Generating Realistic Test Datasets for Duplicate Detection at Scale Using Historical Voter Data

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ABSTRACT
The detection of duplicates is an essential task in data cleaning and integration and has steadily gained importance especially for companies that need to process and integrate large volumes of potentially unclean data on a daily basis. To evaluate the quality and performance of duplicate detection algorithms, labeled test data is required that provide information on the contained duplicates. Current approaches for generating test data, however, are either not scalable (and therefore limited to small datasets) or not able to generate realistic data values and errors, especially outdated values. In this paper, we propose a scheme for generating test datasets that addresses both these issues and present a test dataset generated with it. Our approach relies on using historical data from the North Carolina voter registration which (i) is realistic as it contains actual voter data and (ii) facilitates generating realistic duplicates through the fact that current data values were collected at every election through manually filled out applications. The generated test dataset comprises more than 120 million records with up to 90 attribute values each. To the best of our knowledge, we are the first who provide realistic test data for duplicate detection at this scale.

1 INTRODUCTION
Duplicates are data records (e.g., tuples in the relational case) that refer to the same real-world object. They can result from errors in data management, but also occur because separately developed data sources overlap in their universes of discourse (e.g., many actors and movies are stored in both IMDB\(^1\) and TMDb\(^2\)). The detection of duplicates is an important task in data cleaning \([9, 24]\) and integration \([5, 6]\). Detecting duplicates is quite simple when they are exact, i.e. they agree in all of their values. However, it can be extremely difficult if some of their values disagree due to typos, phonetic or transformation errors, heterogeneous forms of presentation as well as missing or outdated values \([11]\).

The challenge of detecting such so-called fuzzy duplicates has opened up its own field of research and has since been studied intensively \([3, 8, 19, 31]\). However, the best approach to find them strongly depends on (i) the considered domain (e.g., movies, persons, or proteins), (ii) the characteristics of the given data (e.g., volume, data model and heterogeneity), and (iii) the quality and cost requirements of the user (e.g., good results vs. short runtimes and recall vs. precision). Due to the resulting diversity of use cases, none of the existing algorithms has turned out to be a generally applicable and superior solution. Instead, in every use case, it remains a difficult (and expensive) task to choose and configure them so that they provide adequate results.

Such a configuration process requires the evaluation and comparison of different algorithms and parameter settings. This in turn requires test datasets that do not only provide a gold standard (a.k.a. ground truth) labeling the dataset’s duplicates \([17]\), but resemble the required real-life properties as well as possible. Current approaches to test data generation either (i) struggle with the generation of realistic data values and errors (especially outdated values), (ii) cannot guarantee the soundness of the gold standard\(^3\), or (iii) scale badly and thus can only be used to generate small datasets. However, realistic values and errors as well as correctly labeled duplicates are an important prerequisite for a test dataset. Moreover, in times of big data many duplicate detection algorithms focus on scalability (e.g., \([10, 13, 20, 21, 28]\)) so that an evaluation of their key functionalities requires large test datasets with millions of records.

Using a historical dataset to generate test data seems to be a straightforward solution to some of the aforementioned problems, because the mapping between records and real-world objects is part of the data so that the duplicates are already labeled. Thus, it scales much better than, for example, labeling the duplicates in an unclean dataset manually. In addition, historical datasets are perfectly suited to generate outdated values, because these values are an inherent part of the data. One of these historical datasets is provided by the State of North Carolina \([29]\). This dataset contains information on voters registered to the individual elections and—at the time of our study—consisted of 44 snapshots covering a time period of 15 years with a total number of over 500 million records and a large schema with 90 attributes. These numbers make it a perfect candidate to evaluate the suitability of the aforementioned idea, because the large number of records allows us to generate a test dataset of large size and the large time span provides us many outdated values (even more than one for the same object property). Furthermore, since voters often have to re-register at regular intervals by manually filled out forms\(^4\), the registration data contain typos, values confused between attributes, heterogeneous forms of presentation and missing values which makes this dataset particularly useful for

\(^1\)Internet Movie Database: https://www.imdb.com
\(^2\)The Movie Database: https://www.themoviedb.org
\(^3\)To clearly distinguish between errors in the duplicate labels and the operational data, we use the term soundness w.r.t. the correctness of the gold standard.
\(^4\)https://dl.ncsbe.gov/?prefix=Voter_Registration/
the generation of fuzzy duplicates. Finally, its large size gives us the opportunity to customize the test data to different user requirements by selecting a suitable subset of all records (the more data available, the more flexible the selection process). However, the big amount of redundant data as well as the ongoing publication of new snapshots also pose some challenges to the generation process making it a non-trivial task.

The contributions of this paper can be summarized as:

(i) A comprehensive list of desiderata for test datasets for duplicate detection.

(ii) An approach for generating and storing test data based on a historical voter register from North Carolina.

(iii) A realistic test dataset generated with our approach.

(iv) An extensive experimental evaluation to analyze the quality and prove the usability of the generated test dataset.

The generated dataset is available online\(^3\). It will help other researchers to evaluate their algorithms (such as runtime behavior or robustness) against a varying number of data errors) and to compare with them with those of other research projects. It is particularly valuable to the research community through a combination of properties that is unique to the best of our knowledge:

- It contains more than 120 million records and 640 million duplicate pairs making it suitable to evaluate duplicate detection algorithms at scale.
- It contains real-life errors of various types including typos, abbreviations, phonetrical errors and outdated values.
- Its large size qualifies it to customize the test data to different user requirements without losing necessary volume.
- It provides precalculated plausibility and heterogeneity scores, which support the user to remove (or repair) potentially unsound duplicate clusters and adapt the data's heterogeneity to her own requirements, and
- It provides meta information that allows the user to reproduce experiments using previous versions of this dataset.

The rest of this paper is structured as follows: In Section 2, we describe the input to our study, i.e., the voter registration dataset from North Carolina. Thereafter, in Section 3, we discuss several aspects affecting the test data's quality, usability and reproducibility. In Sections 4 and 5, we describe our approach for using the historical voter data for test data generation. This includes the generation process, but also the storage of the resulting test dataset as well as mechanisms for guaranteeing reproducibility. In Section 6 we present an experimental study that evaluates the quality and usability of the generated test dataset. Finally, we discuss related work in Section 7 before we conclude the paper and give an outlook on future research in Section 8.

2 NORTH CAROLINA VOTER REGISTER

The historical voter registration dataset from North Carolina [30] (short NC) was created and is still maintained by the State Board of Elections in NC according to the Help America Vote Act (HAVA) of 2002. This data set is available for non-commercial purposes only and may be used by the public, state residents, registered voters, non-profits, researchers, political committees and candidates [16]. In NC, different statutory limitations exist that exclude following and may be used by the public, state residents, registered voters, in NC according to the.

<table>
<thead>
<tr>
<th>year</th>
<th>#snapshots</th>
<th>total records</th>
<th>new records</th>
<th>new objects</th>
<th>rate of new objects</th>
</tr>
</thead>
<tbody>
<tr>
<td>2008</td>
<td>1</td>
<td>9.7 M</td>
<td>9.7 M</td>
<td>9.4 M</td>
<td>100%</td>
</tr>
<tr>
<td>2009</td>
<td>1</td>
<td>9.7 M</td>
<td>0.7 M</td>
<td>37 K</td>
<td>6.8%</td>
</tr>
<tr>
<td>2010</td>
<td>2</td>
<td>20.2 M</td>
<td>13.1 M</td>
<td>189 K</td>
<td>64.9%</td>
</tr>
<tr>
<td>2011</td>
<td>4</td>
<td>10.3 M</td>
<td>2.3 M</td>
<td>225 K</td>
<td>22.2%</td>
</tr>
<tr>
<td>2012</td>
<td>2</td>
<td>41.8 M</td>
<td>19.9 M</td>
<td>820 K</td>
<td>47.6%</td>
</tr>
<tr>
<td>2013</td>
<td>1</td>
<td>11.4 M</td>
<td>11.1 M</td>
<td>41 K</td>
<td>97.1%</td>
</tr>
<tr>
<td>2014</td>
<td>4</td>
<td>4.3 M</td>
<td>7.5 M</td>
<td>432 K</td>
<td>15.8%</td>
</tr>
<tr>
<td>2015</td>
<td>4</td>
<td>49.0 M</td>
<td>6.6 M</td>
<td>223 K</td>
<td>13.5%</td>
</tr>
<tr>
<td>2016</td>
<td>4</td>
<td>50.9 M</td>
<td>7.7 M</td>
<td>587 K</td>
<td>15.1%</td>
</tr>
<tr>
<td>2017</td>
<td>4</td>
<td>54.1 M</td>
<td>3.6 M</td>
<td>245 K</td>
<td>6.7%</td>
</tr>
<tr>
<td>2018</td>
<td>3</td>
<td>41.7 M</td>
<td>23.7 M</td>
<td>374 K</td>
<td>56.9%</td>
</tr>
<tr>
<td>2019</td>
<td>7</td>
<td>99.8 M</td>
<td>5.3 M</td>
<td>354 K</td>
<td>5.5%</td>
</tr>
<tr>
<td>2020</td>
<td>4</td>
<td>60.8 M</td>
<td>8.0 M</td>
<td>596 K</td>
<td>13.1%</td>
</tr>
<tr>
<td>total</td>
<td>40</td>
<td>506.7 M</td>
<td>119.3 M</td>
<td>13.49 M</td>
<td>23.5%</td>
</tr>
</tbody>
</table>

\(^3\)https://vsis-www.informatik.uni-hamburg.de/download/ncvoter_testdata/(user = `ncvoter`, password = `5k][fj3`)"
are represented by more than one record. A closer look revealed that at most only one of them has not the voter status removed (and hence is not outdated). This means that every snapshot already corresponds to a historical dataset.

3 TEST DATA DESIDERATA

Before we describe in which way we used the NC voter register to generate test data in Section 4, we will take a closer look on the desired properties of such a test dataset. A suitable test dataset has to ensure a high

- **quality**, i.e. the test data should enable meaningful evaluation results,
- **usability**, i.e. the user should be able to customize the test data according to her requirements, and
- **reproducibility**, i.e. the user should be able to reproduce the results of past evaluations that used previous versions of this dataset in order to achieve adequate comparability.

In the rest of this section, we will discuss these requirements and the problems that are related with them in more detail. The way we handled them in our generation process will be described in the remaining course of this paper.

3.1 Quality

A test dataset is of good quality if its gold standard is sound, its data contain realistic errors of different types and it contains only few exact duplicates.

3.1.1 Soundness of the Gold Standard. A test dataset for duplicate detection consists of a set of data records and the corresponding gold standard that specifies the duplicate status between the individual records. While errors in the actual data are quite desirable (see Section 3.1.2), it is extremely important that the gold standard is sound, because even a small number of incorrectly labeled record pairs (false positives and false negatives) can render evaluation results completely useless.

In a perfect world, the mappings between the voter records and actual voters are sound. However, almost no dataset is free of errors. Thus, it make sense to perform a soundness check on the test data because, as we illustrate in Figure 3, there may be clusters whose records do not seem to represent the same voter although they share the same NCID. Marking those clusters allows the user to remove or repair them before using the test dataset. Since we often cannot distinguish between sound and unsound clusters with absolute confidence, it does not seem wise to use a Boolean flag as a marker, but to compute similarities which reflect a kind of likelihood that these clusters are sound (i.e., all their records represent the same voter). The user can then use these similarities to decide which risk she wants to take to include unsound clusters into her test data. We refer to this similarity as plausibility in the rest of this paper and discuss a calculation of plausibility scores for the NC test dataset in Section 6.2.

3.1.2 Error Diversity. The results of an evaluation with a test dataset are only representative if this dataset contains real-life (or at least realistic) data values and errors. In our case, both are real because they originate from a real-life dataset. Moreover, users want to evaluate algorithms that should later be applied to error-prone data. This requires that the test dataset contains errors of various kinds and not only outdated values. This includes typos, abbreviations, invalid values, inconsistencies and different forms of representation. In other words a test dataset of high quality has to contain several problems of data quality.

3.1.3 Amount of Exact Duplicates. Another aspect that affects the usefulness of evaluation results is the number of exact duplicates contained in the test data. The detection of such duplicates is rather simple and every duplicate detection algorithm – no matter how primitive – should be able to detect them. Thus if this number dominates the number of fuzzy duplicates by far, an accurate detection of the latter becomes less relevant in order to achieve a good evaluation result. For example, if 90% of all duplicate pairs are exact, even the most primitive algorithm would achieve a recall of 0.9 or higher if it is able to compare values on equivalence. Moreover, an algorithm that classifies only the exact duplicates as such and all other pairs as non-duplicates (precision is 1.0) would even achieve a $F_1$-score of 0.9 which is a pretty good result. However those algorithms are completely useless when it comes to real-life use cases where fuzzy duplicates need to be detected. While this aspect is of little relevance in many approaches to test data generation (the number of exact duplicates is usually very small there), it is of great importance when using historical data, since many of the given snapshots overlap to a large extent, so that their combination leads to many exact duplicates. As we will see in Section 4, by simply combining the individual snapshots of the NC voter register we produced a relative amount of exact duplicates of over 90%.

The actual definition of an exact duplicate pair is that both records share the exact same value in every attribute. However, in the case of the NC voter data, solely removing those duplicates which are completely identical does not solve this problem, because often many of the remaining duplicate records only differ in some minor aspects, such as:

- **Meta Data Attributes**: Many duplicate records only differ in some date values, such as snapshot or registration date, that are less relevant for the duplicate detection process.
- **Time-related Attributes**: The voters’ age values increase by one every year and thus cause that some duplicates are no longer exact, although none of the other characteristics of the corresponding person changed.
- **Whitespaces**: Many values contain leading and trailing whitespaces that are simply to detect and remove by trimming all data values in an initial preparation step.

We describe how we addressed this problem in Section 4.

3.2 Usability

Since we aim to provide test data for duplicate detection at scale, the resulting dataset should contain several million records. Besides size, the requirements of the individual users can vary from one evaluation to another. Therefore it is advantageous if the test data can be adapted to the needs of the respective use case in terms of several data characteristics, such as the number of clusters, the cluster sizes, or the degree of heterogeneity (a.k.a. dirtiness). This can be accomplished by applying a postprocessing step, which selects a subset of all data carefully. Further options for customization are the removal and merge of attributes, changing the character of the attributes’ values. A flexible and unconstrained customization requires that the test dataset contains (i) many duplicate clusters of various sizes and (ii) duplicate records of different degrees of heterogeneity, so that the user has a large set to choose from. In addition, it requires that the user can adjust the characteristics of the test data with relative ease. This can be supported by storing all records of one cluster together and providing precalculated heterogeneity scores.
The number of objects (i.e., clusters) was always 13.51 M.

Table 2: Statistical results of the generation process

<table>
<thead>
<tr>
<th>duplicate removal</th>
<th>#records</th>
<th>#dupl. pairs</th>
<th>cluster size</th>
<th>#removed</th>
<th>#dupl. pairs</th>
</tr>
</thead>
<tbody>
<tr>
<td>no</td>
<td>506.6 M</td>
<td>11,422.6 M</td>
<td>37.50</td>
<td>386</td>
<td>0%</td>
</tr>
<tr>
<td>exact</td>
<td>165.5 M</td>
<td>1,215.2 M</td>
<td>12.25</td>
<td>104</td>
<td>67.3%</td>
</tr>
<tr>
<td>trimming</td>
<td>120.0 M</td>
<td>640.8 M</td>
<td>8.88</td>
<td>77</td>
<td>76.3%</td>
</tr>
<tr>
<td>person data</td>
<td>58.4 M</td>
<td>135.4 M</td>
<td>4.32</td>
<td>51</td>
<td>88.5%</td>
</tr>
</tbody>
</table>

The number of historical information stored in the individual snapshots is rather low compared to the historical information of the whole dataset and thus only provides small clusters (see Figure 1a). Therefore, we also evaluated the whole dataset containing all records from every snapshot available. By doing so, we examined a total of 506,599,545 records.

One of our goals was to analyze the amount of (near) exact duplicates within the resulting test data. Moreover, we think that most potential users are only interested in the personal data of the voters, since the election and district attributes are very case-specific. Therefore, we executed our approach four times: (i) one time without removing any duplicate, (ii) one time with removing all exact duplicates, (iii) one time with removing all duplicates that were exact after their values have been trimmed (i.e., leading and trailing whitespaces were deleted), and (iv) one time with removing all duplicates whose personal data were equivalent (after trimming attribute values). To check the equivalence of duplicate records efficiently, we used the Message-Digest Algorithm 5 (short MD5) to calculate a hash value for each record. A record was then not imported into the test dataset when it already contained a record with the same hash value. Of course, collisions between non-exact records cannot be excluded for sure, but such a collision only means that the test dataset loses a duplicate record and thus does not have severe consequences if it happens a few times.\(^6\) The input to the hash function is the concatenation of the values of all relevant attributes to a single large string. As mentioned in Section 3.1.3, some meta data and time-related attributes can reduce the number of near exact duplicates drastically and therefore were not included into the concatenation. These attributes are the different dates (snapshot, load, registration and cancellation) and the age.\(^7\)

The number of resulting objects (i.e., duplicate clusters), records, duplicate pairs, the average and maximal number of records per object (i.e., duplicate cluster size), as well as the number of removed records and duplicate pairs are listed in Table 2. The number of distinct duplicate clusters (i.e., objects) per cluster size (i.e., number of records per object) are presented in Figure 1b (one time for all attributes and one time for the personal data only). Using a single snapshot did not produce any exact duplicate which was even less than expected. In contrast, using all snapshots produced hundred of millions of exact duplicates. Obviously, the average number of records per voter decreased when these exact duplicates were removed (e.g., 8.88 without whitespaces) and decreased further on when we restricted the data to the personal attributes (4.32), but was still large compared to the single-snapshot approach (1.18). In total, the number of

\(^6\)MD5 produces 128 bit hashes, which means that it is relatively unlikely that the hash values of two different inputs collide.

\(^7\)In the case of age values, the most obvious solution is to transform them into years of birth because the latter do not change. However, such a calculation would encode a part of the date of birth values, which were originally removed from the data due to privacy reasons. We have therefore decided to use them only for internal calculations (e.g., plausibility) and not to store them in the resulting test data.
records that were removed because of being exact duplicates was up to 76.3% and the number of removed duplicate pairs was up to 94.4% when all attributes where taken into account and up to 88.5% and 98.8% if only person data was considered. These large amounts of exact duplicates illustrate the importance of their removal, because otherwise every evaluation of duplicate detection algorithms using these data would suffer from the effects described in Section 3.1.3.

To estimate the value of future snapshots, we counted the number of new clusters (i.e., the snapshot contains an NCID that was never used before) and new records (i.e., the snapshot contains a record that was not part of any of the previous snapshots) per snapshot-year. These numbers are presented in Table 1 (note that the second includes the first). The last two columns show the percentage of rows that result in a new record (new record rate) and the percentage of new records that lead to a new cluster (new object rate). As expected, the number of new clusters and new records is the highest for the first snapshot. Surprisingly, the numbers of the following snapshots vary enormously (we expected an almost constant number). Investigations revealed that in some snapshots the formats of one or two attributes changed (e.g., from ‘64TH HOUSE’ to ‘NC HOUSE DISTRICT 64’) so that each of their records were considered to be ‘new’ even if – apart from that – they were identical with one of the already existing ones. This again shows us the importance of providing the user with an instrument that allows him to filter out records based on their similarities (see Section 6.5). However, as one can see, even the last four snapshots contributed a significant amount of new clusters and records to the test data. Thus, we can expect the same for future snapshots.

5 TEST DATA STORAGE
The voter data is originally given as a set of TSV files. However, we want to store our test dataset by using a data model that is more suitable with respect to its later usage, i.e. to evaluate duplicate detection algorithms as well as potential extensions with data from new snapshots. As a consequence, the data model has to satisfy three essential requirements:

- To customize the test data, we need to select and reduce duplicate clusters based on user-defined specifications. Moreover, when we integrate additional snapshot data, we need to calculate statistics by comparing duplicate records (e.g., plausibility and heterogeneity). Both require fast and collective access to all records of the same cluster.
- Every record of the voter data has 90 attributes, but only a few records have values for district-related attributes. This means that millions of records have missing values in at least 38 attributes. Thus, we require the underlying data model to provide an efficient handling of sparse data.
- Working on hundreds of gigabytes requires scalable software solutions.

Schemalsses NoSQL data models are much better suited to store sparse data than the relational model which requires the definition of a rigid schema. Moreover, many NoSQL data stores are designed to handle aggregates each of which is a collection of related data that we wish to treat as a unit [27]. Thus, they allow an easy and efficient way to access all the records of a certain person as we need it for customization and future extensions. Among all the available NoSQL data stores, we decided to use the document store MongoDB [18]. In contrast to the relational data model which is aggregate-ignorant, document stores are strongly aggregate-oriented [27] because they allow to (i) group records by storing them within the same document and (ii) nest different documents hierarchically. Furthermore, MongoDB is highly scalable. Besides its schemless structure, MongoDB has three features that are especially helpful for this work [18]:

- **Indexes**: Since our test dataset contains millions of nested documents, indexes are very important to efficiently select those documents from the dataset.
- **Aggregation Pipeline**: Multi-stage pipelines can be used to transform documents into an aggregated result. Available pipeline stages provide tools for filtering, transformation, grouping and sorting. These pipelines enable users to extract relevant subsets of the data and thus to customize their own test datasets.
- **Compass**: MongoDB has a powerful GUI called Compass. It enables the user to easily interact with the stored data with full CRUD functionality. It is very helpful for exploring, generating, adjusting and using the test data. Moreover, it allows to monitor load jobs of new snapshots and helps to identify mistakes at an early stage.

In our case, we created one document for every person (i.e., duplicate cluster) that in turn contains a document for every record of this person (which are grouped into an array) and in addition a document containing some relevant meta data including the hash values of the stored records. Since most users will be interested in the personal data only, we split every record into four parts (person, district, election and meta) and stored them into different subdocuments.
we associate every record with three version-similarity maps (one for plausibility and two for heterogeneity). Every value of parallel or sequential import of one or more new snapshots. In value is greater than the value of the desired version.

For doing this, we additionally store an array with the dates of inserted per snapshot. The reconstruction of similarity scores is cate cluster by a map that counts how many new records were snapshots per cluster, we enriched the meta data of every dupli-

ture 2 and consists of three steps. The first step corresponds to a (parallel or sequential) import of one or more new snapshots. In the second step, current statistics are updated and (if required) new statistics are calculated. In the last step, a new version number is associated to the test data, versioning-related meta data are updated and the new version is published. As illustrated by this figure, the update process (and thus the creation of a new version) can be triggered by two reasons:

• new snapshots are available, or

• new statistics are required.

Logically, in the second case, the first step is skipped and the update process starts immediately with the second one.

5.1.1 Update Process. The update process is depicted in Figure 2 and consists of three steps. The first step corresponds to a (parallel or sequential) import of one or more new snapshots. In the second step, current statistics are updated and (if required) new statistics are calculated. In the last step, a new version number is associated to the test data, versioning-related meta data are updated and the new version is published. As illustrated by this figure, the update process (and thus the creation of a new version) can be triggered by two reasons:

• new snapshots are available, or

• new statistics are required.

5.1 Future Updates & Reproducibility
The state of North Carolina publishes a new snapshot at every election and every New Year’s Day. Moreover, we have observed that they publish some old snapshots belatedly (e.g., the snapshot from 2010-11-02 was published in May 2019). To improve our test dataset both in size and heterogeneity, we will update it regularly.

5.1.1 Update Process. The update process is depicted in Figure 2 and consists of three steps. The first step corresponds to a (parallel or sequential) import of one or more new snapshots. In the second step, current statistics are updated and (if required) new statistics are calculated. In the last step, a new version number is associated to the test data, versioning-related meta data are updated and the new version is published. As illustrated by this figure, the update process (and thus the creation of a new version) can be triggered by two reasons:

• new snapshots are available, or

• new statistics are required.

Logically, in the second case, the first step is skipped and the update process starts immediately with the second one.

5.1.2 Reproducibility. As described in Section 3.3, the import of new data poses some challenges to the repeatability of evaluation processes that were using previous versions of our test data. Since no record is ever removed from the test dataset, it grows monotonically, which means that the set of all records of the current version will always be a subset of all records of every future version. Thus, theoretically, it is sufficient to add a field to every record which is monotonically increasing with every new update. This field can be the date of import or the number of the first version containing this record (the snapshot date is not suitable because it is not monotonically increasing due to belatedly published snapshots). To reconstruct an earlier version, the user can use this field to filter out all records whose field value is greater than the value of the desired version.

However, we also want to allow users to limit their evaluation to an arbitrary subset of snapshots (e.g., a certain time interval). For doing this, we additionally store an array with the dates of all snapshots containing the corresponding record.

To reconstruct statistics, such as the number of records or snapshots per cluster, we enriched the meta data of every duplicate cluster by a map that counts how many new records were inserted per snapshot. The reconstruction of similarity scores is discussed in Section 5.2.

5.2 Storing Similarity Scores
To support the user in customizing its data by (i) removing further near exact duplicates, (ii) repairing potentially unsound clusters, and (iii) restricting the data to a certain range of heterogeneity, we associate every record with three version-similarity maps (one for plausibility and two for heterogeneity). Every value of each of these maps corresponds to another map that assigns a similarity score to each of the previously existing records of the same cluster. Since the order of these records never change, this approach does not only avoid expensive recalculations, it also en-

Table 3: Examples of erroneous and unsound clusters

<table>
<thead>
<tr>
<th>NCID</th>
<th>last_name</th>
<th>first_name</th>
<th>midd_name</th>
<th>sex</th>
<th>age</th>
</tr>
</thead>
<tbody>
<tr>
<td>r1</td>
<td>WILLIAMS</td>
<td>DEBRA</td>
<td>OEHRIE</td>
<td>FEMALE</td>
<td>45</td>
</tr>
<tr>
<td>r2</td>
<td>WILLIAMS</td>
<td>DEBRA</td>
<td>OEHRIE</td>
<td>FEMALE</td>
<td>47</td>
</tr>
<tr>
<td>r3</td>
<td>OEHRIE</td>
<td>DEBRA</td>
<td>ANN</td>
<td>FEMALE</td>
<td>49</td>
</tr>
<tr>
<td>r4</td>
<td>FIELDS</td>
<td>MARY</td>
<td>ELIZABETH</td>
<td>FEMALE</td>
<td>61</td>
</tr>
<tr>
<td>r5</td>
<td>BETHEA</td>
<td>JOSHUA</td>
<td>ELIZABETH</td>
<td>MALE</td>
<td>93</td>
</tr>
</tbody>
</table>

6 EXPERIMENTAL EVALUATION
When we explored the snapshots, we noticed several errors within the data. Some records contain typos, abbreviations, or have values confused between attributes. Moreover, some notations have changed over time (e.g., ‘1ST CONGRESSIONAL’ vs. ‘CO. DISTRICT 1’ or ‘66 AND ABOVE’ vs. ‘Age Over 66’). One example is presented in Figure 3. The name values of at least one record of voter DB175272 got mixed up. Furthermore, at least one of the middle name values contains a typo (‘OEHRIE’ vs. ‘OEHRLE’). Remember that a proper evaluation of duplicate detection algorithms requires errors of many different types and not just outdated values. Thus, such real-life data errors are very welcome in our test dataset, since they challenge the detection of duplicates, but do not corrupt the gold standard. However, we also detected some duplicate clusters that contain records that hardly refer to the same person. An excerpt of one of those examples is depicted in Figure 3 where the two records r4 and r5 share the same NCID, but obviously describe different persons. Such unsound clusters are a real threat to the quality of our test data because they spoil the gold standard and thus will negatively affect every future evaluation if they remain in the dataset.

In order to evaluate the quality and usability of our test dataset beyond these first impressions, we conducted a series of experiments, which are described in the rest of this section.

6.1 Evaluated Datasets
To better understand the results of the evaluation of our test dataset, we compare them with those of three manually labeled test datasets that are commonly used in the literature. We ac-

Note, the plausibility is already limited to personal attributes and thus does not need to be stored twice.

We used the non-prepared versions where special characters are not removed.

http://hpi.de/naumann/projects/repeatability/datasets
For all three datasets the duplicate information is provided by a list of pairs. Several characteristics of these datasets are presented in Table 3. Interestingly, the duplicate distributions of these sets are quite different. Whereas the Cora set contains very large clusters (up to 238 records) and its average cluster size is 10.32, the maximal and average cluster size of the Census and CDDB datasets are quite small (1.74 or 1.03 respectively). The three datasets NC1 to NC3 are described in Section 6.5.

### 6.2 Plausibility Check

As we have explained in Section 3.1.1, it is very important that the gold standard of the test data is sound. However, as shown in Figure 3, we have also seen that this does not always seem to be the case. To keep the threat of an unsound gold standard to a minimum, we performed a quality check by calculating a plausibility score for every pair of duplicate records. Since we cannot make any decision with absolute confidence, we did not remove any cluster (or record) from the data, but instead equipped each of them with their respective plausibility score so that the user can decide herself what risk she wants to take to include a potentially unsound cluster into her test data.

In the plausibility check, we have the basic assumption that all records of the same cluster are duplicates and the similarity scores should only reflect (significant) contradictions to this assumption. Consequently, the similarity measure should compensate simple errors and differences in data representation as we know it from duplicate detection algorithms. Due to our basic assumption, this compensation can be even stricter. Therefore, word confusions within an single attribute value or between different values as well as missing or abbreviated values should not reduce similarity at all, because they are more an indication of unknown or erroneous values than a clear indication of a non-duplicate. Moreover, to compensate outdated values we should only use attributes whose values rarely change and that are either very identifying (i.e., two records with the same value are likely duplicates) or discriminating (i.e., two records with different values are likely no duplicates). In our use case, we decided to use:

- the three name values (first, middle and last),
- the sex code,
- the year of birth (which we derived from the snapshotdate and the age value), and
- the place of birth.

It is not uncommon that values are confused between the three name attributes. Thus, we computed a single name similarity before combining it with the similarity scores of the other attributes. To capture errors in the name order, but also within the individual name values (e.g., typos), we computed the name similarity by using the hybrid measure Generalized Jaccard Coefficient [5] with an extended version of the Damerau-Levenshtein Similarity as the internal token similarity measure, i.e.:

\[
\text{sim}_{\text{name}}(o_i, o_j) = \text{GenJacc}_{\text{DumLe}}(\text{name}(o_i), \text{name}(o_j))
\]

where \(\text{name}(o_i) = (\text{FN}(o_i), \text{MN}(o_i), \text{LN}(o_i))\). The Damerau-Levenshtein Similarity was extended to a proper handling of missing and abbreviated values. The comparison to a missing value results in a similarity of 1. The same holds true if one token is a prefix of the other because in both cases we do not have any evidence to mistrust the given data. In the case of the sex code, typos and different representations can almost be excluded for sure. Thus there are actually only four possibilities: The compared values agree (i.e., \(\text{sim}_{\text{sex}} = 1\)), disagree (i.e., \(\text{sim}_{\text{sex}} = 0\)), one of them is undesignated (i.e., it has the value ‘U’) or missing. Since we do not have any contradiction in the later two cases, we set the similarity to \(\text{sim}_{\text{sex}} = 1\).

We computed the year of birth (short YoB) as snapshotdate – age. Since the actual YoB can be one year earlier if the person has not yet had birthday when the snapshot was made, we introduced a tolerance of 1. Moreover, we assumed a similarity of 0 if the age difference was 10 or greater. This led to the following formula:

\[
\text{sim}_{\text{YoB}}(o_i, o_j) = 1 - \min\left(1, \frac{\max(0, |\text{YoB}(o_i) - \text{YoB}(o_j)| - 1)}{10}\right)
\]

In the case of the place of birth we simply computed the extended Damerau-Levenshtein Similarity between the two values. The final similarity score was then calculated as the weighted average of the previously presented scores where we considered the name similarity to be more important (weight 0.5) than the others (each 0.15). A cluster is already unsound, if only one of its records refers to another voter regardless of how plausible the other records are actually duplicates. Thus, we computed the plausibility of a cluster as the minimal plausibility of its records.

We performed our plausibility check on the dataset with 120 million records (exact duplicates were removed after trimming). The results show that only a few clusters of this dataset are highly suspicious to be unsound. The average cluster plausibility is 0.99, 92.8% of all clusters (and 93.6% of all duplicate pairs) have the maximum possible value 1.0. The distribution of the remaining 7.2% (or 6.4% resp.) are presented in Figure 4a. The minimal plausibility of all clusters (and pairs) is 0.06. 5.5% of all clusters have a plausibility lower than 0.9, 0.43% (=58,083 clusters) lower than 0.8 and only 0.0045% (=608 clusters) lower than 0.5. The pair-based values are similar. As a comparison, the two clusters from Figure 3 have a plausibility of 0.81 (DB175272) and 0.33 (DR19657)

---

Table 3: Characteristics of evaluated datasets

<table>
<thead>
<tr>
<th>dataset</th>
<th>Cora</th>
<th>Census</th>
<th>CDDB</th>
<th>NC1</th>
<th>NC2</th>
<th>NC3</th>
</tr>
</thead>
<tbody>
<tr>
<td>#records</td>
<td>1,879</td>
<td>841</td>
<td>9,763</td>
<td>24,761</td>
<td>22,739</td>
<td>22,402</td>
</tr>
<tr>
<td>#attributes</td>
<td>17</td>
<td>6</td>
<td>7</td>
<td>38</td>
<td>38</td>
<td>38</td>
</tr>
<tr>
<td>#duplicate pairs</td>
<td>64,578</td>
<td>376</td>
<td>300</td>
<td>19,916</td>
<td>15,993</td>
<td>15,263</td>
</tr>
<tr>
<td>#clusters</td>
<td>182</td>
<td>483</td>
<td>9,508</td>
<td>10,000</td>
<td>10,000</td>
<td>10,000</td>
</tr>
<tr>
<td>#non-singletons</td>
<td>118</td>
<td>345</td>
<td>221</td>
<td>10,000</td>
<td>10,000</td>
<td>10,000</td>
</tr>
<tr>
<td>max. cluster size</td>
<td>238</td>
<td>4</td>
<td>6</td>
<td>7</td>
<td>8</td>
<td></td>
</tr>
<tr>
<td>avg. clustersize</td>
<td>10.32</td>
<td>1.74</td>
<td>1.03</td>
<td>2.45</td>
<td>2.27</td>
<td>2.24</td>
</tr>
<tr>
<td>max. heterog.</td>
<td>0.63</td>
<td>0.46</td>
<td>0.65</td>
<td>0.2</td>
<td>0.46</td>
<td>0.83</td>
</tr>
<tr>
<td>avg. heterog.</td>
<td>0.171</td>
<td>0.15</td>
<td>0.218</td>
<td>0.09</td>
<td>0.304</td>
<td>0.487</td>
</tr>
</tbody>
</table>

---

respectively which matches our intuition that the differences in the first cluster are probably the result of data errors in the name values while the second cluster contains obvious non-duplicates.

An appropriate scoring of plausibility heavily depends on the domain of the data, since we should only use attributes that are less volatile and are either very identifying or discriminating. Moreover, it also depends on the quality of the data, since typical error patterns (e.g., an incorrect encoding of special symbols) are no significant evidence for an unsound cluster and should therefore be compensated in the scoring process. It is therefore difficult to make comparisons between the plausibility of datasets defined on different schemes without creating any noticeable bias. For this reason, we decided not to include such a plausibility calculation for the Cora, Census and CDDB datasets.

6.3 Duplicate Heterogeneity

To score the heterogeneity between two duplicate records we want to take all their differences into account and thus do not actually want to apply measures that compensate them. At the same time, there are differences that should cause a larger heterogeneity than others. For example, difference in upper and lower case or confusions of tokens are less significant than replacing the original strings with completely different letters or tokens.

To address the problem resulting from uppercase letters, we decided to compare every two values one time with and one time without lowercasing them. To address the problem of token confusions, we decided to compare every two values one time with a sequential and one time with a hybrid similarity measure. Together this results in four comparisons for every two values. We used the average of the four resulting scores as final similarity. As the sequential similarity measure, we chose Damerau-Levenshtein. Since the Generalized Jaccard Coefficient is computationally too expensive when working on 90 attributes, we used the Monge-Elkan Similarity\textsuperscript{13} \cite{19} as the hybrid measure instead (using Damerau-Levenshtein as an internal token similarity measure)\textsuperscript{14}. To calculate the heterogeneity between two records, we used the weighted average of their inverse value similarities (i.e., the records are the more heterogeneous, the less similar they are).

If the resulting heterogeneity scores should reflect any kind of difficulty to detect the corresponding fuzzy duplicates, identifying attributes, such as names, should be weighted higher than others. At the same time and in contrast to plausibility, we want to compare the heterogeneity (i.e., dirtiness) of different datasets in order to enable an evaluation of algorithms with respect to a varying dirtiness of the data. To ensure a fair comparison, no context knowledge and external expertise should be included into the scoring process and all necessary information should be taken from the dataset itself. To accomplish that we used the same similarity measure for all attributes and weighted every attribute by its uniqueness, where we quantified this uniqueness by the attribute’s entropy \cite{15}. Since duplicate records distort these uniqueness scores (e.g., an otherwise unique id can occur multiple times), we used only one record per duplicate cluster to compute them. The heterogeneity of a cluster was then computed as the average heterogeneity of its records. Since a consideration of clusters of size one is pointless, we restricted our analyses to clusters with at least two records.

The results of our analysis are depicted in Figure 4b. They show that – despite of outdated values and data errors – most of the duplicate records are very similar and the dataset as a whole is quite clean and homogeneous. The average heterogeneity is 0.09 (cluster) and 0.16 (pairs) respectively. However, since we removed exact duplicates, almost none of the clusters is completely homogeneous and most of them have a heterogeneity of around 0.04 (41.3%). In the case of the duplicate pairs, we do not have such a large peak, but also here you can see, that most values are in the range between 0.02 and 0.07. The maximal heterogeneity is 0.64 (cluster) and 0.9 (pairs). As a comparison, the two clusters from Figure 3 have a heterogeneity of 0.38 (DB175272) and 0.35 (DR19657) respectively. Interestingly, the more plausible cluster is also more heterogeneous. That is because although cluster DR19657 contains records referring to different persons, these records form two very homogenous groups (one with six records similar to $r_6$ and one with four records similar to $r_5$).

To score the heterogeneity of the Cora, Census und CDDB datasets, we used the same settings (i.e., same similarity measures and all attributes are weighted based on their entropy). The pair-based heterogeneity distributions of all three datasets are depicted in Figure 4c. The heterogeneity scores of the Cora set are almost normally distributed. Most of them have a value of 0.15, the maximal value is 0.63 and the average is 0.171. The Census set has three peaks at 0, 0.07 and 0.1, a maximal heterogeneity of 0.46 and an average of around 0.15. In general, except of the large number of very homogenous pairs, its distributions has some similarity to this of the Cora set, but is a little less regular. The

\textsuperscript{13}Since this measure is non-symmetric, we computed it in both directions and used the average as final score.

\textsuperscript{14}There are millions of possible similarity measures (including various settings) and we had to choose one even though this could possibly generate a bias in the evaluation processes using this test data. However, as illustrated in Section 6.5, this bias was almost not existing in our experiments.
6.4 Diversity of Error Types

To allow an extensive evaluation of the capabilities of duplicate detection algorithms, we chose source data whose collection process promises many different types of errors. To test this assumption, we conducted a statistical analysis on the personal attributes of our test dataset by searching for several kinds of irregularities within these data (see Table 4). Here, we distinguished between irregularities that can be identified by analyzing single records (so-called singletons) and those that can only be detected by comparing two duplicate records (so-called pair-based irregularities).

In the first case, we evaluated every record individually leading to a frequency that can be normalized using the total number of records. In the second case, we compared every two duplicate records, counted the number of times the individual irregularities occur and normalized them using the total number of duplicate pairs. Moreover, we distinguished between irregularities concerning a single attribute and irregularities concerning multiple attributes. We evaluated the following singletons:

- **outlier**: A value that is outside a predefined range (e.g., age > 110) or contains a character that is unusual for its associated domain (e.g., the first name ‘X ÆA-12’).
- **abbreviation**: A value that consists of a single letter, possibly followed by a punctuation mark.
- **missing**: A value that is null, an empty string or any other value indicating missing information (e.g., ‘-’ or ‘unknown’).

As pair-based irregularities, we analyzed:

- **typos**: Two values whose lowercase versions differ only in one character or contain a character transposition. These are exactly those values having a Damerau-Levenshtein distance of 1. In order not to interpret a complete replacement of one value by the other as a typing error, we only considered values longer than two.
- **phonetical error**: Two values that are not identical after removing non-letter characters, are both longer than two and have the same soundex code.
- **token transpositions**: Two values whose token sets are identical, but their order is different.
- **prefix/postfix**: Two values where one of them is a prefix/postfix of the other after removing a potential punctuation mark from the end of the shorter value. Such prefix/postfix situations indicate abbreviations or forgotten tokens/characters.
- **OCR-error**: Two distinct string values which only differ at those positions where one of them has a digit. If both characters are digits, they need to be identical.
- **different representation**: Two values that only differ in nonalphabetical characters (e.g., a hyphen, space or punctuation mark between tokens).

### Table 4: Statistics of different irregularities with the NC, the Cora and the Census datasets

<table>
<thead>
<tr>
<th>error type</th>
<th>example1</th>
<th>NC (total 135 M pairs)</th>
<th>Cora (total 65 K pairs)</th>
<th>Census (total 376 pairs)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>most common attribute</td>
<td>frequency total</td>
<td>perc.</td>
<td>most common attribute</td>
</tr>
<tr>
<td>outlier</td>
<td>age = 5069</td>
<td>age</td>
<td>281 K</td>
<td>0.5%</td>
</tr>
<tr>
<td>abbreviation</td>
<td>midl_name = A</td>
<td>midl_name</td>
<td>7.4 M</td>
<td>12.7%</td>
</tr>
<tr>
<td>missing</td>
<td>midl_name = null</td>
<td>mail_addr 3</td>
<td>58 M</td>
<td>99%</td>
</tr>
<tr>
<td>typo</td>
<td>ADELL</td>
<td>midl_name</td>
<td>1.2 M</td>
<td>0.9%</td>
</tr>
<tr>
<td>OCR-error</td>
<td>DICOL3</td>
<td>last_name</td>
<td>1.8 K</td>
<td>0.0%</td>
</tr>
<tr>
<td>phonetic</td>
<td>BAILEY</td>
<td>midl_name</td>
<td>1 M</td>
<td>0.7%</td>
</tr>
<tr>
<td>prefix</td>
<td>KIM</td>
<td>midl_name</td>
<td>5.3 M</td>
<td>3.9%</td>
</tr>
<tr>
<td>postfix</td>
<td>BRAGG</td>
<td>last_name</td>
<td>232 K</td>
<td>0.2%</td>
</tr>
<tr>
<td>formatting</td>
<td>JRS RIDGE</td>
<td>midl_name</td>
<td>213 K</td>
<td>0.2%</td>
</tr>
<tr>
<td>token transp.</td>
<td>ANH THI</td>
<td>race_desc</td>
<td>743 K</td>
<td>0.5%</td>
</tr>
<tr>
<td>value confusion</td>
<td>(JOSE,JUAN)</td>
<td>first/midl_name</td>
<td>20 K</td>
<td>0.0%</td>
</tr>
<tr>
<td>integrated value</td>
<td>(MAN LL )</td>
<td>midl/last_name</td>
<td>242 K</td>
<td>0.2%</td>
</tr>
<tr>
<td>scattered value</td>
<td>(AN LE MA)</td>
<td>midl/last_name</td>
<td>21 K</td>
<td>0.0%</td>
</tr>
</tbody>
</table>

1 Selected from some attribute of the NC dataset.
2 Singletons are normalized using the total number of records (NC = 58.4 M, Cora = 1,879, Census = 841).
3 Pair-based irregularities are normalized using the total number of duplicate pairs (NC = 135.4 M, Cora = 65 K, Census = 376).
value confusion: Two records whose values are confused between two different attributes (e.g., the first and last name of one record are transposed).

integrated value: Two records where in one of them the value of one attribute is integrated into another (e.g., a middle name stored as a second token in the first name).

scattered values: Two records having the same set of tokens assigned differently to two attributes. To avoid possible overlaps with the previous two types, we only counted scattered values that are not integrated or confused.

Obviously some of these irregularities overlap with (or sometimes even include) others so that we counted some of the errors for more than one type. For example, some OCR-errors are also typos. Moreover, it is important to note that we consider these irregularities as indications of particular error types, but cannot always classify them with absolute confidence. For example, not every two distinct values that have the same soundex code are an actual phonetical error. Nevertheless, the errors are real even if their assignment to the individual types may be disputed. It should also be mentioned that our definitions of the individual error types do not cover them completely so that our analysis was not able to find every actual error. For example, OCR errors that do not contain digits (e.g., `Tim' vs. `Tirn') were not recognized as such. However, despite these minor inaccuracies, we think that our experiment provides a good overview of the wide variety of different errors contained in the voter data.

The results of this analysis are presented in Table 4 (grouped by singletons and pair-based irregularities), were we list the absolute values in combination with their percentages. To achieve greater comparability, we also evaluated the Census and the Cora datasets. We selected the Census dataset because it has a similar domain as our voter data and selected the Cora dataset because it has similarly large clusters. As we can see, the percentages of the Census and the Cora datasets are much higher than those of the NC dataset. For example, the percentage of typos in the attribute last_name of the Census dataset is 65%. This means that out of 376 duplicate pairs, 243 differed in this attribute by only one character or had two consecutive characters transposed. Although the percentages of the NC dataset are much smaller than those of the Cora and the Census datasets, the absolute numbers are many times larger. This shows the potential for customizing smaller datasets with higher error percentages, but still containing million of records. Moreover, the NC dataset contains irregularities that are (almost) not contained in the Cora and Census datasets. Examples are OCR-errors or errors that affect more than one attribute.

6.5 Usability

There are various ways to customize a test dataset by using the precalculated heterogeneity scores. In our experiments, we sketched a very simple approach and let the development of more sophisticated approaches to the user (or future research resp.). This approach consists of three steps. In the first step, we defined a lower and an upper bound \( h_L \) and \( h_T \) for the heterogeneity scores. In the second step, we randomly selected over 100 thousand clusters from our whole dataset, scanned over all records of every cluster in their sorting order and removed every record whose heterogeneity to its preceding (not removed) records was not in the requested range \( [h_L, h_T] \). In the third step, we sorted the reduced clusters by their size and selected the 10 thousand largest clusters as test input. We applied this approach for the three settings \( (h_L, h_T) \in \{(0.06, 0.2), (0.2, 0.4), (0.4, 1.0)\} \) while restricting the schema to the personal attributes. The result are the three test datasets NC1, NC2 and NC3. The characteristics of these datasets are depicted in Table 3. As the numbers show, even though we only used 0.07% of all clusters as input, these datasets are larger than the Cora, Census and CDBB datasets.

To evaluate the difficulty of detecting fuzzy duplicates within the individual datasets, we applied three duplicate detection algorithms each using another similarity measure (the same for all attributes). The first measure (short ME/Lev) was the same combination of Monge-Elkan and Damerau-Levenshtein as we used it to calculate the heterogeneity scores (see Section 6.3). The other two measures\(^{15} \) were the Jaro Winkler (sequential) and the Jaccard Similarity using trigrams (token-based) [5]. The similarity of two records was always computed as the weighted average similarity of their values. Since we observed that the name values are often confused between the individual attributes, we matched every combination of them and used the 1:1 matching with the highest similarity for aggregation. To weight the individual attributes we used again their entropy. In this case, however, we calculate it using all records (i.e., including the duplicates), since the user cannot know these duplicates in advance. In addition, the entropy was calculated solely based on the 20 thousand records of the customized datasets. Thus, the resulting weights differ from the ones we used to calculate the heterogeneity scores (e.g., 0.54 vs. 0.5 for the first name). To reduce the search space, we applied a multi pass of the Sorted Neighborhood Method [19] by using one pass for each of the five most unique attributes and a window of size \( w = 20 \). None of the true duplicates were lost through this reduction.

The results of these duplicate detection algorithms applied to the different test datasets are depicted in Figure 5. As we can see in Figure 5a to 5c, the quality of the duplicate detection algorithms decreased when we increased the heterogeneity of the test data, since the more difficult it was to separate the duplicate from the non-duplicate pairs. In the first case, the test dataset was very clean and we could achieve almost a perfect \( F_1 \)-score for all four measures. Moreover, for two out of three measures, this score was high for every threshold between 0.55 and 0.85, which made it easier to select an appropriate value for this threshold without knowing these numbers. In the case of the second dataset, the maximal \( F_1 \)-score was still pretty solid (i.e., close to 0.8), but the threshold had to be set much more carefully and the quality of a setting already depended on the individual measures. For example, for Jaccard the best threshold was 0.57, but for Jaro Winkler it was 0.74 and there was no threshold that worked well for all measures. Finally, in the case of the last dataset, the maximal \( F_1 \)-score decreased significantly and even a score of 0.4 was hard to achieve. Moreover, the differences between the individual similarity measures became more noticeable which means that the selection of this measure was much more important than in the previous two cases. All this shows that the precalculated heterogeneity scores can be perfectly used to adjust the test data, since the more difficult it was to separate the duplicate from the non-duplicate pairs.

15 Here we tried to cover a wide range of measures by using hybrid, sequential and token-based measures.
Almost all existing test datasets have been either artificially generated test data is the ERIQ dataset [2] is that the resulting standard is realistic. Data pollution tools, such as GeCo [4], TDGen [1], or DaPo [12], are the best option to generate test data with realistic values and value patterns because real-life data can be used as input. Moreover, if a broad spectrum of error types is supported they are nearly domain-independent. Except DaPo, however, existing pollution tools are strongly limited with respect to their scalability so that generating large datasets is either impossible or extremely time consuming [12]. A major problem that all these tools have in common is an appropriate simulation of outdated values and the complex error patterns that result from them.

Data of the NC voter register have been already used as test data in several works (see Table 5). Sadly, in most cases these uses are not fully documented, so we cannot say exactly which datasets were used there. The small size of the first dataset lets us to create test data that are cleaner (NC1), equally clean (NC2) and dirtier (NC3) than these real-life use cases giving us the opportunity to design our test data in the way our evaluation goals require. We repeated this experiment with different parts of our original test dataset as input. Since the compositions of the generated datasets differ slightly, there were also slight differences in the resulting graphs, but the findings were always the same.

7 RELATED WORK

Almost all existing test datasets have been either artificially generated (e.g., by using an automatic generation tool) or been manually labeled. While artificial datasets have the obvious disadvantage of not containing actual real-life duplicates and errors (including outdated values), labeling duplicates within a dataset manually is extremely expensive so that this approach is only an option for very small datasets. For example, the commonly used Cora, Census and CDDB datasets contain less than ten thousand records each (see Section 6.1). The essential shortcoming of an automatic labeling approach [32] is that the resulting standard is often not sound and biased towards the used algorithms, which can significantly distort evaluation results. One example of artificially created test data is the ERIQ dataset [31] which contains 100 thousand records of customer data. Although this number is much larger than those of the manually labeled datasets, it is still small compared to typical big data applications. Further test datasets (manually labeled or artificially generated) can be found on the websites of the Hasso-Plattner Institute[16] and the University of Leipzig[16]. To the best of our knowledge, none of the existing test datasets provides precalculated similarity scores that help users to generate customized datasets on their basis.

Automatic test data generation tools can be divided into two classes: Data synthetization tools that generate all data values – including duplicates and errors – from scratch and data pollution tools that get a clean dataset as input and pollute it with duplicates, errors and inhomogeneities. Data synthetization tools, such as DBGen [11] or the Febrl Data Set Generator [2], are very efficient so that large datasets can be generated in short time [12]. However, since every data value is fictional, it is almost impossible to guarantee that the resulting data (values and patterns) are realistic. Data pollution tools, such as GeCo [4], TDGen [1], or DaPo [12], are the best option to generate test data with realistic values and value patterns because real-life data can be used as input. Moreover, if a broad spectrum of error types is supported they are nearly domain-independent. Except DaPo, however, existing pollution tools are strongly limited with respect to their scalability so that generating large datasets is either impossible or extremely time consuming [12]. A major problem that all these tools have in common is an appropriate simulation of outdated values and the complex error patterns that result from them.

Figure 5: $F_1$-scores in relation to similarity threshold for several similarity measures and test datasets

Cora, Census and CDDB datasets, we see that they show similar patterns as NC2 in terms of the maximally achieved $F_1$-score and the shapes and positions of the individual graphs. Only the Census dataset differs a little bit, because here the Jaro Winkler scored much better than for the other datasets and the single graphs correspond less to a bell shape. In summary, this shows that the sheer amount of data of our original test dataset enables us to create test data that are cleaner (NC1), equally clean (NC2) and dirtier (NC3) than these real-life use cases giving us the opportunity to design our test data in the way our evaluation goals require. We repeated this experiment with different parts of our original test dataset as input. Since the compositions of the generated datasets differ slightly, there were also slight differences in the resulting graphs, but the findings were always the same.

16https://dbs.uni-leipzig.de/en/research/projects/object_matching/benchmark_datasets_for_entity_resolution
17https://dls.ncsbe.gov/index.html?prefix=datalab
duplicate distributions of the first three datasets, however, are nowhere near the numbers of the test dataset we generated in our study. The last two datasets have been artificially polluted with duplicates and errors using the GeCo tool [4]. Thus, compared to our dataset, it does not contain real-life errors and especially lacks in realistic outdated values. Moreover, none of these authors enhanced their data with useful statistics, such as plausibility and heterogeneity scores, as we did in this paper and do not provide mechanisms that enable the reproducibility of past usage. Finally, to the best of our knowledge we are the first who discuss the aspects of quality, usability and reproducibility in the context of test data for duplicate detection in such a depth.

8 CONCLUSION & FUTURE WORK

In this paper, we presented a large-scale dataset for evaluating duplicate detection algorithms and the approach behind its generation. Extracted from an historical voter register from North Carolina, our test dataset contains more than 120 million records and 640 million duplicate pairs making it uniquely suitable for evaluating duplicate detection at scale. Besides the dataset’s size, our study focused on its quality, usability and reproducibility. The record’s historical nature means that they contain many outdated values and – since data was often entered manually – also a large variety of other error types, such as typos, phonetical errors or confusions between attributes. While the data values themselves contain many errors, the underlying gold standard is largely error-free which is a mandatory requirement to ensure meaningful evaluation results. In general, its large size as well as the large number of data errors makes the dataset perfectly suitable for users who want to customize their own datasets based on the requirements of their respective evaluation goals. To support such customizations, we equipped the individual records with similarity scores modeling their plausibility and heterogeneity. Finally, we integrated several mechanisms to ensure reproducibility when the dataset is growing with future updates. In summary, our approach enables generating large-scale test datasets with realistic errors including outdated values (which are hard to synthesize) and without the need for labeling duplicates manually (which is extremely labor-intensive).

Our plans for future work targets two different ways to extend our approach. First, we intend to generalize the procedure described here and apply it to historical corpora from other domains. This will provide the research community with large-scale test datasets beyond use cases that revolve around personal data. Second, we plan to combine our approach with a scalable data pollution tool, such as DaPo, to unite the strengths of having real outdated values and being able to inject additional errors at will. Our goal here is to increase the flexibility for customization and thereby facilitate generating test datasets geared for specific user demands. We think our presented work is useful to other researchers and we hope that our current line of research will pave the way for novel solutions that combine approaches using historical data with methods of data pollution in creative ways.

REFERENCES