dapper: Data Augmentation for Private Posterior Estimation in R

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Abstract This paper serves as a reference and introduction on using the dapper R package. The goal of this package is to provide some tools for exploring the impact of different privacy regimes on a Bayesian analysis. A strength of this framework is the ability to target the exact posterior in settings where the likelihood is too complex to analytically express.

1 Introduction

Differential privacy provides a rigorous framework for protecting confidential information. In this framework, privacy is obtained through the injection of random noise in the data analysis workflow. It has served as the theoretical foundation for recent advances in privacy technology. Several high profile examples include Apple (), Google (), Microsoft (), and the U.S. Census Bureau ().

There are several approaches for constructing or modifying a data analysis workflow to provide privacy guarantees, which loosely fall into three categories: direct, query, and dissemination. The approaches mainly differ in where in the workflow privacy noise is injected. In the dissemination setting, data needs to be released to the public, and privacy is maintained by first perturbing the data with random noise before being released. There is strong interest in further developing this approach because data curators are often interested in releasing data to the public without knowledge of how their data will be used or analyzed. In this scenario, it can be impossible to apply the direct or query method because they require knowing before hand how the data will be analyzed. For a recent example of a high profile application of the dissemination approach see the the U.S. Census Bureau's TopDown algorithm (Bureau 2023). (Ji, Lipton, and Elkan 2014) provide a nice survey of the direct and query based methods applied to common machine learning algorithms.

Correctly carrying out statistical inference in the dissemination setting requires adjusting statistical workflows to account for privacy noise. As an example, one instance of implementing the dissemination approach for tabular data involves directly adding independent, random error to each cell. For regression models, this corresponds to having data with measurement errors in the covariates. This, unfortunately, violates the assumptions of most statistical models. In the presence of such errors, standard estimators can exhibit significant bias (Gong 2022). Therefore, fitting models without accounting for the added privacy noise can lead to incorrect inference.

Unfortunately, making the necessary adjustments poses formidable mathematical challenges, even for seemingly simple models like linear regression. The difficulty can be seen from considering the marginal likelihood that results from correctly accounting for the injected privacy noise. This function is often analytically intractable and as a result, it is difficult or impossible to apply common maximum likelihood methods to derive estimators. In particular, the marginal likelihood can involve a complex integral where it is not possible to even evaluate the likelihood at a point. This makes gradient based methods, like the Newton-Raphson method, impractical. And approximating the likelihood can be computationally unfeasible since the integral is also of high dimension. Few tools are available to researchers to address these issues, and their absence is a serious barrier to the wider adoption of dissemination methods.

The DAPPER package provides a tool for conducting valid statistical inference in the presence of privacy noise. It implements the Bayesian framework proposed in (Ju et al. 2022). This framework describes how to modify an existing Bayesian model to account for privacy noise. DAPPER serves as user-friendly interface for implementing the framework in R. It allows the user to specify a sampler from an existing Bayesian model and automatically constructing a valid posterior sampler that accounts for the added privacy noise.

In the previously mentioned case where privacy noise is directly added to each cell of a data table, there exist readily available methods from measurement error literature for making adjustments. For textbook length treatments on the topic see (Yi 2017; Carroll et al. 2006). Work in this area mostly focuses on methods which do not require fully specifying the measurement error model, since this is often assumed unknown. However, in differential privacy, the measurement error model is exactly known. This difference, makes feasible some ideas which the measurement error community has not previously considered (Smith 2011; Karwa, Kifer, and Slavković 2015).

The rest of this article is organized as follows. Section 2 covers the necessary background to understand the mathematical notation and ideas used throughout the paper. Section 3 goes over the

main algorithm without going into mathematical detail, for specifics see (Ju et al. 2022). Section 4 provides an overview of the dapper package and discusses important implementation details. Section 5 contains two example of how one might use the package to analyze the impact of adding noise for privacy. The first example goes over a typical odds ratio analysis for a 2×2 table and the second example covers a linear regression model.

2 Background

Let $x = (x_1, \dots, x_n) \in \mathcal{X}^n$ represent a confidential database containing n records. Usually, the goal of collecting data is to learn some characteristic about the underlying population. To accomplish this task, a common approach is to assume the population is represented by some statistical model $f(\cdot \mid \theta)$. It is often the case that some function of θ has relevant meaning to the scientific question at hand. In this setting, learning characteristics of a population reduces to learning about θ .

In the Bayesian statistical framework, learning about θ is accomplished by drawing samples from the posterior $p(\theta \mid x) \propto f(x \mid \theta)p(\theta)$. For large data sets, it is common to work with a summary statistic s=s(x) that has much smaller dimension than the original data. Doing so can greatly simplify calculations. In general, there can be information loss with using summary statistics, but for models with a sufficient statistic, there is no loss. Curators of large databases often use summary statistics to publicize data since it allows them to efficiently communicate information contained in large data sets. For this reason, summary statistics are a natural target for dissemination based privacy approaches.

Differential Privacy

While a summary statistic can already partially anonymize data, it is still possible to deduce information about an individual entry depending on the distribution of x. Differential privacy solves this problem by taking a summary statistic s, and adding noise to it to produce a noisy summary statistic s_{dp} . While this method is not new [insert citation on swapping and randomized responses], differential privacy provides a rigorous framework for specify where and how much noise to add.

We now describe the ϵ -DP privacy framework in more detail. For the noisy summary statistic, we write $s_{dp} \sim \eta(\cdot \mid x)$. Here, η is a known noise infusion process designed to meet a certain property: The privacy mechanism η is said to be ϵ -differentially private (Dwork and Roth 2013) if for all values of s_{dp} , and all "neighboring" databases $(x, x') \in \mathcal{X}^n \times \mathcal{X}^n$ differing by one record (denoted by d(x, x')), the probability ratio is bounded:

$$\frac{\eta(s_{dp} \mid x)}{\eta(s_{dp} \mid x')} \le \exp(\epsilon), \quad \epsilon > 0.$$

The parameter ϵ is called the privacy loss budget, and controls how strong the privacy guarantee is. Larger values of ϵ correspond to weaker privacy guarantees which in turn means less noise being added.

The differential privacy framework is used to create and verify privacy mechanisms. One such mechanism is the Laplace mechanism. It works by taking a deterministic query $s:X\mapsto\mathbb{R}^m$ and constructing the privatized query $s_{dp}:=s(x)+u$ where u is an m-dimensional vector of iid Laplace noise. Using the ratio bound, if we draw each $u_i\sim Lap(\Delta(s)/\epsilon)$, we can show s_dp is ϵ -differentially private. Here $\Delta(s)$ is the global sensitivity of s. Example 2, will cover an application of the Laplace mechanism to linear regression.

Data Augmentation

The idea behind data augmentation is to run a Gibbs sampler on a coupling of two random variables where one of the marginals is the target distribution. Suppose we wish to sample from a density f_A which is difficult. The data augmentation method instead considers sampling from a joint distribution f(a,b). Since we are ultimately interested in samples from f_A , the joint distribution should be chosen so that (i) the marginal distribution with respects to a is f_A and (ii) $f(a \mid b)$ and $f(b \mid a)$ are easy to sample from. The choice f is not unique and can require some foresight.

3 Methodology

Given data s_{dp} , the goal of Bayesian inference is to sample from the posterior distribution $p(\theta \mid s_{dp})$. Since the observed likelihood, $p(s_{dp} \mid \theta)$, often has no simple closed form expression, most standard

sampling schemes do not apply. To conduct privacy-aware Bayesian inference, the dapper package implements the data augmentation algorithm which allows us to sample from $p(\theta \mid s_{dn})$ without needing to specify $p(s_{dp} \mid \theta)$.

The algorithm considers the joint distribution $p(\theta, x \mid s_{dp})$ and alternates sampling from the two distributions

- $p(\theta \mid x, s_{dp})$ $p(x \mid \theta, s_{dp})$

Since s_{dp} is derived from x, we have $p(\theta \mid x, s_{dp}) = p(\theta \mid x)$ which is just the usual posterior distribution given the confidential data x. The dapper package assumes the user has access to a sampler for $p(\theta \mid x)$. This can come from any R package such as fmcmc. For the second distribution, $p(x \mid \theta, s_{dp})$, may only be known up to a constant. The dapper package samples from this distribution by running a Gibbs-like sampler. Each of the n components of x is individually updated. However unlike the standard Gibbs sampler, each component is updated using a Metropolis-Hasting algorithm. This method is sometimes called the Metropolis within Gibbs sampler (Robert and Casella 2004).

In some cases, sampling from $p(x \mid \theta, s_{dp})$ can be made more efficient when the privacy mechanism can be written as a function of s_{dp} and a sum consisting of contribution from each individual record. More precisely, we say the privacy mechanism satisfies the record additivity property if

$$\eta(s_{dp} \mid x) = g(s_{dp}, \sum_{i=1}^{n} t_i(x_i, s_{dp}))$$

for some known and tractable functions g, t_1, \ldots, t_n . The sample mean is a example of a summary statistic satisfying record additivity where $t_i(x_i, s_{dp}) = x_i$.

The algorithm is in the following pseudo code:

- 1. Sample θ^{t+1} from $p(\cdot \mid x^{(t)})$.
- 2. Sample from $p(x \mid \theta, s_{dp})$ using a three step process
 - Propose $x_i^* \sim f(\cdot \mid \theta)$.
 - If s satisfies the record additive property then update $s(x^*, s_{dp}) = t(x, s_{dp}) t_i(x_i, s_{dp}) + t_i(x_i, s_{dp})$
 - Accept the proposed state with probability $\alpha(x_i^* \mid x_i, x_{-i}, \theta)$ given by:

$$\alpha(x_i^* \mid x_i, x_{-i}, \theta) = \min \left\{ 1, \frac{\eta(s_{dp} \mid x_i^*, x_{-i})}{\eta(s_{dp} \mid x_i, x_{-i})} \right\}.$$

The Structure of dapper

The package is structured around the two functions dapper_sample and new_privacy. The first function is used to draw samples from the posterior. The second function is used to create a privacy data model object which the dapper_sample function requires as input. The purpose of the data model object is to collect all the components specific to the data augmentation algorithm into one bundle. This way, the other arguments into dapper_sample pertain only to sampling parameters such as the number of iterations.

Since the input to these functions are R functions, there is a great deal of freedom in implementation. The next two sections describe in detail the inputs into these functions and highlight some considerations that should be taken into account in order avoid slow or unexpected behavior.

Before delving into the specifics of each component, it is necessary to clearly define how the confidential data is represented. Internally, the confidential database is encoded as a 2D matrix. There can be multiple ways of doing this and is often the case when dealing with likelihood functions that has a sufficient statistic. For example, if our data consist of 100 responses from a two question, yes/no, survey. Then we can either encode the data as a 2×2 matrix, or a 100×2 matrix. Both are mathematically equivalent, but the 2×2 matrix will be much more memory efficient. In general, the representation that uses the least amount of memory should be used. Correctly specifying the privacy model will require a consistent representation among all of its components.

Privacy Model

Creating a privacy model is done using the new_privacy constructor. The main arguments consist of the four components as outlined in the methodology section.

The internal implementation of the DA algorithm in dapper_sample requires some care in how each component is constructed.

- latent_f is an R function that samples from the latent process. Its syntax should be lik_smpl(theta) where theta is a vector representing the likelihood model parameters being estimated. This function must work with the supplied initial parameter provide in the init_par argument of dapper_sample function. The output should be a $n \times k$ matrix where k is the dimension of the vector of likelihood parameters.
- post_f is a function which makes draws from the posterior sampler. It should have the syntax post_f(dmat, theta). Here dmat is the hypothetical data set representing the confidential data. This sampler can be generated by wrapping mcmc samplers generated from other R packages (e.g. rstan, fmcmc, adaptMCMC). If using this approach, it is recommended to avoid using packages with a large initialization overhead such as mcmc since the sampler is reinitialized every loop iteration. In the case of mcmc, the Metropolis-Hastings loop is implemented in C so there is a significant initialization cost when calling from an R function. The purpose of the theta argument is to serve as the initialization point if samples from post_f are draws from say a Metropolis-Hastings sampler.
- priv_f is an R function that represents the log of the privacy mechanism density, $\eta(s_{sdp} \mid x)$.
- st_f is an R function which calculates a summary statistic. The optional argument add is a flag
 which represents whether st_f should be interpreted as satisfying record additivity or not.

Sampling

The main function in **DAPPER** is the dapper_sample function. The syntax of the function is:

The three required inputs into dapper_sample function are the privacy model (data_model), the value of the observed privatized statistic (sdp), and the total number of observations in the complete data (nobs). The dapper package is best suited for problems where the complete data can be represented in tabular form. This is because internally, it is represented as a matrix.

The optional arguments are the number of mcmc draws (niter), the burn in period (warmup), number of chains (chains) and character vector that names the parameters. Running multiple chains can be done in parallel using the furrr package. Additionally, progress can be monitored using the progressr package.

The data_model input is a privacy object that can be constructed using the new_privacy constructor. The process of constructing a privacy object will be discussed in the next section.

5 Examples

2x2 Contingency Table

As a demonstration, we analyze the UC Berkeley admissions data, which is often used as an illustrative example of Simpson's paradox. The question posed is whether the data suggest there is bias against females during the college admissions process. Below is a table of the aggregate admissions result from six departments based on sex for a total of N=4526 applicants. The table on the left represents the true admissions data and the table on the right is the result of adding independent, $N(0,100^2)$ error to each cell. Throughout the example we assume we only have access to the original total count, N, and the noise infused table.

	Male	Female		Male	Female
Admitted		557	Admitted		473
Rejected	1493	1278	Rejected	1511	1438

Below we walk through the process of defining a privacy model.

1. latent_f: Since we can condition on the table total N, we can model the original, unobserved table counts as a multinomial distribution. We can easily draw from this distribution using the rmultinom function in the base stats package. Note, in this example, the return value of one sample from rmultinom is a 4×1 matrix. In order to conform with dapper_sample we must convert the matrix to a vector.

```
latent_f <- function(theta) {
  t(rmultinom(1, 4526, theta))
}</pre>
```

2. post_f: Given confidential data, we can derive the posterior analytically using a Dirichlet prior. In this example, we use a flat prior which corresponds to Dirch(1) distribution. A sample from the Dirichlet distribution can be generated using random draws from the gamma distribution.

```
post_f <- function(dmat, theta) {
  x <- c(dmat)
  t1 <- rgamma(4, x + 1, 1)
  t1/sum(t1)
}</pre>
```

3. st_f: Since the latent model only returns one observation from a multinomial distribution, we can just use the identity function as the summary statistic.

```
st_f <- function(dmat) {
  c(dmat)
}</pre>
```

4. priv_f: The privacy mechanism is Guassian white noise drawn from independent $N(0, 100^2)$ distributions. Hence given confidential table cells $(n_{11}, n_{22}, n_{12}, n_{21})$

$$\eta(s_{dp} \mid x) = \prod \phi(s_{sd}; n_{ij}, 100^2).$$

Here $\phi(\cdot; \mu, \sigma^2)$ is the density of the normal distribution with mean and variance μ, σ^2 .

```
priv_f <- function(sdp, x) {
  dnorm(sdp - x, mean = 0, sd = 100, log = TRUE)
}</pre>
```

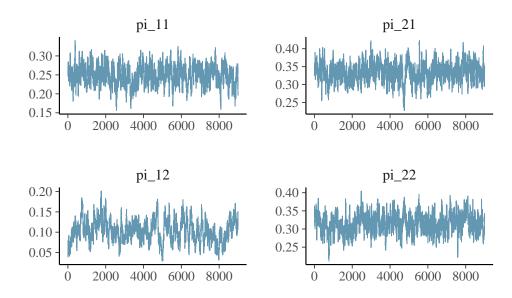
Once we have defined all components of the model we can create a new privacy model object using the new_privacy function and feed this into the dapper_sample function. Below we simulate 10,000 posterior draws with a burn-in of 1000.

```
library(dapper)
dmod <- new_privacy(post_f = post_f,</pre>
                     latent_f = latent_f,
                     priv_f = priv_f,
                              = st_f,
                     add
                              = FALSE,
                     npar
                              = 4,
                     varnames = c("pi_11", "pi_21", "pi_12", "pi_22"))
dp_out <- dapper_sample(dmod,</pre>
                   sdp = c(adm_prv),
                   niter = 10000,
                   warmup = 1000,
                   chains = 1,
                   init_par = rep(.25,4))
```

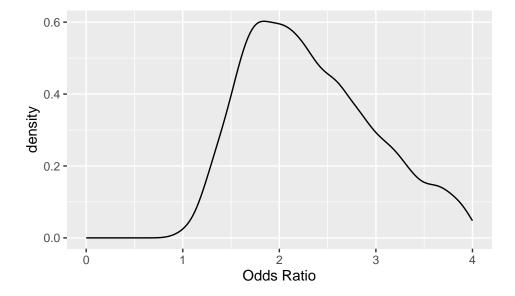
results can be quickly summarized using the summary function which is displayed below. The rhat values in the table are close to 1, which indicates the chain has run long enough to achieve adequate mixing.

```
#> # A tibble: 4 x 10
#>
    variable mean median
                            sd
                                        q5 q95 rhat ess_bulk ess_tail
                                 mad
    <chr>
           <num>
                                                                 <num>
#> 1 pi_11
            0.248   0.249   0.0254   0.0257   0.206   0.288   1.00
                                                         191.
                                                                  388.
                                                         210.
#> 2 pi_21
            0.334 0.333 0.0261 0.0254 0.291 0.377
                                                 1.00
                                                                  329.
            0.103 0.103 0.0272 0.0271 0.0586 0.149 1.02
                                                         76.0
                                                                  159.
#> 3 pi_12
#> 4 pi_22
            0.315  0.315  0.0256  0.0261  0.275  0.358  1.00
                                                         164.
                                                                  460.
```

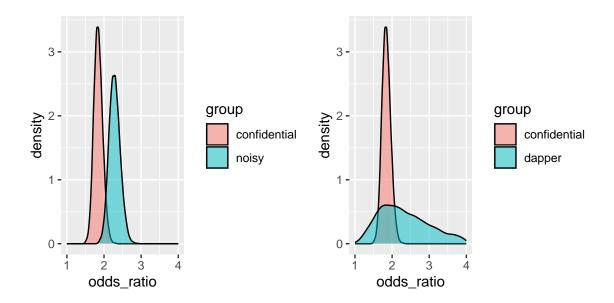
Diagnostic checks using trace plots can be done using the **Bayesplot** package as shown in figure [x]. It is especially important to check for good mixing with dapper since sticky chains are likely to be produced when the amount of injected noise is high. See the discussion section for a more detailed explanation.



To see if there is evidence of gender bias we can look at the odds ratio. Specifically, we look at the odds of a male being admitted to that of female. A higher odds ratio would indicate a bias favoring males. Figure [Z] shows the posterior draws from the dapper model. The large odds ratio values would seem to indicate there is bias favoring the males. See [XXX] for an explanation of the "paradox" of this result.



For comparison, we run a standard Bayesian analysis on the noise infused table ignoring the privacy mechanism. This will correspond exactly to the model defined in the post_f component. Figure [X] shows a density estimate for the odds ratio under the confidential and noisy data. The posterior distribution for the odds ratio under the noisy data is shifted significantly, indicating a large degree of bias. Looking at figure [Y] shows the MAP estimate from DAPPER is similar to that in the case of the confidential data. The width of the posterior is also much larger since it properly accounts for the uncertainty due to the privacy mechanism. This illustrates the dangers of ignoring the privacy mechanism. The naive analysis not only has bias, but also severely underestimate the uncertainty associated with the odds ratio estimate.



Linear Regression

Below we review an application of dapper to an example presented in (Ju et al. 2022) where they apply a Laplace privacy mechanism to a sufficient summary statistic for a linear regression model. Let $\{(x_i, y_i)\}_{i=1}^n$ be the original, confidential data with $x_i \in \mathbb{R}^2$. They assume the true data generating process follows the model

$$y = -1.79 - 2.89x_2 - 0.66x_2 + \epsilon$$
$$\epsilon \sim N(0, 2^2)$$

and consider the scenario where one desires to publicly release the sufficient summary statistics

$$t(x_i, y_i) = ((\tilde{x}^i)^T \tilde{y}_i, \tilde{y}_i^2, (\tilde{x}^i)^T \tilde{x}_i)$$

with ϵ -DP privacy guarantees. This was accomplished by clamping the data and adding Laplace random error. More precisely, we define the clamp function $[z] := \min\{\max\{z, -10\}, 10\}$ which truncates a value z so that it falls into the interval [-10, 10]. Furthermore, we let $\tilde{z} := [z]/10$ denote the normalized clamped value of z. Thus the clamped statistic is

$$t(x_i, y_i) = ((\tilde{x}^i)^T \tilde{y}_i, \tilde{y}_i^2, (\tilde{x}^i)^T \tilde{x}_i)$$

and the ϵ -DP private statistic is obtained by adding Laplace error to the unique elements of ... data generating process:

$$X \sim N_2(\mu, I_2)$$

$$\mu = \begin{pmatrix} 0.9 \\ -1.17 \end{pmatrix}$$

1. lik_f: Conditional on the table total, the table counts follow a multinomial distribution. We can easily draw from this distribution using the rmultinom function in the base stats package. Note, in this example, the return value of one sample from rmultinom is a 4 × 1 matrix. In order to conform with dapper_sample we must convert the matrix to a vector.

```
latent_f <- function(theta) {
  xmat <- MASS::mvrnorm(100 , mu = c(.9,-1.17), Sigma = diag(2))
  y <- cbind(1,xmat) %*% theta + rnorm(1, sd = sqrt(2))
  cbind(y,xmat)
}</pre>
```

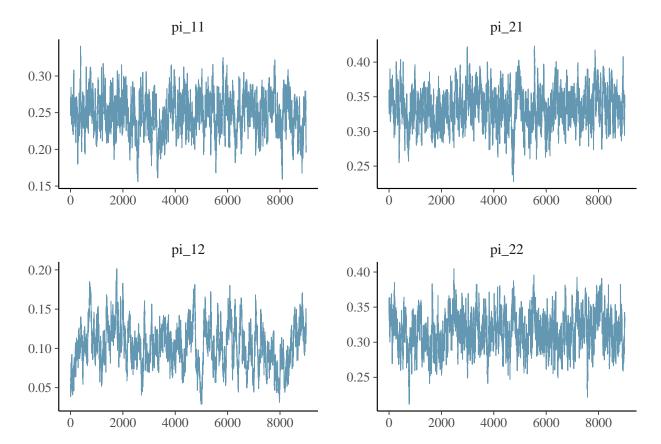
2. post_f: Given confidential data *X* we can derive the posterior analytically using a Dirichlet prior. In this example, we use a flat prior which corresponds to Dirch(1) distribution. A sample from the Dirichlet distribution can be generated using the gamma distribution via the following relation (INSERT)

```
post_f <- function(dmat, theta) {</pre>
        x <- cbind(1,dmat[,-1])</pre>
        y <- dmat[,1]</pre>
        ps_s2 \leftarrow solve((1/2) * t(x) %*% x + (1/4) * diag(3))
        ps_m \leftarrow ps_s2 \%\% (t(x) \%\% y) * (1/2)
        MASS::mvrnorm(1, mu = ps_m, Sigma = ps_s2)
   3. st_f: The complete data can be represented in two ways. Micro vs cell totals. (what section to
      introduce?) This function must return a vector.
      clamp_data <- function(dmat) {</pre>
        pmin(pmax(dmat,-10),10) / 10
      st_f <- function(dmat) {</pre>
        sdp_mat <- clamp_data(dmat)</pre>
        ydp <- sdp_mat[,1, drop = FALSE]</pre>
        xdp <- cbind(1,sdp_mat[,-1, drop = FALSE])</pre>
        s1 <- t(xdp) %*% ydp
        s2 <- t(ydp) %*% ydp
        s3 <- t(xdp) %*% xdp
        ur_s1 \leftarrow c(s1)
        ur_s2 \leftarrow c(s2)
        ur_s3 <- s3[upper.tri(s3,diag = TRUE)][-1]</pre>
        c(ur_s1,ur_s2,ur_s3)
   4. priv_f: Privacy Mechanism Guassian white noise is added to each cell total. Hence given
      confidential data (n_{11}, n_{22}, n_{12}, n_{21})

\eta(s_{dp} \mid x) = \prod \phi(s_{sd}; n_{ij}, 100^2)

      #deltaa <- 13
      #epsilon <- 10
      priv_f <- function(sdp, zt) {</pre>
        sum(VGAM::dlaplace(sdp - zt, 0, 13/10, log = TRUE))
   RUN
deltaa <- 13
epsilon <- 10
n <- 100
xmat <- MASS::mvrnorm(n, mu = c(.9, -1.17), Sigma = diag(2))
beta <- c(-1.79, -2.89, -0.66)
y <- cbind(1,xmat) %*% beta + rnorm(n, sd = sqrt(2))
z <- st_f(cbind(y,xmat))</pre>
z \leftarrow z + VGAM::rlaplace(length(z), location = 0, scale = deltaa/epsilon)
dmod <- new_privacy(post_f = post_f,</pre>
                       latent_f = latent_f,
                       priv_f = priv_f,
                       st_f
                                = st_f,
                       npar
                                = 3,
                       varnames = c("beta0", "beta1", "beta2"))
dp_out <- dapper_sample(dmod,</pre>
                           sdp = z,
                           niter = 3000
                           warmup = 1000,
                           chains = 1,
                           init_par = rep(0,3))
```

```
#> # A tibble: 4 x 10
#>
     variable
                mean median
                                 sd
                                                      q95
                                                           rhat ess_bulk ess_tail
                                        mad
#>
     <chr>
               <num>
                      <num>
                              <num>
                                      <num>
                                             <num>
                                                    <num>
                                                          <num>
                                                                    <num>
                                                                              <num>
#> 1 pi_11
                      0.249 0.0254 0.0257 0.206
                                                            1.00
                                                                    191.
                                                                               388.
               0.248
                                                    0.288
#> 2 pi_21
                                                                    210.
               0.334
                      0.333 0.0261 0.0254 0.291
                                                    0.377
                                                            1.00
                                                                               329.
#> 3 pi_12
               0.103
                      0.103 0.0272 0.0271 0.0586 0.149
                                                            1.02
                                                                     76.0
                                                                               159.
#> 4 pi_22
               0.315
                      0.315 0.0256 0.0261 0.275
                                                    0.358
                                                           1.00
                                                                    164.
                                                                               460.
```



Summary

Currently, there is a dearth of software tools privacy researchers can use to evaluate the impact of privacy mechanisms on statistical analyses. While there have been tremendous gains in the theoretical aspects of privacy, the lack of software resources to deploy and work with new privacy techniques has hampered their adoption. This gap in capability has been noted by several large industry entities who have begun building software ecosystems for working with differential privacy. SmartNoise by Microsoft [insert citation], for example, is a tool for generating synthetic data that has differentially private guarantees. However, the majority of these software tools only address privacy and not the ensuing analysis, or if it does address the analysis, only for specific models. Privacy researchers currently lack good tools for evaluating the impact of privacy mechanisms on a statistical analysis.

Thus DAPPER helps fill an urgent need by providing researchers a way to evaluate how a particular privacy mechanism might effect a statistical analysis. A notable feature is its flexibility which allows the users to specify a custom privacy mechanism. The benefit being that DAPPER can evaluate already established privacy mechanisms and those that have yet to be discovered.

While DAPPER can be a nice addition to a privacy researcher's tool kit, there is still considerably more work that can be done. From a methodological standpoint, DAPPER suffers from slow convergence when the privacy budget is small which is problematic because this is the scenario where a privacy mechanism can have the greatest impact on a statistical analysis. On the computational end, without a statistic that satisfies record additivity, computation time can be unpalatable.

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