Data Science for Linguists

Session 4: Data Cleaning and Preparation

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Data Cleaning and Preparation

- in data science, data preparation (cleaning, transforming, and rearranging of real-world data) often comprises the majority of the work; a common figure is about 80% of an analyst's time
- ad hoc processing of data using a variety of tools (plain Python, Perl, R, Java, sed, awk) is still very common in science, but this approach has many disadvantages
- using Pandas in a Jupyter notebook for these tasks has the advantages of
 - ▷ a high-level, flexible, and fast set of tools that is very accessible to other researchers



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NaN and None in Pandas

- due to their underlying implementation in NumPy, Pandas series represent missing values by means of different **sentinel values**, the choice of which depends on the datatype:
 - by for data with dtype float64, Pandas uses the floating-point value NaN (Not a Number) which is defined by the ISO standard (accessible via np.nan)
 - bython objects (including variable-length strings), it uses the special value None from standard Python (which is very slow to compute with!)
 - because these two options are the only ones available, series of integer datatypes are silently converted into floating point numbers as soon as missing values are inserted
- one of the goals of Pandas is to make working with missing data efficient and smooth
 - there is well-defined default behaviour for any functions executed on Pandas series with missing values; such values will never cause Pandas to throw an exception, though the defalt behaviour might not be extremely useful in some cases
 - many methods (such as the ones for descriptive statistics) are implemented to silently exclude missing data per default
 - it also provides a range of methods which abstract over the somewhat inconsistent underlying implementation of null values

Pandas Nullable Dtypes

- to add support for true integer arrays with missing data, Pandas provides nullable dtypes like pd.Int32 (distinguished from the default dtypes like pd.int32 by capitalisation)
- the null value in series of these types is represented by pd.NA
- the other null values will be normalised without triggering implicit typecasting:

 for more efficient handling of large amounts of string data, there is the specialised extension type pd.StringDtype()

Detecting Null Values

- null values of any kind can be detected using the method data.isnull(), which returns a
 Boolean mask over the data, with True in all positions where there is a null value
- data.notnull() returns a Boolean mask as well, but the opposite of the result of the previous method (False in all positions where there is a null value)
- data.isna() and data.notna() are aliases of data.isnull() and data.notnull(), both sentinel values are matched by both methods (unlike in the underlying implementation)

Dropping Null Values with dropna()

- data.dropna() is a convenience method which combines filtering for and then dropping the null values, i.e. a shorthand for data[data.notna()]
- if data is a Series, the result will simply be a shorter series
- if data is a DataFrame, we can only drop entire rows or columns
 - by default, data.dropna() will drop all rows which contain any null value
 - b to drop columns instead, we can provide the argument axis=1 or axis="columns"
- argument how="all" to only drop rows/columns which consist entirely of null values
- for more fine-grained control (e.g. only weeding out the most gappy records), we can also provide an argument thresh=k to specify that any row/column with at least knon-null values will be kept

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Filling Null Values with fillna()

- data.fillna(x) is a convenience method which combines filtering for and then overwriting
 the null values with a default value x, i.e. a shorthand for data[data.isna()] = x
- x can be a dictionary providing different fill values by column index
- data.fillna(method="ffill") specifies a forward fill, i.e. empty values will be replaced by the previous non-null value (which therefore gets propagated forward)
 - b example: a series with data [1 <NA> 2 <NA> <NA> 3] will become [1 1 2 2 2 3]
- data.fillna(method="bfill") specifies a backward fill, i.e. empty values will be replaced by the subsequent non-null value (which therefore gets propagated backward)
 - b example: a series with data [1 <NA> 2 <NA> <NA> 3] will become [1 2 2 3 3 3]
- if no previous or subseqent value is available, ffill and bfill leave null values at the fringes!
- for a DataFrame, we can again switch to propagation through columns by specifying axis=1



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Removing Duplicates

- data.duplicated() returns a Boolean Series indicating whether each row is a duplicate (all values equal to some previous row) or not
- data.drop_duplicates() returns a Series or DataFrame consisting of only those rows in data where data.duplicated() was False
- the subset argument allows to provide a list of column indices to specify which columns are relevant for duplicate detection (other columns are allowed to have different values)
- if the subset argument is used, the first variant of each duplicate will be used for the values in the irrelevant columns, keep="last" changes this

Replacing Values

- data.replace(oldvals, newvals) substitutes a set of values by replacements

 - b data.replace([-2, -1], 0) sets all cells with value -2 or -1 to 0
 - \triangleright data.replace([-2, -1], [-1, 0]) sets cells with value -2 or -1, and all with -1 to 0
- data.replace(dictionary) replaces each occurrence of a key with its value
- more complex element-wise transformation can be implemented as a function
 (e.g. transform_value(x)), and then executed on every cell by a call to
 data.map(transform value); this works with anonymous functions (lambda) as well

Renaming Axis Indexes

- the axes can be modified in place by executing the map method of their indices:
- data.rename() allows to create a transformed version of a dataset without modifying the original (simple example: data.rename(index=str.title, columns=str.upper))



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Discretisation and Binning

- continuous data is often discretised or otherwise separated into bins for analyis
- pd.cut(data, k) returns a Categorical object which describes the k equal-length bins computed on the basis of the minimum and maximum values within the data
- pd.cut(data, cutoffs) returns a Categorical object which describes the bins computed from the data based on the specified cutoff values between the bins
- pd.qcut(data, k) bins the data into k quantiles (equally sized bins)
- pd.qcut(data, quantiles) bins the data into the provided quantiles (between 0 and 1)
- the argument labels=group names allows to override the default bin names
- a Categorical object cats has the following key applications:
 - > cats.codes returns an array containing the bin index for each datapoint
 - cats.categories shows an IntervalIndex object representing the bins
 - pd.value counts(cats) renders the counts of datapoints in each bin
- these functions will have central importance in Session 7 (aggregation and grouping)



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Detecting and Filtering Outliers

- outlier detection and filtering is typically performed by combinations of simple array operations
- for a normally distributed Series of data, we might define our outliers as all rows where the value exceeds 3 in absolute value: data[data.abs() > 3]
- in a DataFrame full of normally distibuted values, we might be interested in the rows were any of the columns has such a value: data[(data.abs() > 3).any(axis="columns")] (Boolean DataFrame generated by the comparison, on which we apply the any() method)
- to remove the outliers by capping values to the range [-3, 3], we can just do data[data.abs() > 3] = np.sign(data) * 3





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Permutation and Random Sampling

- permutations are sampled using smp = np.random.permutation(k)
- data.take(smp) (= iloc-based indexing) is then a random permutation of the first k lines
- data.sample(n=k) selects a random subset of k rows without replacement
- data.sample(n=k, replace=True) samples k rows with replacement





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Vectorised String Operations: Equivalents of Basic Methods

- cleaning up a messy dataset often requires a lot of string manipulation, but simple element-wise application using data.map() will fail on the null values
- the Pandas Series offers array-oriented and null-aware string operations which are accessible via the str attribut; here is a small sample:
 - data.str.count to count occurrences of a pattern
 - b data.str.contains(s) returns a Boolean mask with the result of calls to s in value on each cell value, mixed with NaN values wherever a value was missing
 - data.str.len to compute the length of each string
 - data.str.strip to trim whitespace (including newlines) from both sides

Vectorised String Operations: Regular Expressions

- there are also vectorised and null-aware versions of the regex capabilities:
 - > matches = data.str.findall(r"somePattern")
 - p first_matches = matches.str.get(1)
- data.str.extract(pattern) returns the captured groups of the regular expression as a new DataFrame

Vectorised String Operations: Miscellaneous Methods

- several methods emulate the capabilities of Python string operations:
 - data.str.cat for element-wise concatenation (with optional delimiter)
 - data.str.get to index each string
 - data.str.repeat to repeat each string
 - data.str.slice for extracting slices from each string
- data.str.get and data.str.slice are also available through the normal indexing syntax (data.str[i] and data.str[i:j])

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Assignment 4: Tasks

- 1) Read up on the CoNLL-U format for dependency treebanks.
- 2) Create a new Jupyter notebook and import pandas. Load the development set of the Basque UD corpus from the file eu_bdt-ud-dev.conllu into a DataFrame object, filtering out lines which are empty or start with #. Specify the CoNLL-U field names as a column index.
- 3) Convert all values consisting of an underscore into an appropriate missing data type, and remove all columns where more than 80% of the values are empty.
- 4) Reduce the dataset to all rows representing forms of the auxiliary *izan* ("to be").
- 5) Apply vectorised string operations which involve regular expressions to convert the morphological features into a more useful format (one new column per feature, feature values in each row; example: column Number[abs] with values Sing and Plur).
- 6) Query the database to see whether there is any syncretism in the paradigm of *izan*.
- (i.e. same form, different morphological features) [Hint: duplicate detection on different subsets]
- 7) How often does each form of *izan* occur in this development set? Create a new dataframe consisting of the form, the columns with the morphological features, and these counts.
- 8) Split the forms of *izan* attested in the corpus into ten bins of equal size, so that each bin groups together forms of roughly equal frequency. Detect and remove outliers if necessary.

Preliminary Course Plan

- 1 27/10 IPython and Jupyter
- 2 03/11 Introduction to NumPy
- 3 10/11 Pandas and Data Frames
- 4 17/11 Data Cleaning and Preparation
- 5 24/11 Linguistic Preprocessing
- 6 01/12 Data Wrangling: Join, Combine, Reshape
- 7 08/12 Data Aggregation and Grouping
- 8 15/12 Visualisation with Seaborn
- 9 22/12 Modeling and Prediction
- 10 12/01 Classification
- 11 19/01 Clustering
- 12 26/01 Pattern Extraction and Density Estimation
- 13 02/02 Statistical Inference
- 14 09/02 Data Science Projects

Questions

Questions?

Comments?

Suggestions?