

Pos: DATA WRANGLING



CONTENTS

In this course we learn to read data from different sources and transform the data using Pandas dataframe techniques and more including:

Transforming data using apply and map functions Transforming data using dfply verbs



CHEAT SHEETS

NumPy Basics Cheat Sheet

https://www.datacamp.com/community/blog/python-numpy-cheat-sheet

Pandas Basics Cheat Sheet

https://www.datacamp.com/community/blog/python-pandas-cheat-sheet

Pandas Data Wrangling Cheat Sheet

https://www.datacamp.com/community/blog/pandas-cheat-sheet-python



DATA WRANGLING

from wikipedia:

Data munging or data wrangling is loosely the process of manually converting or mapping data from one "raw" form into another format that allows for more convenient consumption of the data with the help of semi-automated tools.

In almost every dataset, it is necessary to perform some data munging before you can use the data to perform eg. a classification task. Therefore, as a data scientist, you'll spend a lot of time munging and wrangling the data!



DATA MANIPULATION IN PYTHON

Let us first look at munging techniques available in basic Python:

Renaming columns
Adding or removing columns
Reordering columns
Merging data frames
Finding and removing duplicates



LOAD IRIS DATASET AS DATAFRAME

The iris dataset is included in the sklearn.datasets. However, the dataset was changed in scikit-learn version 0.20 and fixed two wrong data points according to Fisher's paper. The new version is the same as in R, but not as in the UCI Machine Learning Repository.

```
In [77]: # let's import a few libraries
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
```



```
In [79]: # https://stackoverflow.com/questions/38105539/how-to-convert-a-scikit-learn-dataset-to-a-pandas-dataset
# if you'd like to check dataset type use: type(load_iris())
# if you'd like to view list of attributes use: dir(load_iris())

# save load_iris() sklearn dataset to iris_bunch
iris_bunch = load_iris()

# list of attributes
dir(iris_bunch)

Out[79]: ['DESCR',
   ''data',
   'data' module',
   'feature_names',
   'filename',
   'frame',
   'target',
   ''target_names']
```



```
In [80]: # so what's in iris_bunch?
         # sample of the data
         iris_bunch.data[1:10,]
Out[80]: array([[4.9, 3. , 1.4, 0.2],
                [4.7, 3.2, 1.3, 0.2],
                [4.6, 3.1, 1.5, 0.2],
                [5., 3.6, 1.4, 0.2],
                [5.4, 3.9, 1.7, 0.4],
                [4.6, 3.4, 1.4, 0.3],
                [5., 3.4, 1.5, 0.2],
                [4.4, 2.9, 1.4, 0.2],
                [4.9, 3.1, 1.5, 0.1]
In [81]: # list with feature names
         iris_bunch.feature_names
Out[81]: ['sepal length (cm)',
           'sepal width (cm)',
          'petal length (cm)',
           'petal width (cm)']
```





In [84]: df_iris.head()

Out[84]:

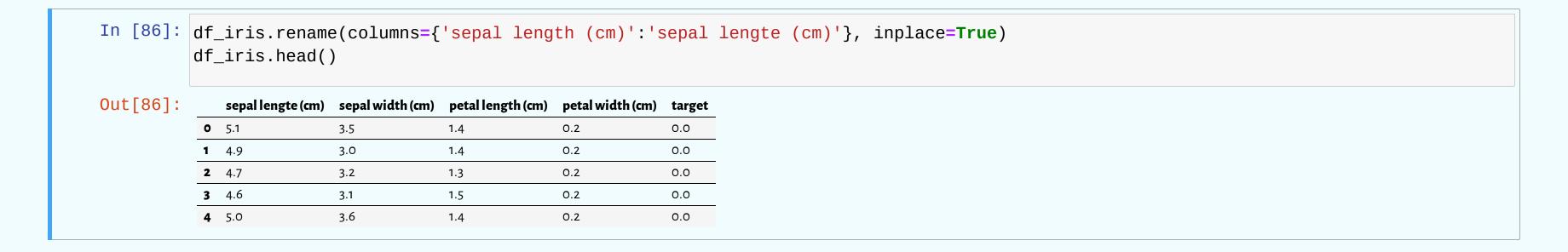
	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
0	5.1	3.5	1.4	0.2	0.0
1	4.9	3.0	1.4	0.2	0.0
2	4.7	3.2	1.3	0.2	0.0
3	4.6	3.1	1.5	0.2	0.0
4	5.0	3.6	1.4	0.2	0.0



RENAMING COLUMNS IN A DATA FRAME

You can do any of the following with Python's built-in functions. Note that these modify iris dataframe directly; that is, you don't have to save the result back into df_iris when using the "inplace" argument with a value set to True.

RENAME THE COLUMN NAMED "SEPAL LENGTH (CM)" TO "SEPAL LENGTE (CM)"





RENAME BY INDEX IN NAMES LIST: CHANGE THIRD ITEM, "PETAL LENGTH (CM)", TO "PETAL LENGTE (CM)"

In [87]: | df_iris.rename(columns={ df_iris.columns[2]: "petal lengte (cm)" }, inplace = True) df_iris.head() Out[87]: sepal lengte (cm) sepal width (cm) petal lengte (cm) petal width (cm) target **o** 5.1 3.5 1.4 0.0 3.0 0.2 **1** 4.9 1.4 0.0 **2** 4.7 3.2 0.2 1.3 0.0 **3** 4.6 3.1 1.5 0.2 0.0 3.6 **4** 5.0 1.4 0.2 0.0



SHOW ALL COLUMN NAMES AS A LIST, INCLUDING THE MODIFIED COLUMN NAMES



ADDING AND REMOVING COLUMNS IN A DATA FRAME

There are many different ways of adding columns to a dataframe and removing columns from a data frame.



DIFFERENT WAYS TO ADD A COLUMN TO A DATAFRAME

```
In [90]: # By declaring a new list as a column
# Declare a list that is to be converted into a column
size = ["small", "large", "medium"]

# Using 'size' as the column name
# and equating it to the list
df['size1'] = size
df
Out[90]: id weight size

O 1 20 small
1 2 27 large
2 3 24 medium
```





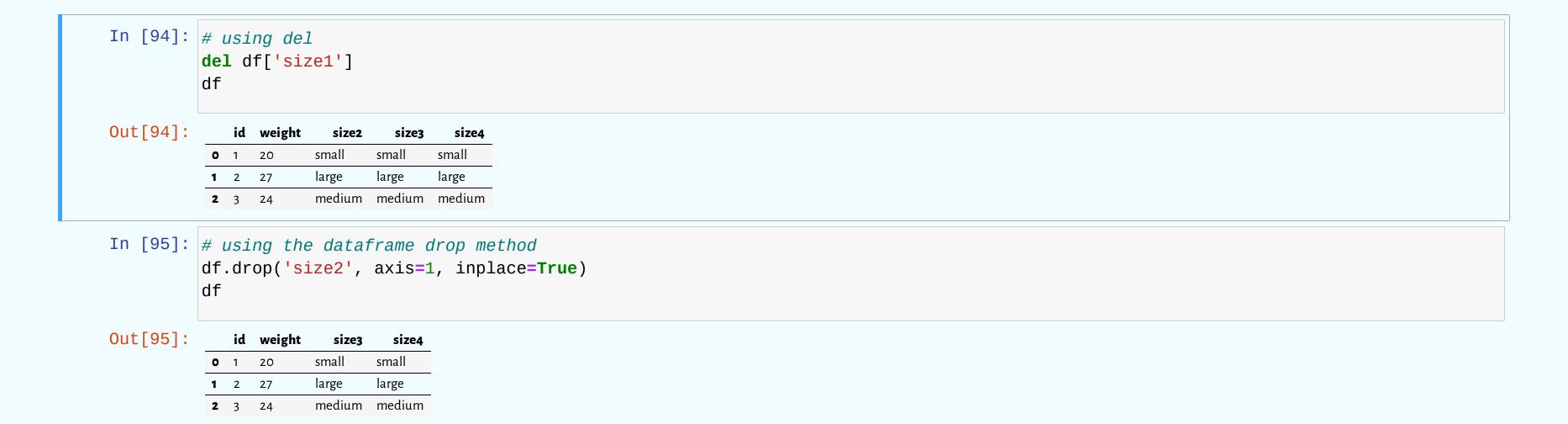
```
In [92]: # Using Dataframe.assign() method
         # This will create a new dataframe with the new column added to the old dataframe.
         # Using 'size3' as the column name and equating it to the list
         df = df.assign(size3 = ["small", "large", "medium"])
          df
Out[92]:
             id weight size1 size2
                                  size3
                            small
                                  small
                     small
          0 1 20
                     large
                                  large
          1 2 27
                            large
                     medium medium medium
          2 3 24
```



```
In [93]: # Using a Python dictionary to add a column
         # Define a dictionary with key values of an existing column and their respective
          # value pairs as the values for our new column.
          size4 = {1: 'small', 2: 'large', 3: 'medium'}
          df['size4'] = df['id'].map(size4)
          df
Out[93]:
             id weight
                        size1
                              size2
                                     size3
                                           size4
                            small
                                  small
                                         small
           0 1 20
                      small
                      large
                            large
          1 2 27
                                   large
                                          large
          2 3 24
                      medium medium medium medium
```



DIFFERENT WAYS TO REMOVE A COLUMN FROM A DATAFRAME





```
In [96]: # by listing the columns you would like to keep
         # and dropping any columns not listed
         df=df[['id','size4']]
         df
Out[96]:
            id size4
          o 1 small
          1 2 large
          2 3 medium
In [97]: # The pop method is used to remove the specified column from the DataFrame and return the removed column as a pandas Series.
         removed_col=df.pop('size4')
         removed_col
Out[97]: 0
               small
               large
              medium
```

Name: size4, dtype: object





REORDERING COLUMNS IN A DATA FRAME

```
In [99]: # Create a sample dataframe as we did earlier
         # Define a dictionary containing data
         data = {
                 'id': [1,2,3],
                 'weight': [20,27,24],
                 'size': ["small", "large", "medium"]
         # Convert the dictionary into DataFrame
         df = pd.DataFrame(data)
         df
Out[99]:
            id weight
                     size
                    small
          0 1 20
          1 2 27
                    large
                    medium
          2 3 24
```



REORDER BY COLUMN NAMES





Out[102]:

	id	weight	size
0	1	20	small
1	2	27	large
2	3	24	medium

In [103]: # get values in a row using the row index to select the row

df.iloc[0]

Out[103]: id 1

weight 20 size small

Name: 0, dtype: object



MERGING DATA FRAMES

You want to merge two data frames on a given column from each. Just like a join in SQL.



```
In [105]: # Make another data frame with the data and story numbers, including ratings
          data = {
                   'subject': [1,1,1,2,2,2],
                   'storyid': [1,2,3,2,3,1],
                   'rating': [6.7,4.5,3.7,3.3,4.1,5.2]
          rating_df = pd.DataFrame.from_dict(data)
          rating_df
Out[105]:
             subject storyid rating
           0 1
                         6.7
           1 1
                         4.5
           2 1
                         3.7
           3 2
                         3.3
           4 2
                   3
                         4.1
           5 2
                         5.2
```



```
In [106]: # Merge the two data frames
           # how='left' is the default (left join), so it is not required to provide it, but for clarity we do
          # Here we provide it in order to make the code more readable
           stories_df.merge(rating_df, on=['storyid'], how='left')
Out[106]:
              storyid title subject rating
                    lions 1
                               6.7
                    lions 2
                               5.2
           1 1
           2 2
                    tigers 1
                               4.5
           3 2
                    tigers 2
                               3.3
                    bears 1
                               3.7
            4 3
           5 3
                    bears 2
                               4.1
```



MERGING DATA FRAMES ON MORE THAN ONE COLUMN

```
In [107]: # Make up more data: create a dataframe with animals and a dataframe with observations
         data = {
                  'size': ['small', 'big', 'small', 'big'],
                 'type': ['cat', 'cat', 'dog', 'dog'],
                 'name': ['lynx', 'tiger', 'chihuahua', 'great dane']
         animals_df = pd.DataFrame.from_dict(data)
          print(animals_df)
              size type
                              name
          0 small cat
                           lynx
              big cat
                             tiger
          2 small dog
                        chihuahua
               big dog great dane
```



```
In [108]: data = {'number':[1,2,3,4], 'size': ['big', 'small', 'small', 'big'], 'type': ['cat', 'dog', 'dog']}
          observations_df = pd.DataFrame.from_dict(data)
          print(f"\n{observations_df}" )
          # Merge the dataframes
          observations_df.merge(animals_df, on=['size', 'type'], how='left')
             number
                       size type
                       big cat
                  2 small dog
                  3 small dog
                        big dog
Out[108]:
             number size type
                               name
                        cat tiger
           0 1
                   small dog chihuahua
           1 2
                   small dog chihuahua
                       dog great dane
```



FINDING AND REMOVING DUPLICATE RECORDS IN A SERIE

You want to find and/or remove duplicate entries from a serie or data frame.



```
In [111]: # For each element: is this one a duplicate (first instance of a particular value not counted)
    x_duplicates = x.duplicated()
    print(*x_duplicates)

# The values of the duplicated entries
    # note that '10' appears in the original vector four times, and so it has three entries here
    x_duplicates_values = x[x.duplicated()]
    print(*x_duplicates_values)
False False False False False False True False True False False True True True True True False False False False
10 10 8 14 19 10 8 7
```



```
In [112]: # create a list with duplicates
x_list = [1, 2, 3, 3, 4, 4, 5]

# a set is another python data structure: it can only contains unique items
# and therefore we can use it to remove duplicates and convert the set back to a list again
list(set(x_list))
Out[112]: [1, 2, 3, 4, 5]
```



FINDING AND REMOVING DUPLICATE RECORDS IN DATA FRAMES

```
In [113]: # Create a sample data frame:
    data = {
        'label': ['A', 'B', 'C', 'B', 'B', 'A', 'A'],
        'type': [4,3,6,3,1,2,4,4]
}
```



```
In [114]: df = pd.DataFrame.from_dict(data)
          print(df)
          # Is each row a repeat?
          x_duplicates = df.duplicated()
          x_duplicates
            label type
                Α
                     1
Out[114]: 0
               False
               False
               False
               True
               False
               False
               True
               True
          dtype: bool
```



```
In [115]: df_duplicates_values = df[df.duplicated()]

Out[115]: label type
3 B 3
6 A 4
7 A 4

In [116]: # Show unique repeat entries
df_duplicates_values_unique = df_duplicates_values.drop_duplicates()
df_duplicates_values_unique

Out[116]: label type
3 B 3
6 A 4
```



```
In [117]: # Original data with repeats removed. These do the same:
         df_unique1 = df.drop_duplicates()
         df_unique1
Out[117]:
            label type
          o A
               3
          2 C
               6
          4 B
          5 A 2
In [118]: df_unique2 = df[~df.duplicated()]
         df_unique2
Out[118]:
            label type
          o A 4
          1 B
               3
          2 C
          4 B
          5 A 2
```



INTRODUCING APPLY

At any Python Q&A site, you'll frequently see an exchange like this one:

Q: How can I use a loop to [...insert task here...]? A: Don't. Use one of the apply functions or the Python list comprehension

So, what are these wondrous apply functions and list comprehensions and how do they work? I think the best way to figure out anything in Python is to learn by experimentation, using embarrassingly trivial data and functions.

Let's examine some of those.



APPLY_ALONG_AXIS

Description: Returns a series or array or list of values obtained by applying a function to margins of an array or matrix.

OK. We know about series/arrays and functions, but what are these "margins"?

Simple: either the rows (0), the columns (1) or both. By both, we mean apply the function to each individual value.







```
In [122]: # Divide all values by 2
# notice that here the axis argument is simply ignored but a value has to be provided
div_2 = np.apply_along_axis(lambda x: x/2, 1, m)
print(div_2.T)

[[ 0.5     5.5]
        [ 1.     6. ]
        [ 1.5     6.5]
        [ 2.     7. ]
        [ 2.5     7.5]
        [ 3.     8. ]
        [ 3.5     8.5]
        [ 4.     9. ]
        [ 4.5     9.5]
        [ 5.     10. ]]
```



GROUPBY

To illustrate, we can load up the classic iris dataset, which contains a bunch of flower measurements:

```
In [123]: # As we saw before, the iris data is available in the seaborn package:
            import seaborn as sns
            # The following script loads the iris data set into a data frame
            df_iris_seaborn = sns.load_dataset('iris')
            df_iris_seaborn
Out[123]:
                 sepal_length sepal_width petal_length petal_width species
             o 5.1
                            3.5
                                                            setosa
             1 4.9
                            3.0
                                       1.4
                                                  0.2
                                                            setosa
             2 4.7
                            3.2
                                                  0.2
                                                            setosa
                                       1.3
             3 4.6
                                                  0.2
                                                            setosa
                            3.1
                                       1.5
                 5.0
                                                  0.2
                                                            setosa
              145 6.7
                            3.0
                                                  2.3
                                                            virginica
                                       5.2
                                                            virginica
             146 6.3
                            2.5
                                       5.0
                                                 1.9
                                                            virginica
             147 6.5
                                                  2.0
                            3.0
                                                            virginica
             148 6.2
                            3.4
                                                  2.3
                                                            virginica
             149 5.9
            150 rows × 5 columns
```



In [124]:	df_iris	_seaborn.	groupby('species').agg('me
Out[124]:		sepal_length	sepal_width	petal_length	petal_width
	species				
	setosa	5.006	3.428	1.462	0.246
	versicolor	5.936	2.770	4.260	1.326
	virginica	6.588	2.974	5.552	2.026

Essentially, groupby provides a way to split your data by factors and do calculations on each subset. It returns an object of class "groupby" and there are many more complex ways to use it.



MAP

Description: map returns a map object of the same length as X, each element of which is the result of applying FUN to the corresponding element of X

Maps are lazily evaluated, meaning the values are only computed on-demand.

A simple example:

```
In [125]: import statistics
1 = [list(range(1,11)), list(range(11,21))]
mean_map=map(statistics.mean, 1)
print(list(mean_map))

[5.5, 15.5]
```



```
In [126]: # the sum of the values in each element
    sum_map=map(sum, 1)
    print(list(sum_map))

[55, 155]

In [127]: # Notice: alternatively, you can also use list comprehension:
    sum_list=[sum(i) for i in 1]
    print(sum_list)

[55, 155]
```



```
In [128]: # calculate multiple values
mult_funcs = [min, statistics.median, max]

mean_sum_list = [[x(i) for x in mult_funcs] for i in 1]
print(mean_sum_list)

[[1, 5.5, 10], [11, 15.5, 20]]
```



REPLICATE WITH NUMPY

NumPy is a package for working with numerical data and used for data science and scientific computing.

An example: let's simulate 5 normal distributions, each with 5 observations:



REPLICATE A FUNCTION ON ELEMENTS WITH MAP

With map we can apply a function to multiple list arguments

```
In [130]: one_to_four = range(1, 5)
    four_to_one = range(4, 0, -1)
    rep = lambda value, times: [value]*times
    list(map(rep, one_to_four, four_to_one))

Out[130]: [[1, 1, 1, 1], [2, 2, 2], [3, 3], [4]]
```



DFPLY

The dfply package makes it possible to do R's dplyr-style data manipulation with pipes in Python on pandas DataFrames.

This is an alternative to pandas-ply and dplython, which both engineer dplyr syntax and functionality in Python. There are probably more packages that attempt to enable dplyr-style dataframe manipulation in Python, but those are the two I am aware of.

dfply uses a decorator-based architecture for the piping functionality and to "categorize" the types of data manipulation functions. The goal of this architecture is to make dfply concise and easily extensible, simply by chaining together different decorators that each have a distinct effect on the wrapped function.

dfply is intended to mimic the functionality of dplyr. The syntax is the same for the most part, but will vary in some cases as Python is a considerably different programming language than R.

from <u>kieferk/dfply</u>



DFPLY

When working with data you must:

- Figure out what you want to do.
- Precisely describe what you want in the form of a computer program.
- Execute the code.



The dfply package makes each of these steps as fast and easy as possible by:

- Elucidating the most common data manipulation operations, so that your options are helpfully constrained when thinking about how to tackle a problem.
- Providing simple functions that correspond to the most common data manipulation verbs, so that you can easily translate your thoughts into code.
- Using efficient data storage backends, so that you spend as little time waiting for the computer as possible.



DATA: NYCFLIGHTS13

To explore the basic data manipulation verbs of dfply, we'll start with the built-in nycflights13 data frame. This dataset contains all 336776 flights that departed from New York City in 2013. The data comes from the <u>US Bureau of Transporation Statistics</u>

```
In [132]: from dfply import *
    import pandas as pd
    flight_data = pd.read_csv('./datasets/flights.csv')
    flight_data.shape
Out[132]: (336776, 19)
```



In [133]: flight_data.head()

Out[133]:

:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
	0	2013	1	1	517.0	515	2.0	830.0	819	11.0	UA	1545	N14228	EWR	IAH	227.0	1400	5	15	2013-01-01T10:00:00Z
-	1	2013	1	1	533.0	529	4.0	850.0	830	20.0	UA	1714	N24211	LGA	IAH	227.0	1416	5	29	2013-01-01T10:00:00Z
	2	2013	1	1	542.0	540	2.0	923.0	850	33.0	AA	1141	N619AA	JFK	MIA	160.0	1089	5	40	2013-01-01T10:00:00Z
-	3	2013	1	1	544.0	545	-1.0	1004.0	1022	-18.0	B6	725	N804JB	JFK	BQN	183.0	1576	5	45	2013-01-01T10:00:00Z
	4	2013	1	1	554.0	600	-6.0	812.0	837	-25.0	DL	461	N668DN	LGA	ATL	116.0	762	6	0	2013-01-01T11:00:00Z



SINGLE TABLE VERBS

Dfly aims to provide a function for each basic verb of data manipulating:

- filter_by() (and row_slice())
- arrange()
- select() (and rename())
- distinct()
- mutate() (and transmute())
- summarize()
- sample()



In [134]: # dfply works directly on pandas DataFrames, chaining operations on the data with the >> operator

flight_data >> head(5)

Out[134]:

	y	/ear	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
	0 2	2013	1	1	517.0	515	2.0	830.0	819	11.0	UA	1545	N14228	EWR	IAH	227.0	1400	5	15	2013-01-01T10:00:00Z
_	1 2	2013	1	1	533.0	529	4.0	850.0	830	20.0	UA	1714	N24211	LGA	IAH	227.0	1416	5	29	2013-01-01T10:00:00Z
	2 2	2013	1	1	542.0	540	2.0	923.0	850	33.0	AA	1141	N619AA	JFK	MIA	160.0	1089	5	40	2013-01-01T10:00:00Z
_	3 2	2013	1	1	544.0	545	-1.0	1004.0	1022	-18.0	B6	725	N804]B	JFK	BQN	183.0	1576	5	45	2013-01-01T10:00:00Z
	4 2	2013	1	1	554.0	600	-6.0	812.0	837	-25.0	DL	461	N668DN	LGA	ATL	116.0	762	6	0	2013-01-01T11:00:00Z



In [135]: # The DataFrame as it is passed through the piping operations is represented by the symbol X
flight_data >> select(X.dep_time, X.origin) >> head(3)

Out[135]:

	dep_time	origin
0	517.0	EWR
1	533.0	LGA
2	542.0	JFK



ARRANGE ROWS WITH ARRANGE()

arrange() works similarly to filter() except that instead of filtering or selecting rows, it reorders them. If you provide more than one column name, each additional column will be used to break ties in the values of preceding columns:

ıt[136]:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
	0	2013	1	1	517.0	515	2.0	830.0	819	11.0	UA	1545	N14228	EWR	IAH	227.0	1400	5	15	2013-01-01T10:00:00Z
	1	2013	1	1	533.0	529	4.0	850.0	830	20.0	UA	1714	N24211	LGA	IAH	227.0	1416	5	29	2013-01-01T10:00:00Z
	2	2013	1	1	542.0	540	2.0	923.0	850	33.0	AA	1141	N619AA	JFK	MIA	160.0	1089	5	40	2013-01-01T10:00:00Z
	3	2013	1	1	544.0	545	-1.0	1004.0	1022	-18.0	B6	725	N804JB	JFK	BQN	183.0	1576	5	45	2013-01-01T10:00:00Z
	Λ	2013	1	1	554.0	600	-6.0	812.0	837	-25.0	DL	461	N668DN	LGA	ATL	116.0	762	6	0	2013-01-01T11:00:00



Use desc() to order a column in descending order:

[n [137]:	flight	_da	ta >>	arr	ange(X.	arr_delay,	ascendi	.ng =Fa1	se) >> hea	d(5)										
Out[137]:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hou
	7072	2013	1	9	641.0	900	1301.0	1242.0	1530	1272.0	НА	51	N384HA	JFK	HNL	640.0	4983	9	0	2013-01-09T14:00:00Z
	235778	2013	6	15	1432.0	1935	1137.0	1607.0	2120	1127.0	MQ	3535	N504MQ	JFK	СМН	74.0	483	19	35	2013-06-15T23:00:00Z
	8239	2013	1	10	1121.0	1635	1126.0	1239.0	1810	1109.0	MQ	3695	N517MQ	EWR	ORD	111.0	719	16	35	2013-01-10T21:00:00Z
	327043	2013	9	20	1139.0	1845	1014.0	1457.0	2210	1007.0	AA	177	N338AA	JFK	SFO	354.0	2586	18	45	2013-09-20T22:00:00
	270376	2013	7	22	845.0	1600	1005.0	1044.0	1815	989.0	MQ	3075	N665MQ	1FK	CVG	96.0	589	16	0	2013-07-22T20:00:00



FILTER ROWS WITH FILTER_BY()

filter_by() allows you to select a subset of the rows of a data frame. The first and subsequent arguments are filtering expressions evaluated in the context of that data frame.

For example, we can select all flights on February 3rd with:

In [138]:	flight	_da	ta >>	fil	.ter_by(X.month==2,	X.day=	= 3) >>	head(5)											
Out[138]:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hou
	112904	2013	2	3	31.0	2359	32.0	458.0	437	21.0	B6	727	N506JB	JFK	BQN	190.0	1576	23	59	2013-02-04T04:00:00
	112905	2013	2	3	536.0	540	-4.0	927.0	850	37.0	AA	1141	N ₅ EBAA	JFK	MIA	164.0	1089	5	40	2013-02-03T10:00:00
	112906	2013	2	3	545.0	540	5.0	1049.0	1017	32.0	B6	725	N571JB	JFK	BQN	203.0	1576	5	40	2013-02-03T10:00:00
	112907	2013	2	3	556.0	600	-4.0	833.0	815	18.0	FL	345	N923AT	LGA	ATL	113.0	762	6	0	2013-02-03T11:00:002
	112908	2013	2	3	556.0	600	-4.0	935.0	912	23.0	B6	135	N508]B	JFK	RSW	163.0	1074	6	0	2013-02-03T11:00:002



MORE FILTER() AND SLICE()

This is equivalent as it can be done using pandas:

n [139]:	flight ₋	_data	[(fl	ight_da	ata["month"]] == 1)	(fli	ght_data["	day"] =	= 1)]	.head	1(5)							
ut[139]:	year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
	0 2013	1	1	517.0	515	2.0	830.0	819	11.0	UA	1545	N14228	EWR	IAH	227.0	1400	5	15	2013-01-01T10:00:00Z
	1 2013	1	1	533.0	529	4.0	850.0	830	20.0	UA	1714	N24211	LGA	IAH	227.0	1416	5	29	2013-01-01T10:00:00Z
	2 2013	1	1	542.0	540	2.0	923.0	850	33.0	AA	1141	N619AA	JFK	MIA	160.0	1089	5	40	2013-01-01T10:00:00Z
	3 2013	1	1	544.0	545	-1.0	1004.0	1022	-18.0	B6	725	N804JB	JFK	BQN	183.0	1576	5	45	2013-01-01T10:00:00Z
	4 2013	1	1	554.0	600	-6.0	812.0	837	-25.0	DL	461	N668DN	IGA	ATL	116.0	762	6	0	2013-01-01T11:00:00Z



To select rows by position, use row_slice(). You can pass single integer indices or a list of indices to select rows as with. This is going to be the same as using pandas' .iloc.

In [140]:	fli	ght_	_data	>>	row_sli	ce([0,7,10])													
Out[140]:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
	0	2013	1	1	517.0	515	2.0	830.0	819	11.0	UA	1545	N14228	EWR	IAH	227.0	1400	5	15	2013-01-01T10:00:00Z
	7	2013	1	1	557.0	600	-3.0	709.0	723	-14.0	EV	5708	N829AS	LGA	IAD	53.0	229	6	0	2013-01-01T11:00:00Z
	10	2013	1	1	558.0	600	-2.0	849.0	851	-2.0	B6	49	N793]B	JFK	PBI	149.0	1028	6	0	2013-01-01T11:00:00Z



SELECT COLUMNS WITH SELECT()

Often you work with large datasets with many columns where only a few are actually of interest to you. select() allows you to rapidly zoom in on a useful subset using operations that usually only work on numeric variable positions:

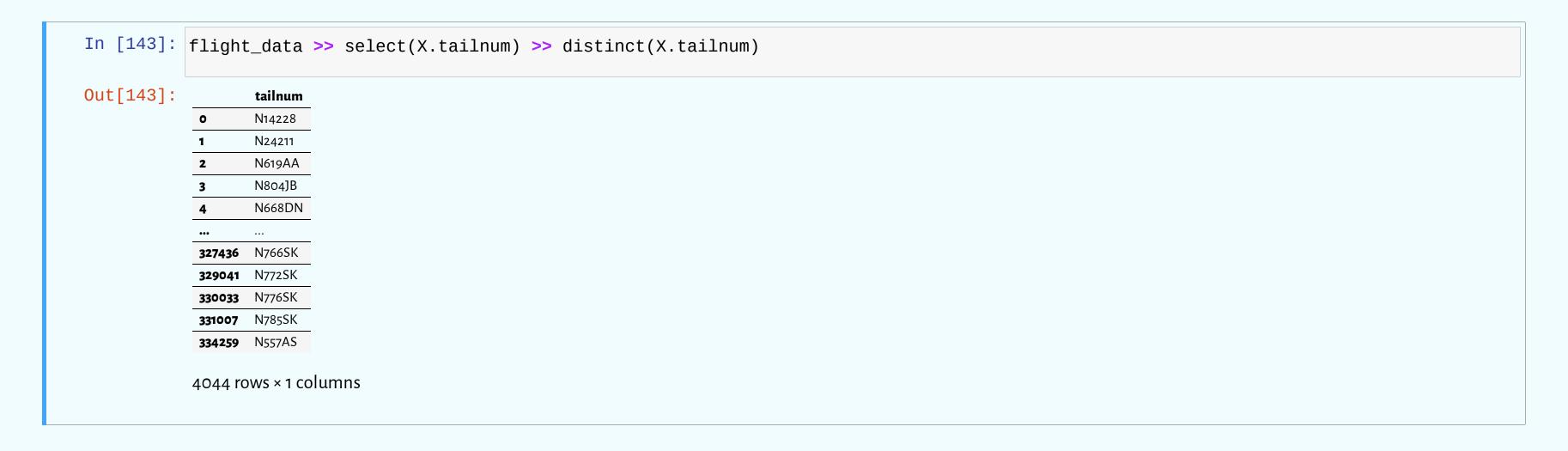


In [142]: # The select function accept string labels, integer positions, and/or symbolically represented column names (X.column) flight_data >> select(X.year, ['month', 'day'],6) >> head(10) Out[142]: year month day arr_time **0** 2013 1 830.0 850.0 **1** 2013 1 **2** 2013 1 923.0 **3** 2013 1 1 1004.0 **4** 2013 1 812.0 **5** 2013 1 1 740.0 **6** 2013 1 1 913.0 709.0 **7** 2013 1 **8** 2013 1 838.0 **9** 2013 1 1 753.0



EXTRACT DISTINCT (UNIQUE) ROWS

A common use of select() is to find out which values a set of variables takes. This is particularly useful in conjunction with the distinct() verb which only returns the unique values in a table.





GROUPED OPERATIONS

These verbs are useful, but they become really powerful when you combine them with the idea of "group by", repeating the operation individually on groups of observations within the dataset.



The verbs are affected by grouping as follows:

- grouped select() is the same as ungrouped select(), except that grouping variables are always retained.
- grouped arrange() orders first by grouping variables
- mutate() and filter() are most useful in conjunction with window functions (like rank(), or min(x) == x), and are described in detail in vignette("window-function").
- sample() samples the specified number/fraction of rows in each group.
- row_slice() extracts rows within each group.
- summarize() and summarize_each() are easy to understand and very useful, and is described in more detail below.



EXAMPLE GROUP BY

In the following example, we split the complete dataset into individual planes and then summarise each plane by counting the number of flights (count = n()) and computing the average distance (distance_mean=X.distance.mean()) and delay (arr_delay_mean=X.arr_delay.mean()).



```
In [144]: delay_df = (flight_data >>
    group_by(X.tailnum) >>
    summarize(tailnum_count=n(X.tailnum), distance_mean=X.distance.mean(), arr_delay_mean=X.arr_delay.mean()) >>
    filter_by(X.tailnum_count >20, X.distance_mean<2000)
)
delay_df >> head(5)
```

Out[144]:

	tailnum	tailnum_count	distance_mean	arr_delay_mean
1	NoEGMQ	371	676.188679	9.982955
2	N10156	153	757.947712	12.717241
3	N102UW	48	535.875000	2.937500
4	N103US	46	535.195652	-6.934783
5	N104UW	47	535.255319	1.804348
•••				
4038	N997DL	63	867.761905	4.903226
4039	N998AT	26	593.538462	29.960000
4040	N998DL	77	857.818182	16.394737
4041	N999DN	61	895.459016	14.311475
4042	N9EAMQ	248	674.665323	9.235294

2961 rows × 4 columns

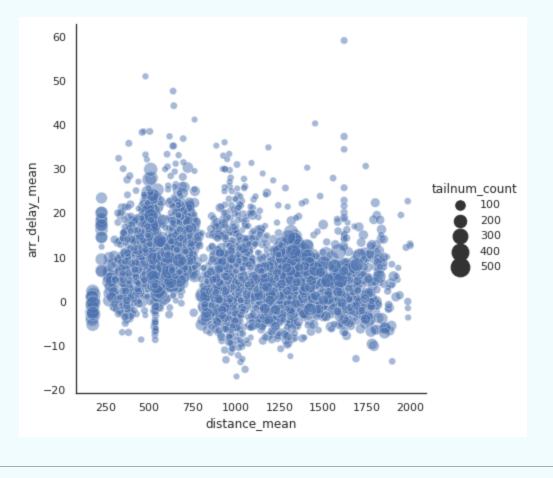


PLOT DELAYS

In [145]: # import the seaborn package for making a visualisation
import seaborn as sns
sns.set_theme(style="white")



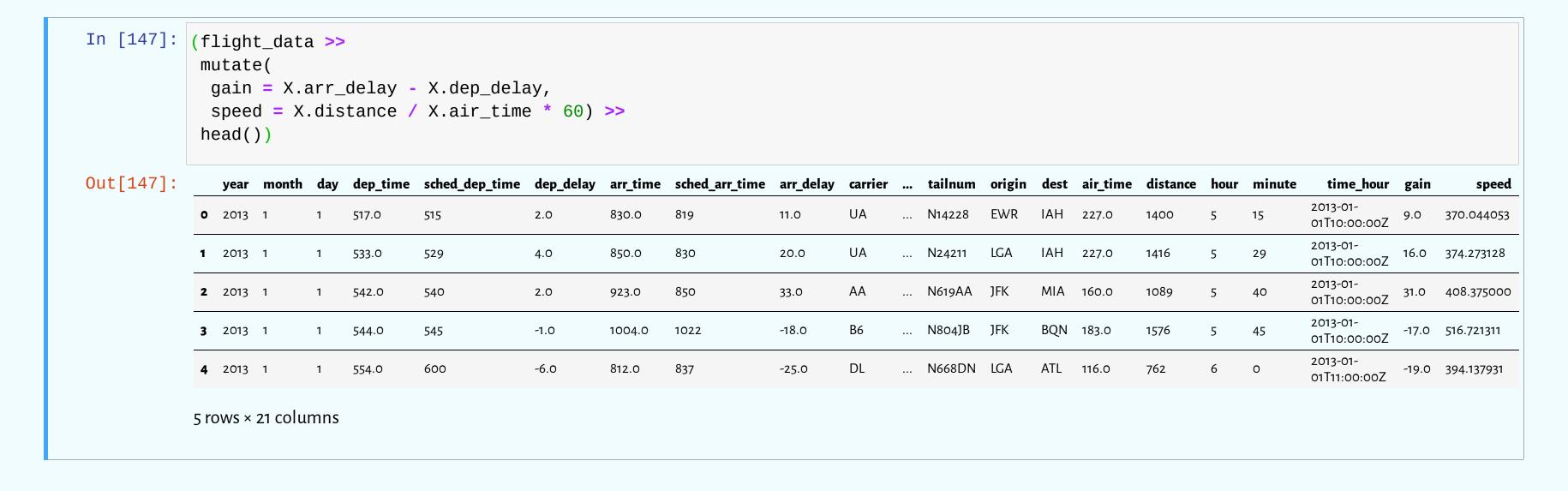
Out[146]: <seaborn.axisgrid.FacetGrid at 0x7fb4308028e0>





ADD NEW COLUMNS WITH MUTATE()

As well as selecting from the set of existing columns, it's often useful to add new columns that are functions of existing columns. This is the job of mutate():





COMBINE DATASETS

- Mutating joins
- Filtering joins
- Set operations
- Binding



RANDOMLY SAMPLE ROWS WITH SAMPLE()

You can use sample() to take a random sample of rows, either a fixed number for sample(n=3, replace=True) or a fixed fraction for sample(frac=0.0002, replace=False).

[n [148]:	flight	t_da1	ta >>	sam	ple(n=3	, replace=1	True)													
out[148]:		year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hou
	328986	2013	9	22	1707.0	1715	-8.0	1937.0	2015	-38.0	AA	2488	N4YHAA	EWR	DFW	181.0	1372	17	15	2013-09-22T21:00:00
	75368	2013	11	21	2055.0	2015	40.0	2201.0	2126	35.0	B6	418	N337JB	JFK	BOS	44.0	187	20	15	2013-11-22T01:00:002
	1046	2013	1	2	845.0	842	3.0	1102.0	1059	3.0	EV	4255	N34110	EWR	CHS	108.0	628	8	42	2013-01-02T13:00:002



In [149]: flight_data >> sample(frac=0.00002, replace=True)

Out[149]:

	year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hour	minute	time_hour
112393	2013	2	2	858.0	900	-2.0	1440.0	1540	-60.0	НА	51	N388HA	JFK	HNL	629.0	4983	9	0	2013-02-02T14:00:00Z
150503	2013	3	16	907.0	910	-3.0	1023.0	1032	-9.0	В6	20	N284JB	JFK	ROC	53.0	264	9	10	2013-03-16T13:00:00Z
214142	2013	5	23	839.0	755	44.0	1023.0	920	63.0	MQ	3737	N3AEMQ	EWR	ORD	111.0	719	7	55	2013-05-23T11:00:00Z
238871	2013	6	18	2050.0	2000	50.0	2239.0	2127	72.0	9E	4287	N836AY	JFK	BWI	39.0	184	20	0	2013-06-19T00:00:00Z
101732	2013	12	20	2158.0	2112	46.0	2343.0	2312	31.0	US	1859	N182UW	EWR	CLT	81.0	529	21	12	2013-12-21T02:00:00Z
82075	2013	11	29	1624.0	1621	3.0	1831.0	1844	-13.0	UA	485	N831UA	EWR	DEN	229.0	1605	16	21	2013-11-29T21:00:00Z
20793	2013	1	24	1927.0	1920	7.0	2343.0	2315	28.0	DL	1345	N3771K	JFK	PHX	308.0	2153	19	20	2013-01-25T00:00:00Z



COMMONALITIES

You may have noticed that all these functions are very similar:

- A data frame is piped to the function as input
- The arguments describe what to do with it, and you can refer to columns in the data frame directly.
- The result is a new data frame

Together these properties make it easy to chain together multiple simple steps to achieve a complex result.