelease <- regressionRelease(y, x, ylower=0, yupper=17, xlower=0, xupper=19, eplison, equal_partit
  results_reg[i,1] <- DPrelease$release.beta
  results_reg[i,2] <- DPrelease$release.alpha
  results_reg[i,3] <- DPrelease$true.beta
  results_reg[i,4] <- DPrelease$true.alpha
  results_reg[i,5] <- DPrelease$release.mean.sq.residuals
  results_reg[i,6] <- DPrelease$true.mean.sq.residuals
}
results_reg_df <- data.frame(results_reg)
names(results_reg_df) <- c("release.beta", "release.alpha", "true.beta",
  "true.alpha", "release.mean.sq.residuals", "true.mean.sq.residuals")

semi.blue <- rgb(0,90,239,50,maxColorValue=255)
semi.red <- rgb(239,90,0,200,maxColorValue=255)
q3_plot2 <- ggplot(data=results_reg_df) +
  geom_density(aes(x=release.mean.sq.residuals, fill="DP regression"), alpha=0.4, colour=NA) +
  geom_density(aes(x=true.mean.sq.residuals, fill="Actual regression"), alpha=0.4, colour=NA) +
  labs(x="Mean Squared residuals", y="") + theme_bw() +
  scale_fill_manual(values=c(semi.blue, semi.red)) +
  theme(legend.position="bottom", legend.title=element_blank())

# c
# set up grid
softmax <- function(x) {
  return(exp(x)/sum(exp(x)))
}
weights = seq(0.1,0.9,by=0.1)
weights_list <- expand.grid(Sxx=weights, Sxy=weights, x.bar=weights, y.bar=weights)
partition_list<- t(apply(weights_list, 1, softmax))
partition_list_df <- data.frame(partition_list)
names(partition_list_df) <- c("Sxx", "Sxy", "x.bar", "y.bar")

```

```

# cross validation
# use same x and y
set.seed(25)
x <- poissonDGP(1000)
y <- noisyLinearDGP(dat, 1000, alpha=1, beta=1, mu=0, sd=1)
# iterate through different partitions
results_cv <- matrix(NA, nrow=nrow(partition_list), ncol=5)
for (i in 1:nrow(partition_list)){
  DPrelease <- regressionRelease(y, x, ylower=0, yupper=17, xlower=0, xupper=19, epsilon, partition_list)
  results_cv[i,1] <- partition_list_df[i,]$Sxx
  results_cv[i,2] <- partition_list_df[i,]$Sxy
  results_cv[i,3] <- partition_list_df[i,]$x.bar
  results_cv[i,4] <- partition_list_df[i,]$y.bar
  results_cv[i,5] <- DPrelease$release.mean.sq.residuals
}
results_cv_df <- data.frame(results_cv)
names(results_cv_df) <- c("Sxx", "Sxy", "x.bar", "y.bar", "release.mean.sq.residuals")

print(results_cv_df[which.min(results_cv_df$release.mean.sq.residuals),])
print(min(results_reg_df$release.mean.sq.residuals))
print(mean(results_reg_df$release.mean.sq.residuals))

```