

A Probabilistic Deduplication, Record Linkage and Geocoding System

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Project web page: <http://datamining.anu.edu.au/linkage.html>

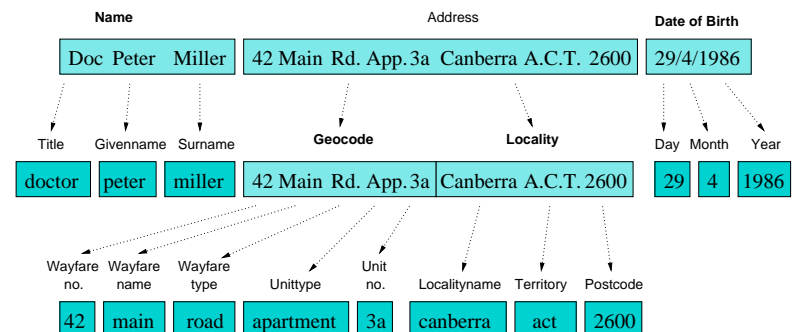
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- Data cleaning and standardisation
- Record linkage / data integration
- *Febri* overview
- Probabilistic data cleaning and standardisation
- Blocking / indexing
- Record pair classification
- Parallelisation in *Febri*
- Data set generation
- Geocoding
- Outlook

Data cleaning and standardisation (1)

- Real world data is often *dirty*
 - Missing values, inconsistencies
 - Typographical and other errors
 - Different coding schemes / formats
 - Out-of-date data
- Names and addresses are especially prone to data entry errors
- Cleaned and standardised data is needed for
 - Loading into databases and data warehouses
 - Data mining and other data analysis studies
 - Record linkage and data integration

Data cleaning and standardisation (2)



- Remove unwanted characters and words
- Expand abbreviations and correct misspellings
- Segment data into well defined *output fields*

Record linkage / data integration

- The task of linking together records representing the same entity from one or more data sources
- If no *unique identifier* is available, *probabilistic linkage techniques* have to be applied
- Applications of record linkage
 - Remove duplicates in a data set (internal linkage)
 - Merge new records into a larger master data set
 - Create customer or patient oriented statistics
 - Compile data for longitudinal studies
 - Geocode data

Data cleaning and standardisation are important first steps for successful record linkage

Record linkage techniques

- Deterministic or exact linkage
 - A *unique identifier* is needed, which is of high quality (precise, robust, stable over time, highly available)
 - For example *Medicare*, *ABN* or *Tax file* number (are they *really* unique, stable, trustworthy?)
- Probabilistic linkage (*Fellegi & Sunter, 1969*)
 - Apply linkage using available (personal) information
 - Examples: *names*, *addresses*, *dates of birth*
- Other techniques (rule-based, fuzzy approach, information retrieval)

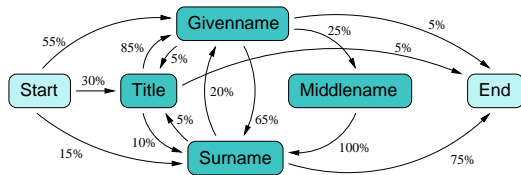
Febrl – Freely extensible biomedical record linkage

- An experimental platform for new and improved linkage algorithms
- Modules for data cleaning and standardisation, record linkage, deduplication and geocoding
- Open source <https://sourceforge.net/projects/febrl/>
- Implemented in *Python* <http://www.python.org>
 - Easy and rapid prototype software development
 - Object-oriented and cross-platform (*Unix*, *Win*, *Mac*)
 - Can handle large data sets stable and efficiently
 - Many external modules, easy to extend

Probabilistic data cleaning and standardisation

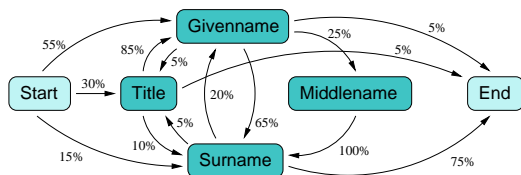
- Three step approach
 1. Cleaning
 - Based on look-up tables and correction lists
 - Remove unwanted characters and words
 - Correct various misspellings and abbreviations
 2. Tagging
 - Split input into a list of words, numbers and separators
 - Assign one or more tags to each element of this list (using look-up tables and some hard-coded rules)
 3. Segmenting
 - Use either rules or a *hidden Markov model (HMM)* to assign list elements to *output fields*

Hidden Markov model (HMM)



- A HMM is a *probabilistic* finite state machine
 - Made of a set of *states* and *transition probabilities* between these states
 - In each state an *observation* symbol is emitted with a certain probability distribution
 - In our approach, the observation symbols are *tags* and the states correspond to the *output fields*

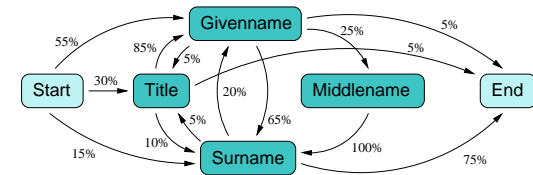
Probabilistic data cleaning and standardisation – Example



- Uncleaned input string: 'Doc. peter Paul MILLER'
Cleaned into string: 'dr peter paul miller'
- Word and tag lists:


```
['dr', 'peter', 'paul', 'miller']
['TI', 'GM/SN', 'GM', 'SN']
```
- Two example paths through HMM
 - 1: Start -> Title (TI) -> Givenname (GM) -> Middlename (GM) -> Surname (SN) -> End
 - 2: Start -> Title (TI) -> Surname (SN) -> Givenname (GM) -> Surname (SN) -> End

HMM data segmentation



- For an observation sequence we are interested in the most likely path through a given HMM (in our case an observation sequence is a *tag list*)
- The *Viterbi* algorithm is used for this task (a dynamic programming approach)
- *Smoothing* is applied to account for unseen data (assign small probabilities for unseen observation symbols)

Blocking / indexing

- Number of possible links equals the product of the sizes of the two data sets to be linked
- Performance bottleneck in a record linkage system is usually the (expensive) evaluation of similarity measures between record pairs
- Blocking / indexing techniques are used to reduce the large amount of record comparisons
- *Febri* contains (currently) three indexing methods
 - Standard blocking
 - Sorted neighbourhood approach
 - Fuzzy blocking using *n*-grams (e.g. bigrams)

Record pair classification

- For each record pair compared a vector containing *matching weights* is calculated

Example:

Record A: ['dr', 'peter', 'paul', 'miller']

Record B: ['mr', 'pete', '', 'miller']

Matching weights: [0.2, 0.8, 0.0, 2.4]

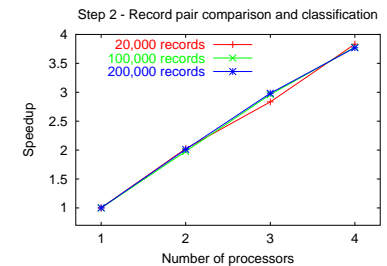
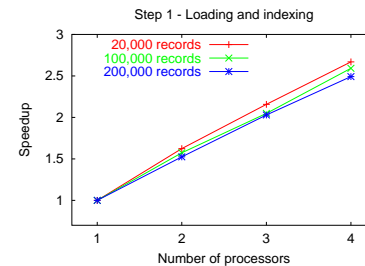
- Matching weights are used to classify record pairs as *links*, *non-links*, or *possible links*
- Fellegi & Sunter* classifier simply sums all the weights, then uses two thresholds to classify
- Improved classifiers are possible (for example using machine learning techniques)

Data set generation

- Difficult to acquire data for testing and evaluation (as record linkage deals with names and addresses)
- Also, linkage status is often not known (hard to evaluate and test new algorithms)
- Febri* contains a data set generator
 - Uses frequency tables for given- and surname, street name and type, suburb, postcode, age, etc.
 - Uses dictionaries of known misspellings
 - Duplicate records* are created via random introduction of modifications (like insert/delete/transpose characters, swap field values, delete values, etc.)

Parallelisation

- Implemented transparently to the user
- Currently using *MPI* via Python module *PyPar*
- Use of super-computing centres is problematic (privacy) → Alternative: *In-house office clusters*
- Some initial performance results (on *Sun SMP*)



Data set generation – Example

- Data set with 4 original and 6 duplicate records

REC_ID,	ADDRESS1,	ADDRESS2,	SUBURB
rec-0-org,	wylly place,	pine ret vill,	taree
rec-0-dup-0,	wyllyplace,	pine ret vill,	taree
rec-0-dup-1,	pine ret vill,	wylly place,	taree
rec-0-dup-2,	wylly place,	pine ret vill,	tared
rec-0-dup-3,	wylly parade,	pine ret vill,	taree
rec-1-org,	stuart street,	hartford,	menton
rec-2-org,	griffiths street,	myross,	kilda
rec-2-dup-0,	griffith sstreet,	myross,	kilda
rec-2-dup-1,	griffith street,	mycross,	kilda
rec-3-org,	ellenborough place,	kalkite homestead,	sydney

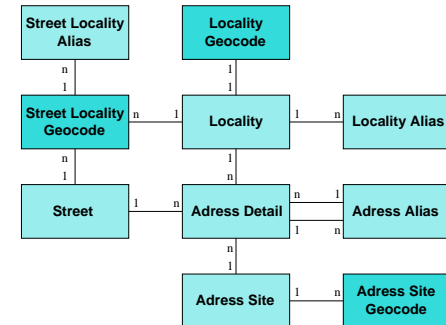
- Each record is given a unique identifier, which allows the evaluation of accuracy and error rates for record linkage

Geocoding

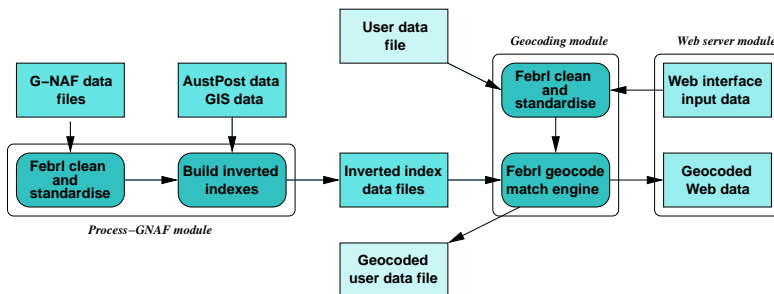
- The process of matching addresses with geographic locations (longitude and latitude)
- Geocoding tasks
 - Preprocess the geocoded reference data (cleaning, standardisation and indexing)
 - Clean and standardise the user addresses
 - (Fuzzy) match of user addresses with the reference data
 - Return location and match status
- Match status: address, street or locality level
- Geocode reference data used: *G-NAF*

Geocoded national address file

- G-NAF: Available since early 2004 (PSMA, <http://www.g-naf.com.au/>)
- Source data from 13 organisations (around 32 million source records)
- Processed into 22 normalised database tables



Febri geocoding system



- Only NSW G-NAF data available (around 4 million address, 58,000 street and 5,000 locality records)
- Additional Australia Post and GIS data used (for data imputing and to compute *neighbouring regions*)

Outlook

- Several research areas
 - Improving probabilistic data standardisation
 - New and improved blocking / indexing methods
 - Apply machine learning techniques for record pair classification
 - Improve performances (scalability and parallelism)
- Project web page
<http://datamining.anu.edu.au/linkage.html>

Febri is an ideal experimental platform to develop, implement and evaluate new data standardisation and record linkage algorithms and techniques