# ACHIEVING k-ANONYMITY PRIVACY PROTECTION USING GENERALIZATION AND SUPPRESSION<sup>1</sup>

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Often a data holder, such as a hospital or bank, needs to share person-specific records in such a way that the identities of the individuals who are the subjects of the data cannot be determined. One way to achieve this is to have the released records adhere to kanonymity, which means each released record has at least (k-1) other records in the release whose values are indistinct over those fields that appear in external data. So, kanonymity provides privacy protection by guaranteeing that each released record will relate to at least k individuals even if the records are directly linked to external information. This paper provides a formal presentation of combining generalization and suppression to achieve k-anonymity. Generalization involves replacing (or recoding) a value with a less specific but semantically consistent value. Suppression involves not releasing a value at all. The Preferred Minimal Generalization Algorithm (MinGen), which is a theoretical algorithm presented herein, combines these techniques to provide k-anonymity protection with minimal distortion. The real-world algorithms Datafly and μ-Argus are compared to MinGen. Both Datafly and μ-Argus use heuristics to make approximations, and so, they do not always yield optimal results. It is shown that Datafly can over distort data and μ-Argus can additionally fail to provide adequate protection.

Keywords: Data anonymity; data privacy; re-identification; data fusion; privacy.

### 1. Introduction

Today's globally networked society places great demand on the collection and sharing of person-specific data for many new uses [1]. This happens at a time when more and more historically public information is also electronically available. When these data are linked together, they provide an electronic image of a person that is as identifying and personal as a fingerprint even when the information contains no explicit identifiers, such as name and phone number. Other distinctive data, such as birth date and postal code, often combine uniquely [2] and can be linked to publicly available information to re-identify individuals.

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<sup>&</sup>lt;sup>1</sup> This paper significantly amends and expands the earlier paper "Protecting privacy when disclosing information: *k*-anonymity and its enforcement through generalization and suppression" (with Samarati) submitted to IEEE Security and Privacy 1998, and extends parts of my Ph.D. thesis [10].

So in today's technically-empowered data rich environment, how does a data holder, such as a medical institution, public health agency, or financial organization, share person-specific records in such a way that the released information remain practically useful but the identity of the individuals who are the subjects of the data cannot be determined? One way to achieve this is to have the released information adhere to *k*-anonymity [3]. A release of data is said to adhere to *k*-anonymity if each released record has at least (*k*-1) other records also visible in the release whose values are indistinct over a special set of fields called the quasi-identifier [4]. The quasi-identifier contains those fields that are likely to appear in other known data sets. Therefore, *k*-anonymity provides privacy protection by guaranteeing that each record relates to at least *k* individuals even if the released records are directly linked (or matched) to external information.

This paper provides a formal presentation of achieving k-anonymity using generalization and suppression. Generalization involves replacing (or recoding) a value with a less specific but semantically consistent value. Suppression involves not releasing a value at all. While there are numerous techniques available<sup>2</sup>, combining these two offers several advantages.

First, a recipient of the data can be told what was done to the data. This allows results drawn from released data to be properly interpreted. Second, information reported on each person is "truthful" which makes resulting data useful for fraud detection, counter-terrorism surveillance, healthcare outcome assessments and other uses involving traceable person-specific patterns<sup>3</sup>. Third, these techniques can provide results with guarantees of anonymity that are minimally distorted. Any attempt to provide anonymity protection, no matter how minor, involves modifying the data and thereby distorting its contents, so the goal is to distort minimally. Fourth, these techniques can be used with preferences a recipient of the released data may have, thereby providing the most useful data possible. In this way, algorithmic decisions about how to distort the data can have minimal impact on the data's fitness for a particular task.

Finally, the real-world systems Datafly [5] and  $\mu$ -Argus [6], which are discussed in subsequent sections, use these techniques to achieve k-anonymity. Therefore, this work provides a formal basis for comparing them.

#### 2. Background

The ideas of k-anonymity and of a quasi-identifier are straightforward. Nevertheless, care must be taken to precisely state what is meant. [3] provides a detailed discussion of k-anonymity. A brief summary is provided in this section as background for the upcoming presentations on generalization and suppression.

<sup>&</sup>lt;sup>2</sup> See Willenborg and De Waal [2] for a list of traditional statistical techniques.

<sup>&</sup>lt;sup>3</sup> In contrast, other techniques (e.g., additive noise) can destroy the "truthfulness" of the information reported on a person even though they can maintain aggregate statistical properties.

Unless otherwise stated, the term data refers to person-specific information that is conceptually organized as a table of rows (or records) and columns (or fields). Each row is termed a tuple. Tuples within a table are not necessarily unique. Each column is called an attribute and denotes a semantic category of information that is a set of possible values; therefore, an attribute is also a domain. Attributes within a table are unique. So by observing a table, each row is an ordered *n*-tuple of values  $\langle d_1, d_2, ..., d_n \rangle$  such that each value  $d_i$  is in the domain of the j-th column, for j=1, 2, ..., n where n is the number of columns. This corresponds to relational database concepts [7].

Let  $B(A_1,...,A_n)$  be a table with a finite number of tuples. The finite set of attributes of B are  $\{A_1,...,A_n\}$ . Given a table  $B(A_1,...,A_n)$ ,  $\{A_i,...,A_j\} \subseteq \{A_1,...,A_n\}$ , and a tuple  $t \in B$ , I use  $t[A_i,...,A_i]$  to denote the sequence of the values,  $v_i,...,v_j$ , of  $A_i,...,A_i$  in t. I use  $B[A_i,...,A_i]$  to denote the projection, maintaining duplicate tuples, of attributes  $A_i, ..., A_i$  in B.

Throughout this work each tuple is assumed to be specific to a person and no two tuples pertain to the same person. This assumption simplifies discussion without loss of applicability. Also, this discussion focuses on protecting identity in person-specific data, but is just as applicable to protecting other kinds of information about other kinds of entities (e.g., companies or governments).

Limiting the ability to link (or match) released data to other external information offers privacy protection. Attributes in the private information that could be used for linking with external information are termed the quasi-identifier. Such attributes not only include explicit identifiers such as name, address, and phone number, but also include attributes that in combination can uniquely identify individuals such as birth date, ZIP<sup>4</sup>, and gender [8]. A goal of this work is to release person-specific data such that the ability to link to other information using the quasi-identifier is limited.

#### **Definition 1. Ouasi-identifier**

Given a population U, a person-specific table  $T(A_1,...,A_n)$ ,  $f_c: U \to T$  and  $f_g: T$  $\rightarrow$  U', where U  $\subseteq$  U'. A quasi-identifier of T, written  $Q_T$ , is a set of attributes  $\{A_i, ..., A_i\} \subseteq \{A_1, ..., A_n\}$  where:  $\exists p_i \in U$  such that  $f_g(f_c(p_i)[Q_T]) = p_i$ .

# Definition 2. k-anonymity

Let  $RT(A_1,...,A_n)$  be a table and  $QI_{RT}$  be the quasi-identifier associated with it. RT is said to satisfy k-anonymity if and only if each sequence of values in  $\mathsf{RT}[QI_{RT}]$  appears with at least k occurrences in  $\mathsf{RT}[QI_{RT}]$ .

<sup>&</sup>lt;sup>4</sup> In the United States, a ZIP code refers to the postal code. Typically 5-digit ZIP codes are used, though 9-digit ZIP codes have been assigned. A 5-digit code is the first 5 digits of the 9-digit code.

# Example 1. Table adhering to k-anonymity

Figure 1 contains table T, which adheres to k-anonymity. The quasi-identifier is  $QI_T = \{Race, Birth, Gender, ZIP\}$  and k=2. Therefore, for each of the tuples contained in T, the values that comprise the quasi-identifier appear at least twice in T. In particular,  $tI[QI_T] = t2[QI_T]$ ,  $t3[QI_T] = t4[QI_T]$ , and  $t5[QI_T] = t6[QI_T] = t7[QI_T]$ .

	Race	Birth	Gender	ZIP	Problem
t1	Black	1965	m	02141	short breath
t2	Black	1965	m	02141	chest pain
t3	Black	1964	f	02138	obesity
t4	Black	1964	f	02138	chest pain
t5	White	1964	m	02138	chest pain
t6	White	1964	m	02138	obesity
t7	White	1964	m	02138	short breath

Figure 1. Example of *k*-anonymity, where k = 2 and  $QI = \{Race, Birth, Gender, ZIP\}$ .

#### Theorem 1

Let  $\mathsf{RT}(A_1,...,A_n)$  be a table,  $QI_{RT} = (A_i,...,A_j)$  be the quasi-identifier associated with  $\mathsf{RT}, A_i,...,A_j \subseteq A_1,...,A_n$ , and  $\mathsf{RT}$  satisfy k-anonymity. Then, each sequence of values in  $\mathsf{RT}[A_x]$  appears with at least k occurrences in  $\mathsf{RT}[QI_{RT}]$  for x=i,...,j.

#### Example 2. k occurrences of each value under k-anonymity

Table T in Figure 1 adheres to k-anonymity. Therefore, each value associated with an attribute of QI in T appears at least k times. |T[Race = "black"]| = 4. |T[Race = "white"]| = 3. |T[Birth = "1964"]| = 5. |T[Birth = "1965"]| = 2. |T[Gender = "m"]| = 5. |T[Gender = "f"]| = 2. |T[ZIP = "02138"]| = 5. And, |T[ZIP = "02141"]| = 2.

It can be trivially proven that if the released data RT satisfies k-anonymity with respect to the quasi-identifier  $Ql_{PT}$ , then the combination of the released data RT and the external sources on which  $Ql_{PT}$  was based, cannot link on  $Ql_{PT}$  or a subset of its attributes to match fewer than k individuals.

#### 3. Methods

In this section I present formal notions of: (1) generalization incorporating suppression; (2) minimal generalization; and, (3) minimal distortion. The Preferred Minimal Generalization Algorithm (MinGen), which ends this section, combines these notions into a theoretical algorithm that uses generalization and suppression to produce tables that adhere to *k*-anonymity with minimal distortion.

#### 3.1. Generalization including suppression

The idea of generalizing an attribute is a simple concept. A value is replaced by a less specific, more general value that is faithful to the original. In Figure 2 the original ZIP codes {02138, 02139} can be generalized to 0213\*, thereby stripping the rightmost digit and semantically indicating a larger geographical area.

In a classical relational database system, domains are used to describe the set of values that attributes assume. For example, there might be a ZIP domain, a number domain and a string domain. I extend this notion of a domain to make it easier to describe how to generalize the values of an attribute. In the original database, where every value is as specific as possible, every attribute is considered to be in a ground domain. For example, 02139 is in the ground ZIP domain, Z<sub>0</sub>. In order to achieve k-anonymity I can make ZIP codes less informative. I do this by saying that there is a more general, less specific domain that can be used to describe ZIPs, say Z<sub>1</sub>, in which the last digit has been replaced by 0 (or removed altogether). There is also a mapping from  $Z_0$  to  $Z_1$ , such as  $02139 \rightarrow 0213^*$ .

Given an attribute A, I say a generalization for an attribute is a function on A. That is, each  $f: A \rightarrow B$  is a generalization. I also say that:

$$A_0 \xrightarrow{f_0} A_1 \xrightarrow{f_1} \dots \xrightarrow{f_{n-1}} A_n$$

is a generalization sequence or a functional generalization sequence.

Given an attribute A of a private table PT, I define a domain generalization **hierarchy** DGH<sub>A</sub> for A as a set of functions  $f_h: h=0,...,n-1$  such that:

$$A_o \xrightarrow{f_0} A_1 \xrightarrow{f_1} \dots \xrightarrow{f_{n-1}} A_n$$

$$A = A_0 \text{ and } |A_n| = 1. \text{ DGH}_A \text{ is over:} \qquad \bigcup_{i=1}^n A_i$$

Clearly, the  $f_h$ 's impose a linear ordering on the  $A_h$ 's where the minimal element is the ground domain  $A_0$  and the maximal element is  $A_n$ . The singleton requirement on  $A_n$  ensures that all values associated with an attribute can eventually be generalized to a single value. In this presentation I assume  $A_h$ , h=0,...,n, are disjoint; if an implementation is to the contrary and there are elements in common, then  $DGH_A$  is over the disjoint sum of  $A_h$ 's and definitions change accordingly.

Given a domain generalization hierarchy DGH<sub>A</sub> for an attribute A, if  $v_i \in A_i$ and  $v_i \in A_i$  then I say  $v_i \le v_j$  if and only if  $i \le j$  and:

$$f_{j-1}(\ldots f_i(v_i)\ldots) = v_j$$

This defines a partial ordering  $\leq$  on:  $\bigcup_{A_h}^{n} A_h$ 

Such a relationship implies the existence of a value generalization hierarchy  $VGH_A$  for attribute A.

I expand my representation of generalization to include suppression by imposing on each value generalization hierarchy a new maximal element, atop the old maximal element. The new maximal element is the attribute's suppressed

value. The height of each value generalization hierarchy is thereby incremented by one. No other changes are necessary to incorporate suppression. Figure 2 and Figure 3 provides examples of domain and value generalization hierarchies expanded to include the suppressed maximal element (\*\*\*\*\*). In this example, domain  $Z_0$  represents ZIP codes for Cambridge, MA, and  $E_0$  represents race. From now on, all references to generalization include the new maximal element; and, hierarchy refers to domain generalization hierarchies unless otherwise noted.

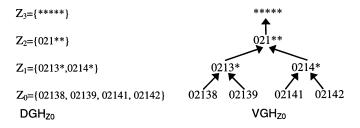


Figure 2. ZIP domain and value generalization hierarchies including suppression.

$$Z_{2=\{******\}}$$
 
$$Z_{1=\{Person\}}$$
 
$$Z_{0=\{Asian,Black,White\}}$$
 
$$Asian Black White 
$$DGH_{E0}$$
 
$$VGH_{E0}$$$$

Figure 3. Race domain and value generalization hierarchies including suppression.

# 3.2. Minimal generalization of a table

Given table PT, generalization can be effective in producing a table RT based on PT that adheres to *k*-anonymity because values in RT are substituted with their generalized replacements. The number of distinct values associated with each attribute is non-increasing, so the substitution tends to map values to the same result, thereby possibly decreasing the number of distinct tuples in RT.

A generalization function on tuple t with respect to  $A_1,...,A_n$  is a function  $f_t$  on  $A_1 \times ... \times A_n$  such that:  $f_t(A_1,...,A_n) = (f_{t1}(A_1),...,f_m(A_n))$ 

where for each i: 1,...,n,  $f_{ti}$  is a generalization of the value  $t[A_i]$ . The function  $f_t$  is a set function. I say  $f_t$  is generated by the  $f_{ti}$ 's.

Given 
$$f, A_1, ..., A_n$$
, a table  $T(A_1, ..., A_n)$  and a tuple  $t \in T$ , i.e.,  $t(a_1, ..., a_n)$ 

$$g(T) = \left\{ k \cdot f(t) : t \in T \text{ and } \left| f^{-1}(f(t)) \right| = k \right\}$$

The function g is a multi-set function and  $f^1$  is the inverse function of f. I say that g is the multi-set function generated by f and by the  $f_i$ 's. Further, I say that g(T) is a generalization of table T. This does not mean, however, that the generalization respects the value generalization hierarchy for each attribute in T. To determine whether one table is a generalization with respect to the value generalization hierarchy of each attribute requires analyzing the values themselves.

Let  $DGH_i$  be the domain generalization hierarchies for attributes  $A_{li}$  where  $i=1,...,A_n$  Let  $T_1[A_{ll},...,A_{lAn}]$  and  $T_m[A_{ml},...,A_{mAn}]$  be two tables such that for each  $i:1,...,n, A_{li},A_{mi}\in DGH_i$ . Then, I say table  $T_m$  is a generalization of table  $T_l$ , written  $T_1 \le T_m$ , if and only if there exists a generalization function g such that  $g[\mathsf{T}_l] = \mathsf{T}_m$  and is generated by  $f_i$ 's where:  $\forall t_l \in \mathsf{T}_l$ ,  $a_{li} \leq f_i(a_{li}) = a_{mi}$  and  $f_i : A_{li} \to A_{mi}$ and each  $f_i$  is in the DGH<sub>i</sub> of attribute  $A_{li}$ . From this point forward, I will use the term *generalization* (as a noun) to denote a generalization of a table.

Race E <sub>0</sub>	ZIP Z <sub>0</sub>	Race E <sub>1</sub>	ZIP Z <sub>0</sub>	Race E <sub>1</sub>	ZIP Z <sub>1</sub>	Race E <sub>0</sub>	ZIP Z <sub>2</sub>	Rac E <sub>0</sub>	e	ZIP Z <sub>1</sub>
Black	02138	Person	02138	Person	0213*	Black	021**	Blac	k	0213*
Black	02139	Person	02139	Person	0213*	Black	021**	Blac	ck	0213*
Black	02141	Person	02141	Person	0214*	Black	021**	Blac	ck	0214*
Black	02142	Person	02142	Person	0214*	Black	021**	Blac	ck	0214*
White	02138	Person	02138	Person	0213*	White	021**	Whi	te	0213*
White	02139	Person	02139	Person	0213*	White	021**	Whi	te	0213*
White	02141	Person	02141	Person	0214*	White	021**	Whi	te	0214*
White	02142	Person	02142	Person	0214*	White	021**	Whi	te	0214*
Р	Т	G	Γ <sub>[1,0]</sub>	G	T <sub>[1,1]</sub>	G7	Γ <sub>[0,2]</sub>		G٦	Γ <sub>[0,1]</sub>

Figure 4. Examples of generalized tables for PT.

#### Definition 3. k-anonymity requirement of a generalized table

Let  $PT(A_1,...,A_n)$  be a table,  $Ql_{PT}=\{A_i,...,A_i\}$ , be the quasi-identifier associated with PT where  $\{A_i,...,A_i\} \subseteq \{A_1,...,A_n\}$ ,  $\mathsf{RT}(A_i,...,A_i)$  be a generalization of PT with respect to  $Ql_{PT}$ ,  $t \in RT[Ql_{PT}]$  and  $k_t$  be the integer denoted in g for f(t). RT is said to satisfy k-anonymity for k with respect to  $Ql_{PT}$  if  $\forall t \in RT[Ql_{PT}], k_t \ge k$  and  $k \ge 2$ .

The k-anonymity requirement guarantees each tuple in the generalized table  $RT[Ql_{PT}]$  is indistinguishable from at least k-1 other tuples in table  $RT[Ql_{PT}]$ .

#### Example 3. k-anonymity requirement of a generalized table

Consider PT and its generalized tables in Figure 4 and the hierarchies in Figure 2 and Figure 3.  $GT_{[0,1]}$  and  $GT_{[1,0]}$  satisfy k-anonymity for k=2; and,  $GT_{[0,2]}$  and  $GT_{[1,1]}$  satisfy k-anonymity for k = 2, 3, 4.

The number of different generalizations of a table T, when generalization is enforced at the attribute level, is equal to the number of different combinations of domains that the attributes in the table can assume. Given domain generalization hierarchies  $DGH_i$  for attributes  $A_i$ , i:1,...,n; the number of generalizations, enforced at the attribute level, for table  $T(A_1,...,A_n)$  is:

$$\prod_{i=1}^{n} \left( |DGH_i| + 1 \right)$$
 Equation 1

Clearly, not all such generalizations are equally satisfactory. A generalization whose values result from generalizing each attribute to the highest possible level collapses all tuples in the table to the same list of values. This provides *k*-anonymity using more generalization than needed if a less generalized table exists which satisfies *k*-anonymity. This concept is captured in the following definition.

### Definition 4. k-minimal generalization

Let  $T_l(A_1,...,A_n)$  and  $T_m(A_1,...,A_n)$  be two tables such that  $T_l[Ql_T] \le T_m[Ql_T]$ , where  $Ql_T=\{A_i,...,A_j\}$  is the quasi-identifier associated with the tables and  $\{A_i,...,A_j\} \subseteq \{A_1,...,A_n\}$ .  $T_m$  is said to be a minimal generalization of a table  $T_l$  with respect to a k anonymity requirement over  $Ql_T$  if and only if:

- 1.  $T_m$  satisfies the k-anonymity requirement with respect to  $QI_T$
- 2.  $\forall T_z: T_1 \le T_z, T_z \le T_m, T_z \text{ satisfies the } k\text{-anonymity requirement}$  with respect to  $QI_T \Rightarrow T_z[A_i, \dots, A_j] = T_m[A_i, \dots, A_j]$ .

# Example 4. k-minimal generalization

Figure 4 shows generalizations, enforced at the attribute level, of PT over  $\{Race, ZIP\}$ . Each generalization satisfies k-anonymity for k=2.  $\mathsf{GT}_{[0,1]}$  generalized ZIP one level.  $\mathsf{GT}_{[1,0]}$  generalized Race one level. So,  $\mathsf{GT}_{[1,1]}$ , and  $\mathsf{GT}_{[0,2]}$  did more generalization than necessary.  $\mathsf{GT}_{[0,2]}$  is a generalization of  $\mathsf{GT}_{[0,1]}$ .  $\mathsf{GT}_{[1,1]}$  is a generalization of both  $\mathsf{GT}_{[1,0]}$  and  $\mathsf{GT}_{[0,1]}$ .

So, table  $T_m$ , generalization of  $T_l$ , is k-minimal if it satisfies k-anonymity and there does not exist a generalization of  $T_l$  satisfying k-anonymity of which  $T_m$  is a generalization.

#### 3.3. Minimal distortion of a table

When different k-minimal generalizations exist, preference criteria can be applied to choose a solution among them. A way of preferring one to another is to select one whose information is most useful. Generalizing a table T results in a table T' that typically has less information than T; and so, T' is considered less useful. In order to capture the information loss, I define an information theoretic metric that

reports the amount of distortion in a generalized table.<sup>5</sup> In a cell of a generalized table, the ratio of the domain of the value found in the cell to the height of the attribute's hierarchy reports the amount of generalization and thereby measures the cell's distortion. Precision of a generalized table is then one minus the sum of all cell distortions (normalized by the total number of cells), as defined below.

#### Definition 5. precision metric *Prec*

Let  $PT(A_1,...,A_{Na})$  be a table,  $t_{Pj} \in PT$ ,  $RT(A_1,...,A_{Na})$  be a generalization of PT,  $t_{Pi} \in PT$ , each DGH<sub>A</sub> be the domain generalization hierarchy for attribute A, and f's be generalizations on A. The precision of RT, written Prec(RT), based on generalization and suppression is:

$$Prec(\mathsf{RT}) = 1 - \frac{\sum\limits_{i=1}^{N_A} \sum\limits_{j=1}^{N} \frac{h}{\left|\mathsf{DGH}_{Ai}\right|}}{\left|\mathsf{PT}\right| \bullet \left|N_A\right|} \quad where \ f_1\left(\dots f_h\left(t_{p_j}\left[A_i\right]\right)\dots\right) = t_{R_j}\left[A_i\right]$$

# Example 5. precision metric

In the case where PT = RT, each value is in the ground domain so each h = 0; therefore, Prec(RT) = 1. Conversely, in the case where each value in RT is the maximal element of its hierarchy, each  $h = |DGH_{Ai}|$ ; and so, Prec(RT) = 0.

#### Example 6. precision metric

Using the hierarchies in Figure 2 and Figure 3, the precision of the generalizations of PT shown in Figure 4 are:  $Prec(GT_{[1,0]}) = 0.75$ ;  $Prec(GT_{[1,1]}) = 0.58$ ;  $Prec(GT_{[0,2]}) = 0.67$ ; and,  $Prec(GT_{[0,1]}) = 0.83$ . Each of these satisfy k-anonymity for k=2, but  $GT_{[0,1]}$  does so with the least distortion. Notice  $GT_{[1,0]}$  and  $GT_{[0,1]}$  each generalize values up one level, but because  $|DGH_{Bace}| = 2$  and  $|DGH_{ZIP}| = 3$ ,  $Prec(GT_{[0,1]}) > Prec(GT_{[1,0]})$ .

Generalizations based on attributes with taller generalization hierarchies typically maintain precision better than generalizations based on attributes with shorter hierarchies. Further, hierarchies with different heights can provide different Prec measures for the same table. So, the construction of generalization hierarchies is part of the preference criteria. Prec best measures the quality of the data when the set of hierarchies used contain only values semantically useful. There is no need to arbitrarily increase the heights of hierarchies solely to prefer one attribute to another. Instead, weights can be assigned to attributes in Prec to make the preference explicit [9].

<sup>&</sup>lt;sup>5</sup> While entropy is the classical measure used in information theory to characterize the purity of data [5], a metric based on the semantics of generalization can be more discriminating than the direct comparison of the encoding lengths of the values stored in the table.

As stated earlier, not all k-minimal generalizations are equally distorted and preference can be based on the k-minimal generalization having the most precision. This concept is captured by the following definition.

#### Definition 6. k-minimal distortion

Let  $T_l(A_1,...,A_n)$  and  $T_m(A_1,...,A_n)$  be two tables such that  $T_l[QI_T] \leq T_m[QI_T]$ , where  $QI_T=\{A_i,...,A_j\}$  is the quasi-identifier associated with the tables and  $\{A_i,...,A_j\} \subseteq \{A_1,...,A_n\}$  and  $\forall x=i,...,j$ ,  $DGH_{Ax}$  are domain generalization hierarchies for  $QI_T$ .  $T_m$  is said to be a minimal distortion of a table  $T_l$  with respect to a k anonymity requirement over  $QI_T$  if and only if:

- 1.  $T_m$  satisfies the k-anonymity requirement with respect to  $QI_T$
- 2.  $\forall T_z: Prec(T_i) \ge Prec(T_z), Prec(T_z) \ge Prec(T_m), T_z \text{ satisfies the } k$ anonymity requirement with respect to  $Ql_T \Rightarrow T_z[A_i,...,A_j] = T_m[A_i,...,A_j].$

Consider the values reported in Example 6 about the tables in Figure 4. Only  $GT_{[0,1]}$  is a k-minimal distortion of PT. A k-minimal distortion is specific to a table, a quasi-identifier, a set of domain generalization hierarchies for the attributes of the quasi-identifier, and Prec (or a weighted Prec).

It is trivial to see that a table that satisfies k-anonymity has a unique k-minimal distortion, which is itself. It is also easy to see that a generalized table RT that is a k-minimal distortion of table PT is also a k-minimal generalization of PT, as stated in the following theorem.

#### Theorem 2

Given tables  $T_1$  and  $T_m$  such that  $T_1 \le T_m$  and  $T_m$  satisfies k-anonymity.  $T_m$  is a k-minimal distortion of  $T_1 \Rightarrow T_m$  is k-minimal generalization of  $T_1$ .

# 3.4. Algorithm for finding a minimal generalization with minimal distortion

The algorithm presented in this section combines these formal definitions into a theoretical model against which real-world systems will be compared.

Figure 5 presents an algorithm, called MinGen, which, given a table  $PT(A_x,...,A_y)$ , a quasi-identifier  $QI=\{A_1,...,A_n\}$ , where  $\{A_1,...,A_n\} \subseteq \{A_x,...,A_y\}$ , a k-anonymity constraint, and domain generalization hierarchies  $DGH_{Ai}$ , produces a table **MGT** which is a k-minimal distortion of PT[QI]. It assumes that k < |PT|, which is a necessary condition for the existence of a k-minimal generalization.

The steps of the MinGen algorithm are straightforward. [step 1] Determine if the original table, PT, itself satisfies the *k*-anonymity requirement; and if so, it is the *k*-minimal distortion. In all other cases execute step 2. [step 2.1] Store the set of all possible generalizations of PT over QI into *allgens*. [step 2.2] Store those generalizations from *allgens* that satisfy the *k*-anonymity requirement into

[step 2.3] Store the k-minimal distortions (based on Prec) from protected. protected into MGT. It is guaranteed that  $|MGT| \ge 1$ . [step 2.4] Finally, the function preferred() returns a single k-minimal distortion from MGT based on user-defined specifications<sup>6</sup>.

```
Input: Private Table PT; quasi-identifier QI = (A_1, ..., A_n),
          k constraint; domain generalization hierarchies
          DGH_{Ai}, where i=1,...,n, and preferred() specifications.
Output: MGT, a minimal distortion of PT[QI] with respect to k
          chosen according to the preference specifications
Assumes: |PT| \ge k
Method:
       if PT[QI] satisfies k-anonymity requirement with respect to k then do
        1.1. MGT \leftarrow \{ PT \} // PT  is the solution
  2.
       else do
        2.1. allgen \leftarrow \{T_i : T_i \text{ is a generalization of } PT \text{ over } QI\}
        2.2. protected \leftarrow \{T_i : T_i \in allgen \land T_i \text{ satisfies } k\text{-anonymity of } k\}
               MGT \leftarrow \{T_i : T_i \in protected \land there does not exist T_z \in protected\}
               such that Prec(T_z) > Prec(T_i) }
               MGT ← preferred(MGT) // select the preferred solution
        return MGT
```

Figure 5. Preferred Minimal Generalization (MinGen) Algorithm.

#### Example 7. MinGen produces k-minimal distortions

Let the hierarchies in Figure 2, Figure 3 and Figure 6 for the quasi-identifier  $Ql=\{Race, BirthDate, Gender, ZIP\}$ , the table PT in Figure 7, and a kanonymity constraint of k=2 be provided to MinGen. After step 2.3 of MinGen,  $MGT = \{GT\}$ , where GT is shown in Figure 7. GT is a k-minimal distortion of PT over QI and a k-minimal generalization of PT over QI.

It can be proved that a generalization of a table T over a quasi-identifier QI, that satisfies a given k-anonymity requirement, and has the least amount of distortion of all possible generalizations of T over QI, is a k-minimal distortion of T over QI with respect to Prec. From Theorem 2, the solution is also a k-minimal generalization of T over Ql.

With respect to complexity, MinGen makes no claim to be efficient. If generalization is enforced at the attribute level, the number of possible

<sup>&</sup>lt;sup>6</sup> The preferred() function returns only one table as a solution. The single solution requirement is a necessary condition because the chosen solution becomes part of the join of external information against which subsequent linking must be protected.

generalizations, |allgens|, is expressed in Equation 1. If generalization is enforced at the cell level,  $|allgens| = \prod_{i=1}^{n} (|DGH_{Ai}| + 1)^{|PT|}$ . Equation 2

Clearly, an exhaustive search of all possible generalizations is impractical even on modest sized tables. So, how do real-world systems find solutions in real-time?

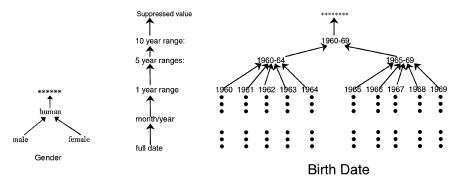


Figure 6. Value generalization hierarchies for {Gender, Birth Date} with suppression.

Race	BirthDate	Gender	ZIP	Problem
black	9/20/1965	male	02141	short of breath
black	2/14/1965	male	02141	chest pain
black	10/23/1965	female	02138	painful eye
black	8/24/1965	female	02138	wheezing
black	11/7/1964	female	02138	obesity
black	12/1/1964	female	02138	chest pain
white	10/23/1964	male	02138	short of breath
white	3/15/1965	female	02139	hypertension
white	8/13/1964	male	02139	obesity
white	5/5/1964	male	02139	fever
white	2/13/1967	male	02138	vomiting
white	3/21/1967	male	02138	back pain
		PT	Prec=1	1.00

Race	BirthDate	Gender	ZIP	Problem
black	1965	male	02141	short of breath
black	1965	male	02141	chest pain
person	1965	female	0213*	painful eye
person	1965	female	0213*	wheezing
black	1964	female	02138	obesity
black	1964	female	02138	chest pain
white	1960-69	male	02138	short of breath
person	1965	female	0213*	hypertension
white	1964	male	02139	obesity
white	1964	male	02139	fever
white	1960-69	male	02138	vomiting
white	1960-69	male	02138	back pain
		GT	Prec=0	0.90

Figure 7. k-minimal distortion for PT where k = 2.

#### 4. Real-World Results

Here are two real-world systems that seek to provide k-anonymity protection using generalization and suppression. They are: (1) my Datafly<sup>7</sup> Systems [5]; and, (2) Statistics Netherlands'  $\mu$ -Argus System [6]. This section shows that Datafly can over distort the data and that  $\mu$ -Argus can fail to provide adequate protection.

<sup>&</sup>lt;sup>7</sup> The systems Datafly and Datafly II refer to two kindred algorithms. The differences between them do not substantially alter the findings reported herein, so in this writing, the term Datafly refers to a simplified abstraction of these algorithms known as the core Datafly algorithm [9].

#### 4.1. The Datafly system

Here is a summary of the setting in which the core Datafly algorithm operates. The data holder (1) declares specific attributes and tuples in the original private table (PT) as being eligible for release. The data holder also (2) groups a subset of attributes of PT into one or more quasi-identifiers (Ql<sub>i</sub>) and assigns (3) a weight from 0 to 1 to each attribute within each Qli that specifies the likelihood the attribute will be used for linking; a 0 value means not likely and a value of 1 means highly probable. The data holder (4) specifies a minimum anonymity level that computes to a value for k. Finally, (5) a weight from 0 to 1 is assigned to each attribute within each Ql<sub>i</sub> to state a preference of which attributes to distort; a 0 value means the recipient of the data would prefer the values not to be changed and a value of 1 means maximum distortion could be tolerated. For convenience in this discussion, I remove many of the finer features. I consider a single quasiidentifier, where all attributes of the quasi-identifier have equal preference and an equal likelihood for linking so the weights can be considered as not being present.

Figure 8 presents the core Datafly algorithm, which, given a table  $PT(A_x,...,A_y)$ , a quasi-identifier  $QI=\{A_1,...,A_n\}$ , where  $\{A_1,...,A_n\}\subseteq\{A_x,...,A_y\}$ , a k-anonymity constraint, and domain generalization hierarchies DGHAi, produces a table MGT which is a generalization of PT[QI] that satisfies k-anonymity. It assumes that k < |PT|, which is a necessary condition for satisfying k-anonymity.

Private Table **PT**; quasi-identifier  $QI = (A_1, ..., A_n)$ , k constraint; hierarchies DGH<sub>Ai</sub>, where i=1,...,n.

Output: MGT, a generalization of PT[QI] with respect to k

Assumes:  $|PT| \ge k$ 

#### Method:

- freq  $\leftarrow$  a frequency list contains distinct sequences of values of PT[QI], along with the number of occurrences of each sequence.
- while there exists sequences in freq occurring less than k times that account for more than k tuples do
  - **let**  $A_i$  be attribute in freq having the most number of distinct values
  - 2.2. freq  $\leftarrow$  generalize the values of  $A_i$  in freq
- 3. freq  $\leftarrow$  suppress sequences in freq occurring less than k times.
- freq  $\leftarrow$  enforce k requirement on suppressed tuples in freq.
- **Return MGT** ← construct table from freq

Figure 8. Core Datafly algorithm.

The core Datafly algorithm has few steps. Step 1 constructs freq, which is a frequency list containing distinct sequences of values from PT[QI], along with the number of occurrences of each sequence. Each sequence in freq represents one or more tuples in a table. Step 2.1 uses a heuristic to guide generalization. The attribute having the most number of distinct values in freq is generalized. Generalization continues until there remains k or fewer tuples having distinct sequences in freq. Step 3 suppresses any sequences of freq occurring less than k times. Complimentary suppression is performed in step 4 so that the number of suppressed tuples satisfies the k requirement. Finally, step 5 produces a table MGT, based on freq such that the values stored as a sequence in freq appear as tuple(s) in MGT replicated in accordance to the stored frequency.

Let the hierarchies in Figure 2, Figure 3 and Figure 6 for the quasi-identifier  $Ql=\{Race, BirthDate, Gender, ZIP\}$ , the table PT in Figure 7, and a k-anonymity constraint of k=2 be provided to Datafly. Figure 9A shows the contents of freq after step 1. Birthdate has the most number of distinct values (12) so its values are generalized. Figure 9B shows the contents of freq after step 2. Tuples t7 and t8 will be suppressed. Table MGT in Figure 10 is the final result. MGT[QI] satisfies k-anonymity for k=2 with Prec(MGT[QI]) = 0.75. However, from Example 7 and Figure 7, GT is a k-minimal distortion for PT with Prec(GT[QI])=0.90. So, Datafly distorted the results more than needed.

				_	
Race	BirthDate	Gender	ZIP	#occurs	
black	9/20/65	male	02141	1	t1
black	2/14/65	male	02141	1	t2
black	10/23/65	female	02138	1	t3
black	8/24/65	female	02138	1	t4
black	11/7/64	female	02138	1	t5
black	12/1/64	female	02138	1	t6
white	10/23/64	male	02138	1	t7
white	3/15/65	female	02139	1	t8
white	8/13/64	male	02139	1	t9
white	5/5/64	male	02139	1	t10
white	2/13/67	male	02138	1	t11
white	3/21/67	male	02138	1	t12
2	12	2	3		

Race	BirthDate	Gender	ZIP	#occurs	
black	1965	male	02141	2	t1,t2
black	1965	female	02138	2	t3, t4
black	1964	female	02138	2	t5, t6
white	1964	male	02138	1	t7
white	1965	female	02139	1	t8
white	1964	male	02139	2	t9, t10
white	1967	male	02138	2	t11, t12
2	2 3	3 2	: 3	- "	
	R				

Figure 9. Intermediate stages of the core Datafly algorithm.

Race	BirthDate	Gender	ZIP	Problem
black	1965	male	02141	short of breath
black	1965	male	02141	chest pain
black	1965	female	02138	painful eye
black	1965	female	02138	wheezing
black	1964	female	02138	obesity
black	1964	female	02138	chest pain
white	1964	male	02139	obesity
white	1964	male	02139	fever
white	1967	male	02138	vomiting
white	1967	male	02138	back pain
				_

Figure 10. Table MGT resulting from Datafly, k = 2,  $Ql = \{Race, Birthdate, Gender, ZIP\}$ .

The core Datafly algorithm does not necessarily provide k-minimal generalizations or k-minimal distortions, even though it can be proved that its solutions always

satisfy k-anonymity. One of the problems is that Datafly makes crude decisions – generalizing all values associated with an attribute and suppressing all values within a tuple. MinGen makes decisions at the cell-level and by doing so, can provide results with more precision. Another problem is the heuristic that selects the attribute with the greater number of distinct values as the one to generalize. This may be computationally efficient, but can be shown to perform unnecessary generalization. In summary, Datafly produces generalizations that satisfy kanonymity, but such generalizations may not be k-minimal distortions.

#### 4.2. The \(\mu\)-Argus system

In  $\mu$ -Argus, the data holder provides a value for k and specifies which attributes are sensitive by assigning a value between 0 and 3 to each attribute. These correspond to "not identifying," "most identifying," "more identifying," and "identifying," respectively. µ-Argus then identifies rare and therefore unsafe combinations by testing 2- and 3-combinations of attributes. Unsafe combinations are eliminated by generalizing attributes within the combination and by cell suppression. Rather than removing entire tuples, μ-Argus suppresses values at the cell-level. The resulting data typically contain all the tuples and attributes of the original data, though values may be missing in some cell locations.

Figure 11 presents the  $\mu$ -Argus algorithm. Given a table  $PT(A_x,...,A_v)$ , a quasi-identifier  $Ql = \{A_1, ..., A_n\}$ , where  $\{A_1, ..., A_n\} \subseteq \{A_x, ..., A_v\}$ , disjoint subsets of QI known as *Identifying*, *More*, and *Most* where QI = *Identifying*  $\cup$  *More*  $\cup$  *Most*, a k-anonymity constraint, and domain generalization hierarchies DGH<sub>Ai</sub>, μ-Argus produces a table MT which is a generalization of PT[Ql].

The basic steps of the  $\mu$ -Argus algorithm are provided in Figure 11. This algorithm results from my reverse engineering an implementation [9]. Shortcomings of the actual  $\mu$ -Argus implementation were found. So, results from In general, the constructed  $\mu$ -Argus algorithm generates both are reported. solutions that are better protected than those released by the actual program.

The program begins in step 1 by constructing freq, which is a frequency list containing distinct sequences of values from PT[QI], along with the number of occurrences of each sequence. In step 2, the values of each attribute are automatically generalized until each value associated with an attribute in the quasi-identifier QI appears at least k times. This is a necessary condition for k-In step 3, the program automatically tests anonymity (see Theorem 1). combinations of attributes to identify those combinations of attributes whose assigned values in combination do not appear at least k times; these combinations are stored in outliers. Afterwards, the data holder, in steps 4 and 5, decides whether to generalize an attribute in QI that has values in outliers and if so, selects the attribute to generalize. Finally, in step 6,  $\mu$ -Argus automatically suppresses a value from each combination in *outliers*. Precedence is given to the value occurring most often in order to reduce the total number of suppressions.

Input: Private Table PT; quasi-identifier  $QI = (A_1, ..., A_n)$ , disjoint subsets of QI known as *Identifying*, *More*, and *Most* where  $QI = Identifying \cup More \cup Most$ , k constraint; domain generalization hierarchies  $DGH_{Ai}$ , where i=1,...,n.

Output: MT containing a generalization of PT[QI]

**Assumes:** |PT| ≥ k

Method:

- freq ← a frequency list containing distinct sequences of values of PT[QI], along with the number of occurrences of each sequence.
- 2. Generalize each  $A_i \in \mathbb{Q}$  in freq until its assigned values satisfy k.
- 3. Test 2- and 3- combinations of *Identifying*, *More* and *Most* and **let** *outliers* store those cell combinations not having *k* occurrences.
- 4. Data holder decides whether to generalize an  $A_j \in QI$  based on *outliers* and if so, identifies the  $A_i$  to generalize. freq contains the generalized result.
- 5. **Repeat** steps 3 and 4 until the data holder no longer elects to generalize.
- 6. Automatically suppress a value having a combination in *outliers*, where precedence is given to the value occurring in the most number of combinations of *outliers*.

Figure 11. μ-Argus algorithm.

One shortcoming of the actual  $\mu$ -Argus implementation appears in step 3 of Figure 11. The actual program does not test all 2- and 3- combinations; this may be a programming error. Figure 12 reports which combinations  $\mu$ -Argus tests. Six combinations (not listed) are not tested at all. It is easy to have tables in which values appear in combinations not examined by  $\mu$ -Argus.

Combinations Always Tested	
Identifying $\times$ More $\times$ Most, Identifying $\times$ Most $\times$ Most, Most $\times$ Most, Identifying $\times$ More, Identifying $\times$ Most, More $\times$ Most, Most $\times$ Most	
Combinations Tested only if   Identifying   > 1	
$More \times More \times Most$ , $Most \times Most \times More$ , $More \times More$	

Figure 12. Combinations of *More*, *Most*, *Identifying* tested by  $\mu$ -Argus.

Let the hierarchies in Figure 2, Figure 3 and Figure 6 for the quasi-identifier  $Ql=\{Race, BirthDate, Gender, ZIP\}$  where  $Most=\{BirthDate\}$ ,  $More=\{Gender, ZIP\}$  and  $Identifying=\{Race\}$ , the table PT in Figure 7, and a k-anonymity constraint of k=2 be provided to  $\mu$ -Argus. In Figure 13, V shows the result of testing  $Most \times More$  at step, and freq is updated to show  $\{BirthDate, ZIP\}$  for t8 did not satisfy k. Figure 14 shows freq before step 6; the values to be suppressed are underlined. Table MT in Figure 15 is the final result from the  $\mu$ -Argus

algorithm provided in Figure 11. The k-anonymity requirement is not enforced on suppressed values, making it vulnerable to linking as well as to an inference attack using summary data<sup>8</sup>. For example, knowing the total number of men and women allows the suppressed values for Gender to be inferred.

				Race	Birth	Sex	ZIP	occurs	sid	outliers
Birth ZIP of	ccurs	sid	outliers	black	1965	male	02141	2	{t1,t2}	{}
1965 02141	2		Outhers	black	1965	female	02138	2	{t3,t4}	{}
1965 02138	2	{t1,t2} {t3,t4}	t) n	black	1964	female	02138	2	{t5,t6}	{}
1964 02138	3	{t5,t6,t7}	υ υ	white	1964	male	02138	1	{t7}	()
1965 02139	1	(t8)	u B	white	1965	female	02139	1	{t8}	{{birth,zip}}
1964 02139	2	{t9,t10}	{}	white	1964	male	02139	2	{t9,t10}	{}
1967 02138	2	{t11,t12}	ő	white	1967	male	02138	2	{t11,t12}	i)
/	V						fre	eq		

Figure 13. Most × More combination test and resulting freq.

Race	Birth	Sex	ZIP	occurs sid	outliers
black	1965	male	02141	2 {t1,t2}	0
black	1965	female	02138	2 {t3,t4}	Ö
black	1964	female	02138	2 {t5,t6}	0
					{{ <b>birth</b> ,sex,zip},
white	1964	male	02138	1 {t7}	{race, <b>birth</b> ,zip}}
					{{birth,zip}, {sex,zip},
					{birth,sex,zip},
					{race, <b>birth</b> , <b>sex</b> },
					{race, birth, zip}, {race, sex},
white	1965	female	02139	1 {t8}	{race, <b>birth</b> }}
white	1964	male	02139	2 {t9,t10}	0
white	1967	male	02138	2 {t11,t12}	Ö

Figure 14. freq before suppression.

id	Race	BirthDate	Gender	ZIP	id	Race	BirthDate	Gender	ZIP
t1	black	1965	male	02141	t1	black	1965	male	021
t2	black	1965	male	02141	t2	black	1965	male	021
t3	black	1965	female	02138	t3	black	1965	female	021
t4	black	1965	female	02138	t4	black	1965	female	021
t5	black	1964	female	02138	t5	black	1964	female	021
t6	black	1964	female	02138	t6	black	1964	female	021
t7	white		male	02138	t7	white	1964	male	021
t8	white			02139	t8	white		female	021
t9	white	1964	male	02139	t9		1964	male	021
t10	white	1964	male	02139	t10	white	1964	male	021
t11	white	1967	male	02138	t11	white	1967	male	021
t12	white	1967	male	02138	t12		1967	male	021
		MT					MT actua		

Figure 15. Results from the  $\mu$ -Argus algorithm and from the program.

The actual μ-Argus program provides MTactual shown in Figure 15. The tuple identified as t7 is ["white", "1964", "male", "02138"], which is unique over MTactual[QI]. Therefore, MTactual does not satisfy the requirement for k=2.

<sup>&</sup>lt;sup>8</sup> See [3] for a detailed discussion of attacks on k-anonymity.

A shortcoming of  $\mu$ -Argus stems from not examining all combinations of the attributes in the quasi-identifier. Only 2- and 3- combinations are examined. There may exist 4-combinations or larger that are unique. Directly extending  $\mu$ -Argus to compute on all combinations loses its computational efficiency. So, generalizations from  $\mu$ -Argus may not always satisfy k-anonymity, even though all generalizations from Datafly do satisfy k-anonymity.

Both algorithms may provide generalizations that are not k-minimal distortions because they both enforce generalization at the attribute level. This renders crude Prec measures. There may exist values in the table that when generalized at the cell level, satisfy k without modifying all values in the attribute. In summary, more work is needed to correct these heuristic-based approaches.

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