

Bioinformatical problem solving with Python



Wednesdays 17:30-19:00, M801

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Not covered topics so far

- Error handling (exceptions)
- Numerical and scientific libraries (SciPy, NumPy, pandas, etc.)
- Plotting libraries (Matplotlib)
- Software design

- Data structures with labeled axes supporting automatic data alignment
- Integrated time series functionality
- Arithmetic operations pass on the axis labels
- Flexible handling of missing data
- pandas data structures: Series and DataFrame from pandas import Series, DataFrame import pandas as pd

- One-dimensional array-like container with associated array of data labels (index).
- Default index (0 ... N-1):
 obj = Series([4, 7, -5, 3])
 obj.values → array([4, 7, -5, 3])
 obj.index → RangeIndex(start=0, stop=4, step=1)
- User-defined index:
 obj2 = Series([4, 7, -5, 3], index=['d', 'b', 'a', 'c'])
 obj2.values → array([4, 7, -5, 3])
 obj2.index → Index([d, b, a, c], dtype=object)

Series can be constructed directly from a Python dictionary:

```
sdata = {'Ohio': 35000, 'Texas': 71000, 'Oregon': 16000}
obj3 = Series(sdata)
obj3.values → array([35000, 71000, 16000])
obj3.index → Index([Ohio, Texas, Oregon], dtype=object)
```

Values can be accessed by their indices:

```
obj3['Ohio'] → 35000
obj3[['Texas', 'Oregon']]
obj3['Oregon'] = 25000
obj3['Utah'] = 5000
```

Given the two Series objects

Create a new Series object with the added values for each location. What happens with the locations that are only present in one of the objects?

• We can bring different Series object in line by calling the 'reindex' method with the union of the two indices: new_index = obj1.index.union(obj2.index) obj3 = obj1.reindex(new_index, fill_value=0) obj4 = obj2.reindex(new_index, fill_value=0) obj3 + obj4

• Alternatively, we can use the 'add' method of Series objects with the 'fill_value' argument: obj1.add(obj2, fill_value=0) There are other useful methods to summarize Series objects:

```
obj1.sum() → 127000
obj1.min() → 5000
obj1.max() → 71000
obj1.mean() → 31750.0
obj1.std() → 28952.54738
```

- Two-dimensional data structure containing an ordered collection of columns (Series).
- Columns can be of different type.
- A DataFrame has both a row and a column index.

DataFrame objects can be constructed from a dict of equal-length lists:

Specifying a list of column names will put the columns in the desired order:

```
df2 = DataFrame(data, columns=['year', 'state', 'pop'],
index=['one', 'two', 'three', 'four', 'five'])
```

Columns can be accessed by dict-like notation or attribute:

```
df['state']
df.year
```

Rows can be accessed with the 'ix' indexing field:

```
df.ix[1]
  df2.ix['three']
or by slicing/boolean indexing the object:
  df[:2]
  df2[data['year'] > 2001]
```

Subset of rows and columns with the 'ix' field: df2.ix['three', ['year', 'state']] Columns can be modified by assigning scalars, lists/arrays, or Series objects:

```
df['size'] = 6000
df['debt'] = [5.2, 6.7, 8.3, 9.0, 1.2]
df2['growth'] = Series([-1.2, 3.0, 2.7], index=['two', 'four', 'five'])
```

- The 'del' keyword removes columns: del df2['growth']
- The 'T' attribute can be used to transpose DataFrame objects.
- The 'sort_index' method can be used to sort DataFrame objects.

Using the two Series objects from the first exercise, create a DataFrame using appropriate column labels. Create a third column containing the mean of the first two columns. Create a new DataFrame containing only the rows with means smaller than 20000.