# 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 privacy noise is injected in the workflow. The DAPPER package address approaches falling under the dissemination category. In this setting, data needs to be released to the public and privacy is maintained by first perturbing the data with random noise before being released. The U.S. Census Bureau's TopDown algorithm (Bureau 2023) is a recent high profile application of differential privacy in the dissemination setting. For a nice survey of the direct and query based methods, see the paper by (Ji, Lipton, and Elkan 2014) which provides details on how to modify several popular machine learning algorithms to have privacy guarantees.

While the aforementioned modifications will guarantee some form of privacy, they will not, however, guarantee correct statistical inference. As an example, one instance of implementing the dissemination approach for tabular data involves directly adding independent, random error to each cell. In the regression setting, 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.

However, the marginal likelihood that would result from correctly accounting for the injected privacy noise, is often analytically intractable. 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.

The DAPPER package provides a tool for conducting valid statistical inference in the presence of privacy noise. It implements a 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 gives the user a great deal of flexibility by allowing the user to input a sampler from an existing Bayesian workflow, unadjusted for privacy noise, and automatically constructing a valid posterior sampler that accounts for the added privacy noise.

With respects to the dissemination approach described in the previous paragraph, there is a large body of literature establishing methods for fitting regression models with measurement errors in the covariates. Though, work in this area mostly focused on methods which did not require fully specifying the measurement error model, since this was often assumed unknown.

The statistical workflow typically assumes no measurement error in the observed covariate data. In the presence of such errors, standard estimators can exhibit significant bias (Gong 2022). Therefore, fitting standard statistical models after adding noise directly to the data for privacy can lead to incorrect inference. Adjusting models to take into account noisy covariates has a rich history spanning several decades. For textbook length treatments see (Yi 2017; Carroll et al. 2006). Prior work mostly focuses on methods which do not require fully specifying the measurement error model, since this was often 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.

# 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  $\theta$  has relevant meaning to the scientific question at hand. In this setting, learning characteristics of a population reduces to learning about  $\theta$ .

In the Bayesian framework, this 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. In the context of privacy, providing a summary statistic can offer a level of anonymity.

### **Differential Privacy**

While a summary statistic can already anonymize the data, it is still possible to deduce information about an individual entry depending on the distribution of x. For additional anonymity, one idea is to add noise to the summary statistic s. More formally we write  $s_{dp} \sim \eta(\cdot \mid x)$ . Here,  $s_{dp}$  is the noise infused version of s and  $\eta$  is a known noise infusion process. The privacy mechanism  $\eta$  is said to be  $\epsilon$ -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 (denoated by d(x, x')), the probability ratio is bounded:

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

The parameter  $\epsilon$  is called the privacy loss budget, and controls how informative  $s_{dv}$  is about x. Larger values of  $\epsilon$  correspond to less noise being added.

#### **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.

## 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,  $p(s_{dy} \mid heta)$  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  $p(\theta \mid s_{dp})$  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

- $\begin{array}{ll} \bullet & p(\theta \mid x, s_{dp}) \\ \bullet & p(x \mid \theta, s_{dp}) \end{array}$

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,  $p(x \mid \theta, s_{dv})$ , 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 summary statistic can be written as the sum of individual contributions from each observation. More precisely, we say a statistic 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 algorithm is in the following pseudo code:

- 1. Sample  $\theta^{t+1}$  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:

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

# 4 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 left up to user. 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 are often multiple ways of doing this. 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  $2 \times 2$  matrix, or a  $100 \times 2$  matrix. Both are mathematically equivalent, but the  $2 \times 2$  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 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.

- lik\_smpl is an R function that samples from the likelihood. 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\_smpl is a function which represents the posterior sampler. It should have the call signature post\_smpl(dmat, theta). Where dmat is the complete 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 such as mcmc whose implementation clashes with gdp\_sample. In the case of mcmc, the Metropolis-Hastings loop is implemented in C which incurs a very large overhead in gdp\_sample since it is reinitialized

every iteration. In general, repeatedly calling an R function that hooks into C code is slow. (NOT QUITE ACCURATE FIX LATER)

- 11\_priv\_mech is an R function that represents the log-likelihood of  $\eta(s_{sdp} \mid x)$ . The function can output the log likelihood up to an additive constant.
- st\_calc is an R function which calculates the summary statistic. The optional argument add is a flag which represents whether T is additive 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

A common procedure when analyzing contingency tables is to estimate the odds ratio. Something something about safetab to connect back to DP (dont forget citation!). As a demonstration, we analyze the UC Berkeley admissions data, which is often used as an illustrative example of Simpson's paradox. The question 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.

	Male	Female		Male	Female
Admitted Rejected	1198 1493	557 1278	Admitted Rejected	1135 1511	473 1438
- Trejected	1170	1270	- rejected	1011	1100

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

1. lik\_smpl: 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.

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

 post\_smpl: 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_smpl <- function(dmat, theta) {
  x <- c(dmat)
  t1 <- rgamma(length(theta), x + 1, 1)
  t1/sum(t1)
}</pre>
```

3. st\_calc: The complete data can be represented in two ways. Micro vs cell totals. (what section to introduce?) This function must return a vector.

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

4. 11\_priv\_mech: Privacy Mechanism Guassian white noise is added to each cell total. Hence given confidential data ( $n_{11}$ ,  $n_{22}$ ,  $n_{12}$ ,  $n_{21}$ )

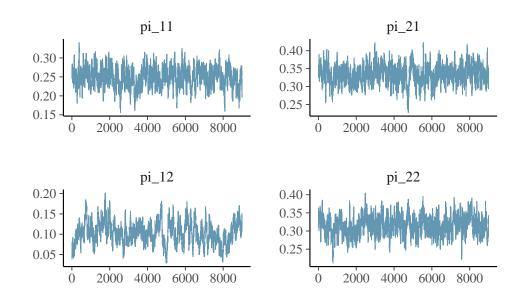
```
\eta(s_{dp}\mid x) = \prod \phi(s_{sd};n_{ij},100^2) ll_priv_mech <- function(sdp, x) { dnorm(sdp - x, mean = 0, sd = 100, log = TRUE) }
```

Once privacy model has been defined we can run gdp\_sample

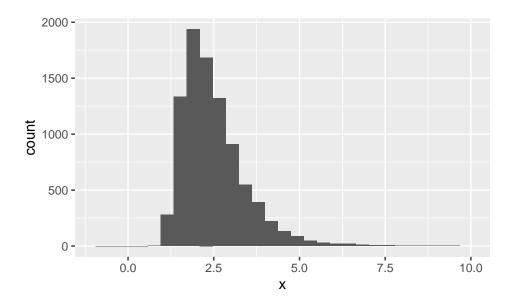
results can be quickly summarized using the summary function

```
#> # A tibble: 4 x 10
#>
    variable mean median
                               sd
                                                      rhat ess_bulk ess_tail
                                             q5
                                                  a95
                                     mad
#>
    <chr>
             <num> <num> <num> <num>
                                                               <num>
                                                                        <num>
                                         <num> <num>
                                                      <num>
#> 1 pi_11
             0.248
                    0.249 0.0254 0.0257 0.206 0.288
                                                       1.00
                                                               191.
                                                                         388.
#> 2 pi_21
             0.334 0.333 0.0261 0.0254 0.291
                                                0.377
                                                       1.00
                                                               210.
                                                                         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.
```

Diagnostic checks can be done using the **Bayesplot** package.



log odds distribution



(Insert gdp Analysis Here)

For clean data, a estimate for the odds ratio and a confidence interval can be constructed using Woolf's method (i.e. A wald confidence interval). It uses the fact that the log of the odds ratio is approximately normal for large sample sizes.

$$\log\left(\frac{n_{11} \cdot n_{22}}{n_{12} \cdot n_{21}}\right) \pm z_{\alpha/2} \sqrt{\frac{1}{n_{11}} + \frac{1}{n_{22}} + \frac{1}{n_{12}} + \frac{1}{n_{21}}}$$

```
or_confint <- function(x, alpha) {
    or <- log(x[1] * x[4]/ (x[2] * x[3]))
    se <- sqrt(sum(1/x))
    c(or - qnorm(alpha/2) * se, or + qnorm(alpha/2) * se)
}
#clean data
exp(or_confint(x, .95))
#> [1] 1.848471 1.833718
#privitized data
exp(or_confint(sdp, .95))
#> [1] 2.293115 2.274220
```

## **Logistic Regression**

## 6 Summary

This package is cool. You should install it.

## References

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