

Programming Principles in Python (CSCI 503/490)

Data

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pandas

- Contains high-level data structures and manipulation tools designed to make data analysis fast and easy in Python
- Built on top of NumPy
- Built with the following requirements:
 - Data structures with labeled axes (aligning data)
 - Support time series data
 - Do arithmetic operations that include metadata (labels)
 - Handle missing data
 - Add merge and relational operations

Series

- A one-dimensional array (with a type) with an **index**
- Index defaults to numbers but can also be text (like a dictionary)
- Allows easier reference to specific items
- `obj = pd.Series([7, 14, -2, 1])`
- Basically two arrays: `obj.values` and `obj.index`
- Can specify the index explicitly and use strings
- `obj2 = pd.Series([4, 7, -5, 3],
index=['d', 'b', 'a', 'c'])`
- Kind of like fixed-length, ordered dictionary + can create from a dictionary
- `obj3 = pd.Series({'Ohio': 35000, 'Texas': 71000,
'Oregon': 16000, 'Utah': 5000})`

Data Frame

- A dictionary of Series (labels for each series)
- A spreadsheet with row keys (the index) and column headers
- Has an index shared with each series
- Allows easy reference to any cell
- ```
df = DataFrame({'state': ['Ohio', 'Ohio', 'Ohio', 'Nevada'],
 'year': [2000, 2001, 2002, 2001],
 'pop': [1.5, 1.7, 3.6, 2.4]})
```
- Index is automatically assigned just as with a series but can be passed in as well via index kwarg
- Can reassign column names by passing columns kwarg

# DataFrame Access and Manipulation

---

- `df.values` → 2D NumPy array
- Accessing a column:
  - `df["<column>"]`
  - `df.<column>`
  - Both return Series
  - Dot syntax only works when the column is a valid identifier
- Assigning to a column:
  - `df["<column>"] = <scalar>` # all cells set to same value
  - `df["<column>"] = <array>` # values set in order
  - `df["<column>"] = <series>` # values set according to match  
# between df and series indexes

# Indexing

---

- Same as with NumPy arrays but can use Series's index labels
- Slicing with labels: NumPy is **exclusive**, Pandas is **inclusive**!
  - `s = Series(np.arange(4))`  
`s[0:2]` # gives two values like numpy
  - `s = Series(np.arange(4), index=['a', 'b', 'c', 'd'])`  
`s['a':'c']` # gives three values, not two!
- Obtaining data subsets
  - `[...]`: get columns by label
  - `loc[...]`: get rows/cols by label
  - `iloc[...]`: get rows/cols by position (integer index)
  - For single cells (scalars), also have `at` and `iat`



# Data Frame

```
df = pd.read_csv('penguins_lter.csv')
```

|     | studyName | Sample Number | Species                             | Region | Island    | Stage              | Individual ID | Clutch Completion | Date Egg | Culmen Length (mm) |
|-----|-----------|---------------|-------------------------------------|--------|-----------|--------------------|---------------|-------------------|----------|--------------------|
| 0   | PAL0708   | 1             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N1A1          | Yes               | 11/11/07 | 39.1               |
| 1   | PAL0708   | 2             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N1A2          | Yes               | 11/11/07 | 39.5               |
| 2   | PAL0708   | 3             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N2A1          | Yes               | 11/16/07 | 40.3               |
| 3   | PAL0708   | 4             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N2A2          | Yes               | 11/16/07 | NaN                |
| 4   | PAL0708   | 5             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N3A1          | Yes               | 11/16/07 | 36.7               |
| ... | ...       | ...           | ...                                 | ...    | ...       | ...                | ...           | ...               | ...      | ...                |
| 339 | PAL0910   | 120           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N38A2         | No                | 12/1/09  | NaN                |
| 340 | PAL0910   | 121           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N39A1         | Yes               | 11/22/09 | 46.8               |
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| 343 | PAL0910   | 124           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N43A2         | Yes               | 11/22/09 | 49.9               |

344 rows x 17 columns



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```

Column Names

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Index

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344 rows x 17 columns

Column: df[ 'Island' ]



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```

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Row: df.loc[2]

Index

344 rows x 17 columns

Column: df['Island']



# Data Frame

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```

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| 4   | PAL0708   | 5             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N3A1          | Yes               | 11/16/07 | 36.7               |
| ... | ...       | ...           | ...                                 | ...    | ...       | ...                | ...           | ...               | ...      | ...                |
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Row: df.loc[2]

Index

Cell: df.loc[341, 'Species']

344 rows x 17 columns

Column: df['Island']

# Data Frame

```
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```

Column Names

|     | studyName | Sample Number | Species                             | Region | Island    | Stage              | Individual ID | Clutch Completion | Date Egg | Culmen Length (mm) |
|-----|-----------|---------------|-------------------------------------|--------|-----------|--------------------|---------------|-------------------|----------|--------------------|
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| 3   | PAL0708   | 4             | Adelie Penguin (Pygoscelis adeliae) | Anvers | Torgersen | Adult, 1 Egg Stage | N2A2          | Yes               | 11/16/07 | NaN                |
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| ... | ...       | ...           | ...                                 | ...    | ...       | ...                | ...           | ...               | ...      | ...                |
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Row: df.loc[2]

Index

Missing Data

Cell: df.loc[341, 'Species']

344 rows x 17 columns

Column: df['Island']



# Assignment 7

---

- Musical Artists Datasets
- Downloading and uncompressing files
- Finding files using OS libraries
- Load per-artist numpy arrays, each saved in the .npy format
- Store per-month dataframes, each in a csv file

# Arithmetic

---

- Add, subtract, multiply, and divide are element-wise like numpy
- ...but use labels to align
- ...and missing labels lead to NaN (not a number) values

```
In [28]: obj3
Out[28]:
Ohio 35000
Oregon 16000
Texas 71000
Utah 5000
dtype: int64
```

```
In [29]: obj4
Out[29]:
California NaN
Ohio 35000
Oregon 16000
Texas 71000
dtype: float64
```

```
In [30]: obj3 + obj4
Out[30]:
California NaN
Ohio 70000
Oregon 32000
Texas 142000
Utah NaN
dtype: float64
```

- also have `.add`, `.subtract`, ... that allow `fill_value` argument
- `obj3.add(obj4, fill_value=0)`

# Filtering

---

- Same as with numpy arrays but allows use of column-based criteria
  - `data[data < 5] = 0`
  - `data[data['three'] > 5]`
- `data < 5` → boolean data frame, can be used to select specific elements
- Multiple criteria, use `&`, `|`, and `~`; remember parentheses!
  - `data[(data['three'] > 5) & (data['two'] < 10)]`
- Also can check for missing values via `isna()/isnull()/notnull()`
  - `data[data['three'].notnull() & data['two'].isnull()]`

# Filtering

```
df[df['Culmen Length (mm)'] > 40]
```

|     | studyName | Sample Number | Species                             | Region | Island    | Stage              | Individual ID | Clutch Completion | Date Egg | Culmen Length (mm) |
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# Filtering

```
df[df['Culmen Length (mm)'] > 40]
```

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| 339 | PAL0910   | 120           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N38A2         | No                | 12/1/09  | NaN                |
| 340 | PAL0910   | 121           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N39A1         | Yes               | 11/22/09 | 46.8               |
| 341 | PAL0910   | 122           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N39A2         | Yes               | 11/22/09 | 50.4               |
| 342 | PAL0910   | 123           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N43A1         | Yes               | 11/22/09 | 45.2               |
| 343 | PAL0910   | 124           | Gentoo penguin (Pygoscelis papua)   | Anvers | Biscoe    | Adult, 1 Egg Stage | N43A2         | Yes               | 11/22/09 | 49.9               |

344 rows x 17 columns



# DataFrame Index

---

- Similar to index for Series
- Immutable
- Can be shared with multiple structures (DataFrames or Series)
- `in` operator works with: `'Ohio' in df.index`
- Can choose new index column(s) with `set_index()`
- `reindex` creates a new object with the data conformed to new index
  - `obj2 = obj.reindex(['a', 'b', 'c', 'd', 'e'])`
  - can fill in missing values in different ways

# Sorting

---

- `sort_values` method on series
  - `obj.sort_values()`
- Missing values (NaN) are at the end by default (`na_position` controls, can be first)
- `sort_values` on DataFrame:
  - `df.sort_values(<list-of-columns>)`
  - `df.sort_values(by=['a', 'b'])`
- Also a `sort_index` method to sort by the index
  - `df.sort_index()`

# Statistics

---

- `sum`: column sums (`axis=1` gives sums over rows)
- missing values are excluded unless the whole slice is `NaN`
- `idxmax`, `idxmin` are like `argmax`, `argmin` (return index)
- `describe`: shortcut for easy stats!

```
In [204]: df.describe()
```

```
Out[204]:
```

|       | one      | two       |
|-------|----------|-----------|
| count | 3.000000 | 2.000000  |
| mean  | 3.083333 | -2.900000 |
| std   | 3.493685 | 2.262742  |
| min   | 0.750000 | -4.500000 |
| 25%   | 1.075000 | -3.700000 |
| 50%   | 1.400000 | -2.900000 |
| 75%   | 4.250000 | -2.100000 |
| max   | 7.100000 | -1.300000 |

```
In [205]: obj = Series(['a', 'a', 'b', 'c'] * 4)
```

```
In [206]: obj.describe()
```

```
Out[206]:
```

|        |        |
|--------|--------|
| count  | 16     |
| unique | 3      |
| top    | a      |
| freq   | 8      |
| dtype: | object |

# Unique Values and Value Counts

---

- `unique()` returns an array with only the unique values (no index)
  - `s = Series(['c', 'a', 'd', 'a', 'a', 'b', 'b', 'c', 'c'])`  
`s.unique()` # `array(['c', 'a', 'd', 'b'])`
- Also `nunique()` to count number of unique entries
- Data Frames use `drop_duplicates`
- `value_counts` returns a Series with index frequencies:
  - `s.value_counts()` # `Series({'c': 3, 'a': 3, 'b': 2, 'd': 1})`

# Reading & Writing Data in Pandas

| Format | Data Description                     | Reader         | Writer       |
|--------|--------------------------------------|----------------|--------------|
| text   | <a href="#">CSV</a>                  | read_csv       | to_csv       |
| text   | Fixed-Width Text File                | read_fwf       |              |
| text   | <a href="#">JSON</a>                 | read_json      | to_json      |
| text   | <a href="#">HTML</a>                 | read_html      | to_html      |
| text   | Local clipboard                      | read_clipboard | to_clipboard |
|        | <a href="#">MS Excel</a>             | read_excel     | to_excel     |
| binary | <a href="#">OpenDocument</a>         | read_excel     |              |
| binary | <a href="#">HDF5 Format</a>          | read_hdf       | to_hdf       |
| binary | <a href="#">Feather Format</a>       | read_feather   | to_feather   |
| binary | <a href="#">Parquet Format</a>       | read_parquet   | to_parquet   |
| binary | <a href="#">ORC Format</a>           | read_orc       |              |
| binary | <a href="#">Msgpack</a>              | read_msgpack   | to_msgpack   |
| binary | <a href="#">Stata</a>                | read_stata     | to_stata     |
| binary | <a href="#">SAS</a>                  | read_sas       |              |
| binary | <a href="#">SPSS</a>                 | read_spss      |              |
| binary | <a href="#">Python Pickle Format</a> | read_pickle    | to_pickle    |
| SQL    | <a href="#">SQL</a>                  | read_sql       | to_sql       |
| SQL    | <a href="#">Google BigQuery</a>      | read_gbq       | to_gbq       |

[[https://pandas.pydata.org/pandas-docs/stable/user\\_guide/io.html](https://pandas.pydata.org/pandas-docs/stable/user_guide/io.html)]



# read\_csv

---

- Convenient method to read csv files
- Lots of different options to help get data into the desired format
- Basic: `df = pd.read_csv(fname)`
- Parameters:
  - `path`: where to read the data from
  - `sep` (or `delimiter`): the delimiter (`,`, `' '`, `'\t'`, `'\s+'`)
  - `header`: if `None`, no header
  - `index_col`: which column to use as the row index
  - `names`: list of header names (e.g. if the file has no header)
  - `skiprows`: number of list of lines to skip

# Writing CSV data with pandas

---

- Basic: `df.to_csv(<fname>)`
- Change delimiter with `sep` kwarg:
  - `df.to_csv('example.dsv', sep='|')`
- Change missing value representation
  - `df.to_csv('example.dsv', na_rep='NULL')`
- Don't write row or column labels:
  - `df.to_csv('example.csv', index=False, header=False)`
- Series may also be written to csv

# Handling Missing Data

---

| Argument             | Description                                                                                                                                 |
|----------------------|---------------------------------------------------------------------------------------------------------------------------------------------|
| <code>dropna</code>  | Filter axis labels based on whether values for each label have missing data, with varying thresholds for how much missing data to tolerate. |
| <code>fillna</code>  | Fill in missing data with some value or using an interpolation method such as <code>'ffill'</code> or <code>'bfill'</code> .                |
| <code>isnull</code>  | Return like-type object containing boolean values indicating which values are missing / NA.                                                 |
| <code>notnull</code> | Negation of <code>isnull</code> .                                                                                                           |

---

[W. McKinney, Python for Data Analysis]

# Derived Data

---

- Create new columns from existing columns
  - `r["PctFail"] = r['Fail'] / r['Total']`
  - `r = r.assign(PctFail= r['Fail'] / r['Total'])`
- Note that operations are computed in a vectorized manner
- Similarities to functional paradigm (map/filter):
  - specify the operation once
  - no loops
  - interpreted as an operation on the entire column

# inplace

---

- Generally, when we modify a data frame, we reassign:
  - `rdf = df.reset_index()`
  - This is usually very **efficient**
  - Allows for method chaining
- There are versions where you can do this "inplace":
  - `df.reset_index(inplace=True)`
  - This means **no reassignment**, but it isn't usually any faster nor better
  - Sometimes still creates a copy
  - Will likely be deprecated



# Documentation

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- pandas [documentation](#) is pretty good
- Lots of recipes on stackoverflow for particular data manipulations/queries