# HARVARD PRIVACY TOOLS PROJECT

# Data Processing Notes

# CONTENTS

1	Overview	1
2	Clamping2.1 Introduction2.2 Algorithm Statement	
3	Imputation3.1 Introduction3.2 Algorithm Statement	
4	Resize         4.1 Motivation	5 5 6 7
	4.7 Accuracy/Error	7

# 1 Overview

This document contains notes on our choices regarding "data processing", which I am using to stand in for all things related to data bookkeeping (data bounds, imputation, sample size calculations, etc.).

# 2 Clamping

#### 2.1 Introduction

Many operations in the library require the definition of a set of possible data values we would like to use for said operation. For numeric variables, this typically takes the form of a closed interval  $[min, max] \in \mathbb{R}$  or  $\mathbb{Z}$ . For categorical variables, this is a discrete set of elements. The clamp component sets these properties for a given analysis. In our system, we have slightly different notions of clamping for numeric and categorical variables. We consider f64 and i64 to be allowable numeric types, and i64, bool, and String to be allowable categorical types.

Our method for clamping numeric types should be familiar; it requires data, a lower bound, and an upper bound, and maps data elements outside of the interval [min, max] to the nearest point in the interval. Null values (represented as NAN for the f64 type and which do not exist for the i64 type) are preserved by numeric clamping.

Our categorical clamping requires data, a set of feasible values, and a null value. Data elements that are outside the set of feasible values are mapped to the null value. Null values are preserved by the categorical clamping.

# 2.2 Algorithm Statement

The component arguments are as follows:

- 1. X: The data of size  $n \times m$  to be clamped.
- 2. categories: A vector of length m, where each element is the set of plausible data values for column m of X. If not provided, defaults to None.
- 3. null\_value: A vector of length m, where each element is the designated "null" for column m of X. That is, elements  $\notin$  categories are mapped to the null\_value. If not provided, defaults to None.
- 4. lower: A vector of length m, where each element is a proposed minimum value for column m of X. If not provided, defaults to None.
- 5. upper: A vector of length m, where each element is a proposed maximum value for column m of X. If not provided, defaults to None.

#### **Algorithm 1** Clamping: $\operatorname{clamp}(X, \operatorname{categories}, \operatorname{lower}, \operatorname{upper})$

```
1: for j \in [m] do
                                                                               ▶ Iterate over columns
        for i \in [n] do

    ▶ Iterate over rows

 2:
            if categories \neq None then
                                                               ▶ Proceed with categorical clamping
 3:
                if X_{i,j} \not\in categories[j] then
 4:
                    X_{i,j} \leftarrow null \ value[j]
 5:
                end if
 6:
            else
                                                                  ▶ Proceed with numeric clamping
 7:
                if X_{i,j} < lower[j] then
 8:
9:
                    X_{i,j} \leftarrow lower[j]
                else if X_{i,j} > upper[j] then
10:
                    X_{i,j} \leftarrow upper[j]
11:
                end if
12:
            end if
13:
        end for
14:
15: end for
16: return X
```

# 3 Imputation

#### 3.1 Introduction

Some operations in the library require the private data to be non-null. The *impute* component sets the non-null property for an analysis.

Much like clamping, we have slightly different notions of imputation for numeric and categorical variables. In this case, we consider only f64 to be an allowable numeric type, while f64, i64, bool, and String are allowable categorical types. We do not allow for numeric imputation on i64 because i64 does not support NAN values and thus are non-null by default.

Numeric imputation is parameterized by min, max, and a supported probability distribution (Gaussian or Uniform) with appropriate arguments. NAN values are replaced with a draw from the provided distribution.

Categorical imputation is parameterized by categories, probabilities, and a null value. Each data element equal to the null value is replaced with a draw from the set of categories, according to the probabilities provided.

#### 3.2 Algorithm Statement

The component arguments are as follows:

- 1. X: The data of size  $n \times m$  for which values should be imputed.
- 2. categories: A vector of length m, where each element is the set of plausible data values for column m of X. If not provided, defaults to None.
- 3. weights: A vector of length m, where each element is a set of imputation weights associated with the categories for column m of X. If not provided, defaults to None.
- 4. null\_value: A vector of length m, where each element is the designated "null" for column m of X. If an element is equal to the corresponding null\_value for its column, it is imputed from the probability distribution over the categories defined by the weights.

- 5. distribution: A string providing the distribution from which elements will be imputed. Currently supported options are "Uniform" and "Gaussian". Used only if *categories* == None. Defaults to "Uniform".
- 6. shift: A vector of length m, where each element is the expectation of the Gaussian distribution used for imputing missing values in column m of X. Used only if distribution == "Gaussian". Defaults to None.
- 7. scale: A vector of length m, where each element is the standard deviation of the Gaussian distribution used for imputing missing values in column m of X. Used only if distribution == "Gaussian". Defaults to None.
- 8. lower: A vector of length m, where each element is a proposed minimum value for column m of X. Acts as a lower bound for the distribution from which elements are imputed. If not provided, defaults to None.
- 9. upper: A vector of length m, where each element is a proposed maximum value for column m of X. Acts as a lower bound for the distribution from which elements are imputed. If not provided, defaults to None.

Let Categorical $(k, (p_1, \ldots, p_k))$  be the categorical distribution, which chooses a single element from the set  $\{1, 2, \ldots, k\}$  where each element i has probability of being chosen  $p_i$ . Then we have

#### Algorithm 2 Imputation:

impute(X, categories, weights, null value, distribution, shift, scale, lower, upper)

```
1: for j \in [m] do
                                                                                 ▶ Iterate over columns
        for i \in [n] do

    ▶ Iterate over rows

 2:
            if categories \neq None then
                                                              ▶ Proceed with categorical imputation
 3:
                if X_{i,j} == null\_value[j] then
 4:
                     sampling_index \leftarrow Categorical(|weights[j]|, weights[j])
 5:
                     X_{i,j} \leftarrow \text{categories}[j][\text{sampling index}]
 6:
 7:
                 end if
            else
                                                                 ▶ Proceed with numeric imputation
 8:
                if X_{i,j} == NAN then
 9:
                     if distribution == "Uniform" then
10:
                         X_{i,j} \leftarrow \text{Uniform(lower[j], upper[j])}
11:
                     else if distribution == "Gaussian" then
12:
                         X_{i,j} \leftarrow \text{Gaussian}(\text{shift}[j], \text{scale}[j])
13:
                         if X_{i,j} < lower[j] then
14:
                             X_{i,j} \leftarrow lower[j]
15:
16:
                         else if X_{i,j} > upper[j] then
                             X_{i,j} \leftarrow upper[j]
17:
                         end if
18:
                     end if
19:
                 end if
20:
            end if
21:
22:
        end for
23: end for
24: return X
```

#### 4 Resize

#### 4.1 Motivation

We want our library to support cases both in which the analyst does and does not have access to the number of records in the data. Doing the latter well is an under-explored problem, so we propose the *resize* framework as a possible solution.

#### 4.2 Goals

The resize framework is a means of jointly achieving a few different goals:

- 1. guarantee known n for analyses that require it,
- 2. give users flexibility in how they trade off between bias and variance, and
- 3. allow for both c-stability and privacy amplification from subsampling within the same privacy calculus.

#### 4.3 Algorithm Statement

11: return  $(X', \epsilon', \delta')$ 

The resize function takes the following inputs:

- 1. X: The private underlying data.
- 2.  $\tilde{n}$ : The size of the private underlying data.

**Algorithm 3** Resize: resize(X, n, p, neighboring, ...)

- 3. n: The desired size of the new data.
- 4. p: The proportion of the underlying data that can be used to construct the new data. p can be > 1.
- 5. ...: Various arguments explaining imputation rules (these will follow the rules from section 3).

Let sample(Y, m) be a function that samples m elements from data set Y without replacement. Let Aug(Y, m, ...) be a function that imputes new elements independent of the data (using imputation parameters given by ...) for a data set Y until it is of size m. The algorithm will look something like the following:

```
1: c \leftarrow \lceil p \rceil
                                                                                      ⊳ sets c-stability property
 2: s \leftarrow p/c
                                                                   > sets subsampled proportion property
 3: X_c \leftarrow \bigcup_{i=1}^c X
                                       \triangleright create new database of size c\tilde{n}, composed of c copies of X
 4: if neighboring == "replace one" then
         m \leftarrow |sc\tilde{n}| > \text{number of records that can be filled using subsampled private data}
 6: else if neighboring == "add/remove one" then
         m \leftarrow Binomial(c\tilde{n}, s)
                                            > number of records that can be filled using subsampled
    private data
 8: end if
 9: (\epsilon', \delta') \leftarrow \left(\log\left(1 + s\left(e^{c\epsilon} - 1\right)\right), s\left(\sum_{i=0}^{c-1} e^{i\epsilon}\right)\delta\right)
                                                                         ▷ privacy amplification via
10: X' \leftarrow sample(X_c, \max(m, n), neighboring) \bigcup (Aug(\emptyset, \max(0, n - m), \ldots))
```

The  $\epsilon'$ ,  $\delta'$  terms come from first applying the group privacy definition with group size c to the database  $X_c$  to get  $\left(c\epsilon, \left(\sum_{i=1}^{c-1} e^{i\epsilon}\right)\delta\right)$  [Vad17] and then applying privacy amplification by subsampling results from Theorems 8 and 9 of [BBG18]. Note that, for the "replace

one" definition, we could be using  $\frac{m}{c\tilde{n}}$  instead of s in the privacy calculation. Using s gives us a very slightly worse privacy guarantee (the only difference is the  $\lfloor \cdot \rfloor$  we used to get m), but is nice for consistency between the methods and not having to keep track of m as an extra property.

# 4.4 Functional Privacy Parameters

We established in section 4.3 that a user asking for an  $(\epsilon, \delta)$ -DP guarantee will get an  $(\epsilon', \delta')$ -DP guarantee with respect to the original private data. What we'd really like, however, is for the user to ask for an  $(\epsilon, \delta)$ -DP guarantee and have the library come up with what we will call a functional  $(\epsilon_f, \delta_f)$  that will ensure  $(\epsilon, \delta)$ -DP on the original data. Any components that operate on the resized data will use the functional  $(\epsilon_f, \delta_f)$  internally instead of the parameters passed by the user.

**Theorem 1.** A mechanism that respects

$$(\epsilon_f, \delta_f) = \left(\frac{1}{c}\log\left(\frac{e^{\epsilon} - 1}{s} + 1\right), \frac{\delta}{s\left(\sum_{i=1}^{c-1} e^{i\epsilon}\right)}\right) - DP$$

on the resized data respects  $(\epsilon, \delta)$ -DP on the true private data.

*Proof.* We know that an  $(\epsilon, \delta)$ -DP on the resized data corresponds to a

$$(\epsilon', \delta') = \left(\log\left(1 + s\left(e^{c\epsilon} - 1\right)\right), s\left(\sum_{i=1}^{c-1} e^{i\epsilon}\right)\delta\right)$$

guarantee on the original data. But we want our mechanism to respect  $(\epsilon_f, \delta_f)$  on the resized data such that it respects  $(\epsilon, \delta)$  on the original data. So we replace  $(\epsilon, \delta)$  with  $\epsilon_f, \delta_f$  and  $(\epsilon', \delta')$  with  $(\epsilon, \delta)$ . Then we can invert the function to get what we want.

Let's start with  $\epsilon, \epsilon_f$ :

$$\epsilon = \log (1 + s (e^{c\epsilon_f} - 1))$$

$$e^{\epsilon} = 1 + s (e^{c\epsilon_f} - 1)$$

$$\frac{e^{\epsilon} - 1}{s} = e^{c\epsilon_f} - 1$$

$$\frac{1}{c} \log \left( \frac{e^{\epsilon} - 1}{s} + 1 \right) = \epsilon_f.$$

We carry out a similar calculation for  $\delta, \delta_f$ :

$$\delta = s \left( \sum_{i=1}^{c-1} e^{i\epsilon} \right) \delta_f$$

$$\frac{\delta}{s \left( \sum_{i=1}^{c-1} e^{i\epsilon} \right)} = \delta_f.$$

So, in order for a mechanism to respect  $(\epsilon, \delta)$ -DP on the original data, it must respect  $\left(\frac{1}{c}\log\left(\frac{e^{\epsilon}-1}{s}+1\right), \frac{\delta}{s\left(\sum_{i=1}^{c-1}e^{i\epsilon}\right)}\right)$ -DP on the resized data.

We now present an mini-algorithm for finding the functional  $(\epsilon, \delta)$ .

# **Algorithm 4** Finding Functional $(\epsilon_f, \delta_f)$ : get\_func\_priv $(p, \epsilon, \delta)$

```
1: c \leftarrow \lceil p \rceil  \triangleright sets c-stability property
2: s \leftarrow p/c  \triangleright sets subsampled_proportion property
3: \epsilon_f \leftarrow \frac{1}{c} \log \left( \frac{e^{\epsilon} - 1}{s} + 1 \right)
4: \delta_f \leftarrow \frac{\delta}{s\left(\sum_{i=1}^{c-1} e^{i\epsilon}\right)}
5: return (\epsilon_f, \delta_f)
```

#### 4.5 Examples

Let X be such that  $\tilde{n} = |X| = 100$ . We can look at a few examples of calls to the resize function (which will return X') and check the behavior.

- 1.  $\operatorname{resize}(X, 150, 1, ...)$ : X' will be made up of the 100 true elements of X and 50 imputed values. The functional privacy parameters are identical to the ones the user provides.
- 2.  $\operatorname{resize}(X, 100, 0.75, ...)$ : X' will be made up of 75 true elements of X and 25 imputed values. The functional privacy parameters will benefit (lower noise) from amplification via subsampling.
- 3. resize(X, 90, 1.5, ...): X' will be a random sample of  $X \cup X$  of size 90. The functional privacy parameters will lead to greater noise than what the user provides, as they have to take into account the new c-stability of 2. This example is illustrative in that is shows that the functional privacy usage is affected only by p it has nothing to do with the relative sizes of the X and X'.

#### 4.6 How to choose n

To this point, we have ignored how a user would go about choosing n. If no trusted public metadata exist, we suggest estimating the size of the private database (e.g. using the Geometric mechanism) and using that release for n.

#### 4.7 Accuracy/Error

The accuracy/error calculations in the library are always relative to the resized data X', rather than the true private data X. We are currently thinking about how best to extend guarantees from X' to X.

# REFERENCES

- [BBG18] Borja Balle, Gilles Barthe, and Marco Gaboardi. Privacy amplification by subsampling: Tight analyses via couplings and divergences. CoRR, abs/1807.01647, 2018.
- [Vad17] Salil Vadhan. The complexity of differential privacy. In *Tutorials on the Foun-dations of Cryptography*, pages 347–450. Springer, 2017.