

Welcome to Programming for Data Science

Welcome to the course manual for CSC310 at URI.

This website will contain the syllabus, class notes and other reference material for the class.

[Course Calendar with zoom links on BrightSpace](#)



Tip

[subscribe to that calendar](#) in your favorite calendar application

Syllabus

Welcome to CSC/DSP310: Programming For Data Science.

In this syllabus you will find an overview of the course, information about your instructor, course policies, restatements of URI policies, reminders of relevant resources, and a schedule for the course.

About

About this course

Data science exists at the intersection of computer science, statistics, and machine learning. That means writing programs to access and manipulate data so that it becomes available for analysis using statistical and machine learning techniques is at the core of data science. Data scientists use their data and analytical ability to find and interpret rich data sources; manage large amounts of data despite hardware, software, and bandwidth constraints; merge data sources; ensure consistency of datasets; create visualizations to aid in understanding data; build mathematical models using the data; and present and communicate the data insights/findings.

This course provides a survey of data science. Topics include data driven programming in Python; data sets, file formats and meta-data; descriptive statistics, data visualization, and foundations of predictive data modeling and machine learning; accessing web data and databases; distributed data management. You will work on weekly substantial programming problems such as accessing data in database and visualize it or build machine learning models of a given data set.

Basic programming skills (CSC201 or CSC211) are a prerequisite to this course. This course is a prerequisite course to machine learning, where you learn how machine learning algorithms work. In this course, we will start with a very fast review of basic programming ideas, since you've already done that before. We will learn how to use machine learning algorithms to do data science, but not how to *build* machine learning algorithms, we'll use packages that implement the algorithms for us.

About this semester

This semester is a lot of new things for all of us. This course will be completely online all semester, so we will get to use a single instructional format all semester, including when all campus activities move remote after Thanksgiving. I recognize that those last two weeks of the semester may change your obligations with siblings, parents, work, etc. In light of that, we will cover all of the most important topics and you will have the opportunity to achieve all of the course learning outcomes before Thanksgiving. The material in the last two weeks of the semester will be more advanced, likely interesting and definitely useful material, but if your ability to participate in class is less at that time, it will not hurt your grade.

About this syllabus

This syllabus is a *living* document and accessible from BrightSpace, as a pdf for download directly online at rhodyprog4ds.github.io/BrownFall20/syllabus. If you choose to download a copy of it, note that it is only a copy. You can get notification of changes from GitHub by "watching" the [repository](#). You can view the date of changes and exactly what changes were made on the Github [commits](#) page.

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Creating an [issue on the repository](#) is also a good way to ask questions about anything in the course it will prompt additions and expand the FAQ section.

[Assignment 6: Naive Bayes](#)
[Assignment 7: Decision Trees](#)
[Assignment 8: Linear Regression](#)

About your instructor

Name: Dr. Sarah Brown Office hours: TBA via zoom, link in BrightSpace

Dr. Brown is a new Assistant Professor of Computer Science, who does research on how social context changes machine learning. Dr. Brown earned a PhD in Electrical Engineering from Northeastern University, completed a postdoctoral fellowship at University of California Berkeley, and worked as a postdoctoral research associate at Brown University before joining URI. At Brown University, Dr. Brown taught the Data and Society course for the Master's in Data Science Program.

The best way to contact me is e-mail or by dropping into my office hours. Please include [\[CSC310\]](#) or [\[DSP310\]](#) in the subject line of your email along with the topic of your message. This is important, because your messages are important, but I also get a lot of e-mail. Consider these a cheat code to my inbox: I have setup a filter that will flag your e-mail if you use one of those in the subject to ensure that I see it. I rarely check e-mail between 6pm and 9am, on weekends or holidays. You might see me post or send things during these hours, but I will not reliably see emails that arrive during those hours.

Note

Whether you use CSC or DSP does not matter.

Tools and Resources

We will use a variety of tools to conduct class and to facilitate your programming. You will need a computer with Linux, MacOS, or Windows. It is unlikely that a tablet or Chromebook will be able to do all of the things required in this course.

All of the tools below are either: - paid for by URI or - freely available online.

BrightSpace

This will be the central location from which you can access all other materials. Any links that are for private discussion among those enrolled in the course will be available only from our course [Brightspace site](#). This is also where your grades will appear.

Zoom

This is where we will meet for synchronous class sessions. You will find the link to class zoom sessions on Brightspace.

URI provides all faculty, staff, and students with a paid Zoom account. It *can* run in your browser or on a mobile device, but you will be able to participate in class best if you download the [Zoom client](#) on your computer. Please [log in](#) and [configure your account](#). Please add a photo of yourself to your account so that we can still see your likeness in some form when your camera is off. You may also wish to use a virtual background and you are welcome to do so.

Class will be interactive, so if you cannot be in a quiet place at class time, headphones with a built in microphone are strongly recommended.

For help, you can access the [instructions provided by IT](#).

Important

TL;DR [\[1\]](#)

- check Brightspace
- Install Zoom
- Setup your URI Zoom Account
- Log in to Prismia Chat
- Make a GitHub Account
- Install Python
- Install Git

Prismia chat

Our class link for Prismia chat is available on Brightspace. We will use this for chatting and in-class understanding checks.

On Prismia, all students see the instructor's messages, but only the Instructor and TA see student responses.

Course Manual

The course manual will have content including the class policies, scheduling, class notes, assignment information, and additional resources. This will be linked from Brightspace and available publicly online at [rhodyprog4ds.github.io/BrownFall20/](#). Links to the course reference text and code documentation will also be included here in the assignments and class notes.

GitHub Classroom

You will need a GitHub Account. If you do not already have one, please create one by the first day of class. There will be a link to our class GitHub Classroom on Brightspace.

Programming Environment

This a programming course, so you will need a programming environment. In order to complete assignments you need the items listed in the requirements list. The easiest way to meet these requirements is to follow the recommendations below. I will provide instruction assuming that you have followed the recommendations.

Requirements:

- Python with scientific computing packages (numpy, scipy, jupyter, pandas, etc)
- [Git](#)
- A web browser compatible with Jupyter Notebooks

Recommendation:

- Install python via [Anaconda](#)
- if you use Windows, install Git with [GitBash \(video instructions\)](#).
- if you use MacOS, install Git with the Xcode Command Line Tools. On Mavericks (10.9) or above you can do this by trying to run git from the Terminal the very first time.`git --version`

Optional:

- Text Editor: you may want a text editor outside of the Jupyter environment. Jupyter can edit markdown files (that you'll need for your portfolio), in browser, but it is more common to use a text editor like Atom or Sublime for this purpose.

 **Note**

all Git instructions will be given as instructions for the command line interface and GitHub specific instructions via the web interface. You may choose to use GitHub desktop or built in IDE tools, but the instructional team may not be able to help.

 **Note**

I use atom, but I decided to use it by downloading both Atom and Sublime and trying different things in each for a week. I liked Atom better after that and I've stuck with it since. I used Atom to write all of the content in this syllabus.

Textbook

The text for this class is a reference book and iwl not be a source of assignments. It will be a helpful reference and you may be directed there for answers to questions or alternate explanations of topics.

Python for Data Science is available free [online](#):

[1] Too long; didn't read.

Grading

This section of the syllabus describes the principles and mechanics of the grading for the course. This course will be graded on a basis of a set of *skills* (described in detail the next section of the syllabus). This is in contrast to more common grading on a basis of points earned through assignments.

Principles of Grading

Learning happens through practice and feedback. My goal as a teacher is for you to learn. The grading in this course is based on your learning of the material, rather than your completion of the activities that are assigned.

This course is designed to encourage you to work steadily at learning the material and demonstrating your new knowledge. There are no single points of failure, where you lose points that cannot be recovered. Also, you cannot cram anything one time and then forget it. The material will build and you have to demonstrate that you retained things.

- Earning a C in this class is intended to be easier than typical grading. I expect everyone to get at least a C.
- Earning a B in this class is intended to be very accessible, you can make a lot of mistakes along the way as you learn, as long as you learn by the end.
- Earning an A in this class will be challenging, but is possible even with making mistakes while you learn.

Grading this way also is more amenable to the fact that there are correct and incorrect ways to do things, but there is not always a single correct answer to a realistic data science problem. Your work will be assessed on whether or not it demonstrates your learning of the targeted skills. You will also receive feedback on how to improve.

How it works

There are 15 skills that you will be graded on in this course. While learning these skills, you will work through a progression of learning. Your grade will be based on earning 45 achievements that are organized into 15 skill groups with 3 levels for each.

These map onto letter grades roughly as follows:

- If you achieve level 1 in all of the skills, you will earn at least a C in the course.
- To earn a B, you must earn all of the level 1 and level 2 achievements.
- To earn an A, you must earn all of the achievements.

You will have at least three opportunities to earn every level 2 achievement. You will have at least two opportunities to earn every level 3 achievement. You will have three types of opportunities to demonstrate your current skill level: participation, assignments, and a portfolio.

Each level of achievement corresponds to a phase in your learning of the skill:

- To earn level 1 achievements, you will need to demonstrate basic awareness of the required concepts and know approximately what to do, but you may need specific instructions of which things to do or to look up examples to modify every step of the way. You can earn level 1 achievements in class, assignments, or portfolio submissions.
- To earn level 2 achievements you will need to demonstrate understanding of the concepts and the ability to apply them with instruction after earning the level 1 achievement for that skill. You can earn level 2 achievements in assignments or portfolio submissions.
- To earn level 3 achievements you will be required to consistently execute each skill and demonstrate deep understanding of the course material, after achieving level 2 in that skill. You can earn level 3 achievements only through your portfolio submissions.

Participation

While attending synchronous class sessions, there will be understanding checks and in class exercises. Completing in class exercises and correctly answering questions in class can earn level 1 achievements. In class questions will be administered through the classroom chat platform Prismia.chat; these records will be used to update your skill progression. You can also earn level 1 achievements from adding annotation to a section of the class notes.

Assignments

For your learning to progress and earn level 2 achievements, you must practice with the skills outside of class time.

Assignments will each evaluate certain skills. After your assignment is reviewed, you will get qualitative feedback on your work, and an assessment of your demonstration of the targeted skills.

Portfolio Checks

To earn level 3 achievements, you will build a portfolio consisting of reflections, challenge problems, and longer analyses over the course of the semester. You will submit your portfolio for review 4 times. The first two will cover the skills taught up until 1 week before the submission deadline.

The third and fourth portfolio checks will cover all of the skills. The fourth will be due during finals. This means that, if you have achieved mastery of all of the skills by the 3rd portfolio check, you do not need to submit the fourth one.

Portfolio prompts will be given throughout the class, some will be structured questions, others may be questions that arise in class, for which there is not time to answer.

TLDR

You could earn a C through in class participation alone, if you make nearly zero mistakes. To earn a B, you must complete assignments and participate in class. To earn an A you must participate, complete assignments, and build a portfolio.

⚠️ Warning

If you will skip an assignment, please accept the GitHub assignment and then close the Feedback pull request with a comment. This way we can make sure that you have support you need.

Detailed mechanics

On Brightspace there are 45 Grade items that you will get a 0 or a 1 grade for. These will be revealed, so that you can view them as you have an opportunity to demonstrate each one. The table below shows the minimum number of skills at each level to earn each letter grade.

Level 3 Level 2 Level 1

letter grade

letter grade	Level 3	Level 2	Level 1
A	15	15	15
A-	10	15	15
B+	5	15	15
B	0	15	15
B-	0	10	15
C+	0	5	15
C	0	0	15
C-	0	0	10
D+	0	0	5
D	0	0	3

For example, if you achieve level 2 on all of the skills and level 3 on 7 skills, that will be a B+.

If you achieve level 3 on 14 of the skills, but only level 1 on one of the skills, that will be a B-, because the minimum number of level 2 achievements for a B is 15. In this scenario the total number of achievements is 14 at level 3, 14 at level 2 and 15 at level 1, because you have to earn achievements within a skill in sequence.

The letter grade can be computed as follows

```
def compute_grade(num_level1,num_level2,num_level3):
    """
    Computes a grade for CSC/DSP310 from numbers of achievements at each level

    Parameters:
    -----
    num_level1 : int
        number of level 1 achievements earned
    num_level2 : int
        number of level 2 achievements earned
    num_level3 : int
        number of level 3 achievements earned

    Returns:
    -----
    letter_grade : string
        letter grade with modifier (+/-)
    """

    if num_level1 == 15:
        if num_level2 == 15:
            if num_level3 == 15:
                grade = 'A'
            elif num_level3 >= 10:
                grade = 'A-'
            elif num_level3 >=5:
                grade = 'B+'
            else:
                grade = 'B'
        elif num_level2 >=10:
            grade = 'B-'
        elif num_level2 >=5:
            grade = 'C+'
        else:
            grade = 'C'
    elif num_level1 >= 10:
        grade = 'C-'
    elif num_level1 >= 5:
        grade = 'D+'
    elif num_level1 >=3:
        grade = 'D'
    else:
        grade = 'F'

    return grade
```

Note

In this example, you will have also achieved level 1 on all of the skills, because it is a prerequisite to level 2.

```
compute_grade(15,15,15)
```

```
'A'
```

```
compute_grade(14,14,14)
```

```
'C-'
```

```
assert compute_grade(14,14,14) == 'C-'
```

```
assert compute_grade(15,15,15) == 'A'
```

```
assert compute_grade(15,15,11) == 'A-'
```

Late work

No late work will be graded. Every skill will be assessed through more than one assignment, so missing assignments occasionally may not hurt your grade. If you do not submit any assignments that cover a given skill, you may earn the level 2 achievement in that skill through a portfolio check, but you will not be able to earn the level 3 achievement in that skill.

Examples

Important

The following will make more sense after you read the next section of the syllabus and see the skills rubric sections.

Note

You may visit office hours to discuss assignments that you did not complete on time to get feedback and check your own understanding, but they will not count toward skill demonstration.

If you always attend and get everything correct, you will earn an A and you won't need to submit the 4th portfolio check or assignment 13.

Getting A Without Perfection

Map to an A

How Achievements were earned

	Level 1	Level 2	Level 3
python	A1	A3	P1
process	A1	P1	P2
access	week 2	A2	P1
construct	week 5	A5	P1
summarize	week 3	A3	P1
visualize	week 3	A3	P2
prepare	week 4	A5	P2
classification	A10	P2	P3
regression	week 8	A11	P2
clustering	week 9	A9	P3
evaluate	week 7	A11	P3
optimize	week 10	A11	P4
compare	week 11	A13	P3
unstructured	week 12	A13	P4
tools	week 11	A13	P3

Activity Legend

In class	Assignment	Portfolio Check
X	AX	PX

Other Activities

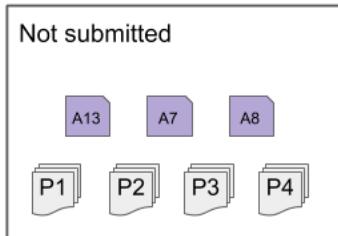
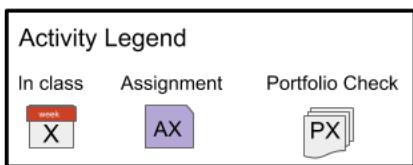
week 1	Attended, but did not understand
A4	Submitted, but incorrect
week 6	Missed class
A6	Not submitted
A7	Submitted, but incorrect
A8	Not submitted
A12	Not submitted
week 13	Attended, but all level 1 complete
week 14	Attended, but all level 1 complete

In this example the student made several mistakes, but still earned an A. This is the advantage to this grading scheme. For the `python`, `process`, and `classification` skills, the level 1 achievements were earned on assignments, not in class. For the `process` and `classification` skills, the level 2 achievements were not earned on assignments, only on portfolio checks, but they were earned on the first portfolio of those skills, so the level 3 achievements were earned on the second portfolio check for that skill. This student's fourth portfolio only demonstrated two skills: `optimize` and `unstructured`. It included only 1 analysis, a text analysis with optimizing the parameters of the model. Assignments 4 and 7 were both submitted, but didn't earn any achievements, the student got feedback though, that they were able to apply in later assignments to earn the achievements. The student missed class week 6 and chose to not submit assignment 6 and use week 7 to catch up. The student had too much work in another class and chose to skip assignment 8. The student tried assignment 12, but didn't finish it on time, so it was not graded, but the student visited office hours to understand and be sure to earn the level 2 `unstructured` achievement on assignment 13.

Getting a B with minimal work

Map to a B easily

	Level 1	Level 2	Level 3
python	week 1	A3	
process	week 1	A1	
access	week 2	A2	
construct	week 5	A5	
summarize	week 3	A3	
visualize	week 3	A3	
prepare	week 4	A4	
classification	week 10	A6	
regression	week 8	A11	
clustering	week 9	A9	
evaluate	week 7	A10	
optimize	week 10	A10	
compare	week 11	A11	
unstructured	week 12	A12	
tools	week 11	A12	

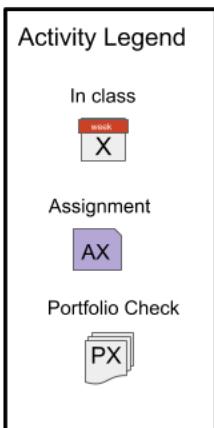


In this example, the student earned all level 1 achievements in class and all level 2 on assignments. This student was content with getting a B and chose to not submit a portfolio.

Getting a B while having trouble

Map to a B, having trouble

	Level 1	Level 2	Level 3
python	A1	P1	
process	A1	P2	
access	A2	P1	
construct	A5	P1	
summarize	A3	P1	
visualize	A3	P2	
prepare	A5	P2	
classification	A10	P3	
regression	A11	P2	
clustering	A9	P3	
evaluate	A11	P3	
optimize	A11	P4	
compare	A13	P3	
unstructured	A13	P4	
tools	A13	P3	



In this example, the student struggled to understand in class and on assignments. Assignments were submitted that showed some understanding, but all had some serious mistakes, so only level 1 achievements were earned from assignments. The student wanted to get a B and worked hard to get the level 2 achievements on the portfolio checks.

Learning Objective, Schedule, and Rubric

Learning Outcomes

There are five learning outcomes for this course.

1. (process) Describe the process of data science, define each phase, and identify standard tools
2. (data) Access and combine data in multiple formats for analysis
3. (exploratory) Perform exploratory data analyses including descriptive statistics and visualization
4. (modeling) Select models for data by applying and evaluating multiple models to a single dataset
5. (communicate) Communicate solutions to problems with data in common industry formats

We will build your skill in the **process** and **communicate** outcomes over the whole semester. The middle three skills will correspond roughly to the content taught for each of the first three portfolio checks.

Schedule

The course will meet MWF 1-1:50pm on Zoom. Every class will include participatory live coding (instructor types, students follow along) instruction and small exercises for you to progress toward level 1 achievements of the new skills introduced in class that day.

Programming assignments that will be due each week Tuesday by 11:59pm. until week 5 they were due Sundays

Note

On the [BrightSpace calendar](#) page you can get a feed link to add to the calendar of your choice by clicking on the subscribe (star) button on the top right of the page. Class is for 1 hour there because of Brightspace/zoom integration limitations, but that calendar includes the zoom link.

topics		skills
week		
1	[admin, python review]	process
2	Loading data, Python review	[access, prepare, summarize]
3	Exploratory Data Analysis	[summarize, visualize]
4	Data Cleaning	[prepare, summarize, visualize]
5	Databases, Merging DataFrames	[access, construct, summarize]
6	Modeling, Naive Bayes, classification performance metrics	[classification, evaluate]
7	decision trees, cross validation	[classification, evaluate]
8	Regression	[regression, evaluate]
9	Clustering	[clustering, evaluate]
10	SVM, parameter tuning	[optimize, tools]
11	KNN, Model comparison	[compare, tools]
12	Text Analysis	[unstructured]
13	Topic Modeling	[unstructured, tools]
14	Deep Learning	[tools, compare]

Skill Rubric

The skill rubric describes how your participation, assignments, and portfolios will be assessed to earn each achievement. The keyword for each skill is a short name that will be used to refer to skills throughout the course materials; the full description of the skill is in this table.

skill		Level 1	Level 2	Level 3
keyword				
python	pythonic code writing	python code that mostly runs, occasional pep8 adherence	python code that reliably runs, frequent pep8 adherence	reliable, efficient, pythonic code that consistently adheres to pep8
process	describe data science as a process	Identify basic components of data science	Describe and define each stage of the data science process	Compare different ways that data science can facilitate decision making
access	access data in multiple formats	load data from at least one format; identify the most common data formats	Load data for processing from the most common formats; Compare and contrast most common formats	access data from both common and uncommon formats and identify best practices for formats in different contexts
construct	construct datasets from multiple sources	identify what should happen to merge datasets or when they can be merged	apply basic merges	merge data that is not automatically aligned
summarize	Summarize and describe data	Describe the shape and structure of a dataset in basic terms	compute summary standard statistics of a whole dataset and grouped data	Compute and interpret various summary statistics of subsets of data
visualize	Visualize data	identify plot types, generate basic plots from pandas	generate multiple plot types with complete labeling with pandas and seaborn	generate complex plots with pandas and plotting libraries and customize with matplotlib or additional parameters
prepare	prepare data for analysis	identify if data is or is not ready for analysis, potential problems with data	apply data reshaping, cleaning, and filtering as directed	apply data reshaping, cleaning, and filtering manipulations reliably and correctly by assessing data as received
classification	Apply classification	identify and describe what classification is, apply pre-fit classification models	fit preselected classification model to a dataset	fit and apply classification models and select appropriate classification models for different contexts
regression	Apply Regression	identify what data that can be used for regression looks like	can fit linear regression models	can fit and explain nonlinear regression
clustering	Clustering	describe what clustering is	apply basic clustering	apply multiple clustering techniques, and interpret results
evaluate	Evaluate model performance	Explain basic performance metrics for different data science tasks	Apply basic model evaluation metrics to a held out test set	Evaluate a model with multiple metrics and cross validation
optimize	Optimize model parameters	Identify when model parameters need to be optimized	Manually optimize basic model parameters such as model order	Select optimal parameters based on multiple quantitative criteria and automate parameter tuning
compare	compare models	Qualitatively compare model classes	Compare model classes in specific terms and fit models in terms of traditional model performance metrics	Evaluate tradeoffs between different model comparison types

	skill	Level 1	Level 2	Level 3
	keyword			
unstructured	model unstructured data	Identify options for representing text data and use them once data is transformed	Apply at least one representation to transform unstructured data for model fitting or summarizing	apply multiple representations and compare and contrast them for different end results
tools	use industry standard data science tools and workflows to solve data science problems	Solve well structured problems with a single tool pipeline	Solve semi-structured, completely specified problems with multiple tools	Scope, choose an appropriate tool pipeline and solve data science problems

Assignments and Skills

Using the keywords from the table above, this table shows which assignments you will be able to demonstrate which skills and the total number of assignments that assess each skill. This is the number of opportunities you have to earn Level 2 and still preserve 2 chances to earn Level 3 for each skill.

	A1	A2	A3	A4	A5	A6	A7	A8	A9	A10	A11	A12	A13	# Assignments
keyword														
python	1	1	1	1	0	0	0	0	0	0	0	0	0	4
process	1	1	0	0	0	0	0	0	0	0	0	0	0	2
access	0	1	1	1	0	0	0	0	0	0	0	0	0	3
construct	0	0	0	0	1	1	0	0	0	0	0	0	0	2
summarize	0	0	1	1	1	1	1	1	1	1	1	1	1	11
visualize	0	0	1	1	0	1	1	1	1	1	1	1	1	10
prepare	0	0	0	1	1	0	0	0	0	0	0	0	0	2
classification	0	0	0	0	0	1	1	0	0	1	0	0	0	3
regression	0	0	0	0	0	0	0	1	0	0	1	0	0	2
clustering	0	0	0	0	0	0	0	0	1	0	1	0	0	2
evaluate	0	0	0	0	0	0	0	0	0	1	1	0	0	2
optimize	0	0	0	0	0	0	0	0	0	1	1	0	0	2
compare	0	0	0	0	0	0	0	0	0	0	1	0	1	2
unstructured	0	0	0	0	0	0	0	0	0	0	1	1	1	2
tools	0	0	0	0	0	0	0	0	0	1	1	1	1	4

Portfolios and Skills

The objective of your portfolio submissions is to earn Level 3 achievements. The following table shows what Level 3 looks like for each skill and identifies which portfolio submissions you can earn that Level 3 in that skill.

Level 3		P1	P2	P3	P4
keyword					
python	reliable, efficient, pythonic code that consistently adheres to pep8	1	1	0	0
process	Compare different ways that data science can facilitate decision making	0	1	1	0
access	access data from both common and uncommon formats and identify best practices for formats in different contexts	1	1	0	0
construct	merge data that is not automatically aligned	1	1	0	0
summarize	Compute and interpret various summary statistics of subsets of data	1	1	0	0
visualize	generate complex plots with pandas and plotting libraries and customize with matplotlib or additional parameters	1	1	0	0
prepare	apply data reshaping, cleaning, and filtering manipulations reliably and correctly by assessing data as received	1	1	0	0
classification	fit and apply classification models and select appropriate classification models for different contexts	0	1	1	0
regression	can fit and explain nonlinear regression	0	1	1	0
clustering	apply multiple clustering techniques, and interpret results	0	1	1	0
evaluate	Evaluate a model with multiple metrics and cross validation	0	1	1	0
optimize	Select optimal parameters based of mutiple quantitateve criteria and automate parameter tuning	0	0	1	1
compare	Evaluate tradeoffs between different model comparison types	0	0	1	1
unstructured	apply mulitple representations and compare and contrast them for different end results	0	0	1	1
tools	Scope, choose an appropriate tool pipeline and solve data science problems	0	0	1	1

Support

Academic Enhancement Center

Located in Roosevelt Hall, the AEC offers free face to face and web-based services to undergraduate students seeking academic support. Peer tutoring is available for STEM-related courses through drop-in centers and small group tutoring. The Writing Center offers peer tutoring focused on supporting undergraduate writers at any stage of a writing assignment. The UCS160 course and academic skills consultations offer students strategies and activities aimed at improving their studying and test-taking skills. Complete details about each of these programs, up-to-date schedules, contact information and self-service study resources are all available on the AEC website.

- **STEM Tutoring** helps students navigate 100 and 200 level math, chemistry, physics, biology, and other select STEM courses. The STEM Tutoring program offers free online and limited in-person peer-tutoring this fall. Undergraduates in introductory STEM courses have a variety of small group times to choose from and can select occasional or weekly appointments. Appointments and locations will be visible in the TutorTrac system on September 14th, 2020. The TutorTrac application is available through [URI Microsoft 365 single sign-on](#) and by visiting aec.uri.edu. More detailed information and instructions can be found on the AEC tutoring page.
- **Academic Skills Development** resources helps students plan work, manage time, and study more effectively. In Fall 2020, all Academic Skills and Strategies programming are offered both online and in-person. UCS160: Success in Higher Education is a one-credit course on developing a more effective approach to studying. Academic Consultations are 30-minute, 1 to 1 appointments that students can schedule on Starfish with Dr. David Hayes to address individual academic issues. Study Your Way to Success is a self-guided web portal connecting students to tips and strategies on studying and time management related topics. For more information on these programs, visit the Academic Skills Page or contact Dr. Hayes directly at davidhayes@uri.edu.
- The **Undergraduate Writing Center** provides free writing support to students in any class, at any stage of the writing process: from understanding an assignment and brainstorming ideas, to developing, organizing, and revising a draft. Fall 2020 services are offered through two online options: 1) real-time synchronous appointments with a peer consultant (25- and 50-minute slots, available Sunday - Friday), and 2) written asynchronous consultations with a 24-

hour turn-around response time (available Monday - Friday). Synchronous appointments are video-based, with audio, chat, document-sharing, and live captioning capabilities, to meet a range of accessibility needs. View the synchronous and asynchronous schedules and book online, visit uri.mywconline.com.

Policies

Anti-Bias Statement

We respect the rights and dignity of each individual and group. We reject prejudice and intolerance, and we work to understand differences. We believe that equity and inclusion are critical components for campus community members to thrive. If you are a target or a witness of a bias incident, you are encouraged to submit a report to the URI Bias Response Team at www.uri.edu/brt. There you will also find people and resources to help.

Disability Services for Students Statement

Your access in this course is important. Please send me your Disability Services for Students (DSS) accommodation letter early in the semester so that we have adequate time to discuss and arrange your approved academic accommodations. If you have not yet established services through DSS, please contact them to engage in a confidential conversation about the process for requesting reasonable accommodations in the classroom. DSS can be reached by calling: 401-874-2098, visiting: web.uri.edu/disability, or emailing: dss@etal.uri.edu. They are available to meet with students enrolled in Kingston as well as Providence courses.

Academic Honesty

Students are expected to be honest in all academic work. A student's name on any written work, quiz or exam shall be regarded as assurance that the work is the result of the student's own independent thought and study. Work should be stated in the student's own words, properly attributed to its source. Students have an obligation to know how to quote, paraphrase, summarize, cite and reference the work of others with integrity. The following are examples of academic dishonesty.

- Using material, directly or paraphrasing, from published sources (print or electronic) without appropriate citation
- Claiming disproportionate credit for work not done independently
- Unauthorized possession or access to exams
- Unauthorized communication during exams
- Unauthorized use of another's work or preparing work for another student
- Taking an exam for another student
- Altering or attempting to alter grades
- The use of notes or electronic devices to gain an unauthorized advantage during exams
- Fabricating or falsifying facts, data or references
- Facilitating or aiding another's academic dishonesty
- Submitting the same paper for more than one course without prior approval from the instructors

URI COVID-19 Statement

The University is committed to delivering its educational mission while protecting the health and safety of our students. At this uncertain time, those concerns include minimizing the potential spread of COVID-19 within our community. While the university has worked this summer to create a healthy learning environment for all, it is up to all of us to ensure our campus stays that way.

As members of the URI community, students are required to comply with standards of conduct and take precautions to keep themselves and others safe. Students are required to comply with Rhode Island state laws, including the Rhode Island Executive Orders related to health and safety, ordinances, regulations, and guidance adopted by the University as it relates to public health crises, such as COVID-19.

An addendum on policies and guidelines concerning your obligations during this crisis has recently been integrated into the Student Handbook. These obligations include:

- Wearing of face masks by all community members when on a URI campus in the presence of others
- Maintaining physical distancing of at least six feet at all times
- Following state rules on the number of individuals allowed in a group gathering

- Completing a daily health self-assessment also available through the Rhody Connect app before coming to campus
- Submitting to COVID-19 testing as the University monitors the health of our community
- Following the University's quarantine and isolation requirements

If you answer yes to any of the questions on the daily health assessment, do not go to campus. YOU MUST STAY HOME/IN YOUR ROOM and notify URI Health Services via phone at 401-874-2246 immediately.

If you are already on campus and start to feel ill, you need to remove yourself from the public and notify URI Health Services via phone immediately at 401-874-2246 and go home/back to your room and self-isolate while you await direction from Health Services.

If you are unable to attend class, please notify me at brownsarahm@uri.edu or through the medium we have established for the class. We will work together to ensure that course instruction and work is completed for the semester.

Class Notes

Class notes will get posted here day by day

- [2020-09-11](#): Jupyter Notebook tour, conditionals, functions
- [2020-09-14](#): Iterables, Pandas
- [2020-09-16](#): Pandas loading and exploring
- [2020-09-18](#): Pandas, Functions as Object, Dictionaries
- [2020-09-21](#): Exploratory Data Analysis, Split, apply, Combine
- [2020-09-23](#): Visualization for EDA
- [2020-09-25](#): Viz & Cleaning
- [2020-09-28](#): Preparing data for analysis
- [2020-09-30](#): Reading complex data and Ray Summit
- [2020-10-02](#): Cleaning Data
- [2020-10-05](#): Merging DataFrames
- [2020-10-07](#): Merging & Databases
- [2020-10-12](#): Intro to ML
- [2020-10-14](#): Naive Bayes Classifier
- [2020-10-16](#): Classification Metrics
- [2020-10-19](#): Mid-Semester Feedback & Naive Bayes Decision rules
- [2020-10-21](#): Decision Trees
- [2020-10-23](#): Decision Trees & Cross validation
- [2020-10-26](#): Regression
- [2020-10-28](#): More Regression & More Evaluation
- [2020-10-30](#): Interpreting Regression Metrics
- [2020-11-02](#): Clustering
- [2020-11-04](#): Evaluating Clustering
- [2020-11-04](#): More Clustering Models
- [2020-11-06](#): Optimizing Clustering Models

Class 2: intro to notebooks and python

Agenda:

1. review Data Science
2. [jupyter notebook](#)
3. [python](#): conditionals and functions

Jupyter Notebooks

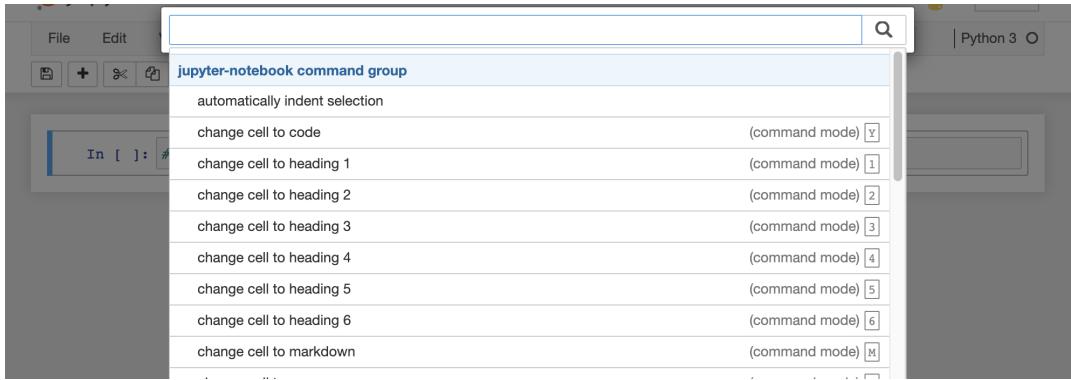
To launch a Jupyter notebook, in your Anaconda prompt on Windows or terminal on Linux or Mac:

```
cd dir/you/want/to/work/in
jupyter notebook
```

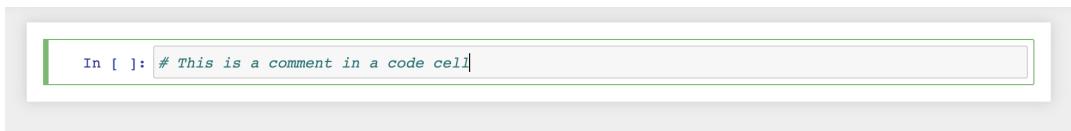
A Jupyter notebook has two modes. When you first open, it is in command mode. The border is blue in command mode.

```
In [ ]: # This is a comment in a code cell
```

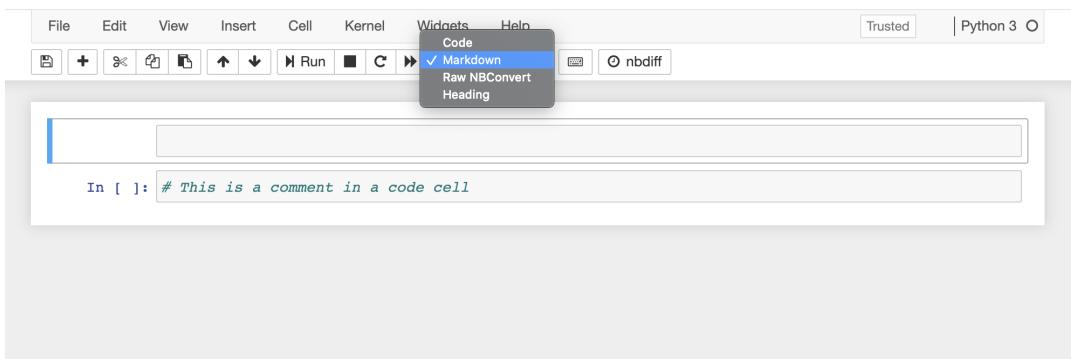
When you press a key in command mode it works like a shortcut. For example **p** shows the command search menu.



If you press **enter** (or **return**) or click on the cell it changes to edit mode. The border is green in edit mode



Type code or markdown into boxes called cells. There are two type of cells that we will used: code and markdown. You can change that in command mode with **y** for code and **m** for markdown or on the cell type menu at the top of the notebook.



You can treat markdown cells like plain text, or use special formatting. Here is a [markdown cheatsheet](#).

Code cells can run like a calculator. If there is a value returned by the last line of a cell, it will be displayed.

```
4+5
```

```
9
```

For example, when we assign, python "returns" **None** so there is no output from this cell

```
a = 9
```

But this one does display the value of the cell

```
a
```

```
9
```

```
b = 4
```

```
b
```

a

9

Getting Help in Python and Jupyter

The standard way to get help in the help function

```
help(print)
```

```
Help on built-in function print in module builtins:

print(...)
    print(value, ..., sep=' ', end='\n', file=sys.stdout, flush=False)

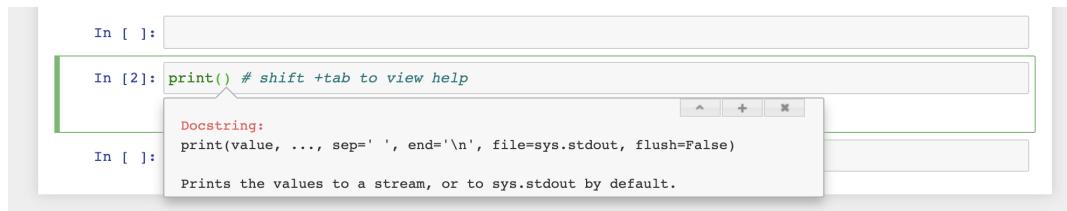
    Prints the values to a stream, or to sys.stdout by default.
    Optional keyword arguments:
    file: a file-like object (stream); defaults to the current sys.stdout.
    sep:   string inserted between values, default a space.
    end:   string appended after the last value, default a newline.
    flush: whether to forcibly flush the stream.
```

There are two special ways to get help in Jupyter, one dynamically while you're working and one that stays displayed for a while

Python comments are indicated by the `#` symbol.

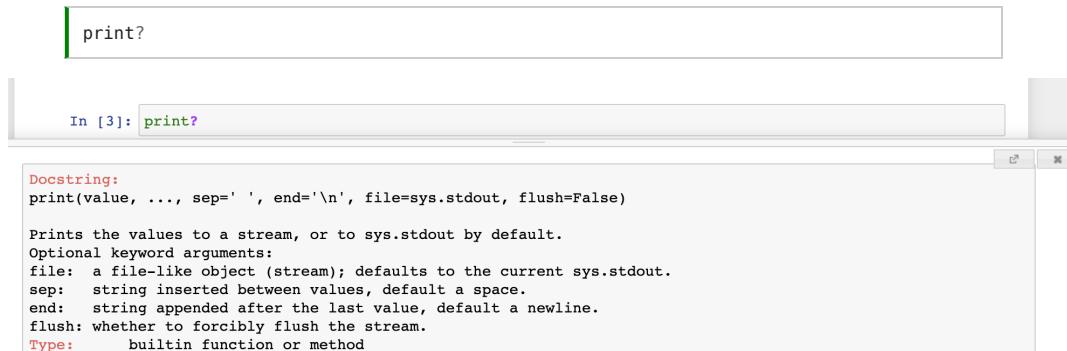
```
print() # shift +tab to view help
```

That will look like this:



Press tab twice for the longer version.

A Question mark puts it in a popup window that stays until you close it



This means you can then use the displayed help to remember how to call the function

```
print(a,b, 'hello',sep='-')
```

9-4-hello

Note

Here in class we changed the value of `a` above and noted that the new value is shown here, but not in the previous cell that had output the value of `a`

Note that these are green in the jupyter notebook because they're python reserved words.

```
a  
b
```

4

If Statements

⚠ Warning

if the concept of an `if` in programming is new to you, you should talk to Dr. Brown. Basic programming is a prerequisite to this course, we're reviewing basic ideas but only at a level of detail to serve as a reminder.

The syntax

```
if a >b:  
    print('greater')
```

greater

```
if b > a:  
    print('greater')
```

💡 Tip

this is updated to include things that were skipped in class and discussed after the breakouts

You can check the contents of a string with the `in` keyword

```
name = 'sarah'  
if 'a' in name:  
    print(name, 'has an a')
```

sarah has an a

💡 Tip

`in` works for all lists, we'll learn more about that next week

if we copy and change the name we get no output

```
name = 'Beibhinn'  
if 'a' in name:  
    print(name, 'has an a')
```

Functions

💡 Tip

this is also updated to include things that were skipped in class and discussed after the breakouts

How to write functions in python:

- the `def` keyword starts a function definition
- then the function name
- then the parameters it accepts in `()`
- end that line with a `:`
- the body of the function is spaced over one tab, but Jupyter will do it automatically for you. if it doesn't you might have forgotten the `:`

```
def greeting(name):
    """
    a function that greets the person name by printing

    parameters
    -----
    name: string
        a name to be greeeted

    Returns
    -----
    nothing
    """
    print('hello', name)
```

```
greeting('sarah')
```

```
hello sarah
```

A better version of that function might be:

```
def greeting(name):
    """
    a function that greets the person name by printing

    parameters
    -----
    name: string
        a name to be greeeted

    Returns
    -----
    nothing
    """
    return 'hello ' + name
```

⚠️ Warning

this is actually not a great function, because the printing is only a *side effect*. It's better to return the output of the function

💡 Tip

you can append strings with `+`

💡 Try it Yourself!

Write a function that checks if a string has a space in it and returns “please rename” if there is a space.

Remember a docstring. Call your function a couple of times to confirm it works.

Unhide the cell below to see the answer.

This is what some calls of the function look like

```
check_string("my data.csv")
```

```
'please rename'
```

If there's no string we see no output

```
check_string("my_data.csv")
```

What does python actually return?

```
NoneType
```

Further Reading

- [How Ipython Works](#)
- [Ipython Overview](#)
- [Jupyter Notebooks Technical Overview](#)

- [Python If Statements](#)
- [Python Functions](#)

Class 3: Welcome to Week 2

This week we will:

- clarify how this grading really works
- learn about accessing data
- use accessing data as motivation to review more python

Grading and Assignment 1

- Solution function posted.
- note: not a sum
- read the rubric
- Brightspace will show grades as they're earned
- In class, respond on prismia
- Portfolio
 - will start posting prompts The docstring functions like a property of the function object. so it has to be inside.

Iterables

Python has a general data type for objects that are designed to facilitate repetition of some sort, they're called **iterables**

We've already seen one. Strings are **Iterables**

```
name = 'sarah'
```

which means we can index them

```
name[3]
```

```
'a'
```

Tip

remember python indexes from 0

Indexing with a negative number counts from the end

```
name[-1]
```

```
'h'
```

Loops in python have similar syntax to the **if** and functions we saw last week:

```
for char in name:
    print(char*3)
```

```
sss
aaa
rrr
aaa
hhh
```

some notes:

- **char** is called the loop variable
- **name** is called the collection- this can be any iterable type object in python
- **print(char*3)** is called the loop body
- python lets us use mathematical operations on strings

Lists and List Comprehensions

We make a list with square brackets

```
names = ['sarah', 'Jose', 'Cam', 'Bri']
```

we can also build lists by folding a loop into the list construction

```
['hello' + n for n in names]
```

```
['hellosarah', 'helloJose', 'helloCam', 'helloBri']
```

this is called a list comprehension

```
greetings = ['hello ' + n for n in names]
```

```
greetings[0]
```

```
'hello sarah'
```

Dictionaries

Dictionaries are a useful datatype in python. It is denoted by {} and contains key: value pairs separated by commas.

```
gh_names = {'brownsarahm': 'Sarah Brown',
            'briannakathrynm1' : 'Brianna MacDonald',
            'jdion62': 'Jacob Dion'}
gh_names
```

```
{'brownsarahm': 'Sarah Brown',
 'briannakathrynm1': 'Brianna MacDonald',
 'jdion62': 'Jacob Dion'}
```

You can think of it like a list of the values with a named index.

```
gh_names['jdion62']
```

```
'Jacob Dion'
```

we can iterate over both the key and the value by using the `items` method on a dictionary. That makes another iterable object that can be used as a loop collection. It functions as a set of pairs now, so we get two loop variables:

```
for key, value in gh_names.items():
    print(value, "'s username is ", key)
```

```
Sarah Brown 's username is brownsarahm
Brianna MacDonald 's username is briannakathrynm1
Jacob Dion 's username is jdion62
```

If we iterate over the dictionary without that method, we get the keys.

```
for val in gh_names:
    print(val)
```

```
brownsarahm
briannakathrynm1
jdion62
```

Libraries

To use libraries in python we import them

We will use [pandas](#) a lot in this class. It's the Python Data Analysis Library.

```
import pandas
```

Once we import we can use the functions, datatypes, and values a library provides by using a `.` after the name. In a notebook, pressing tab will show you the options.

```
pandas.read_csv()
```

```
-----  
TypeError                                 Traceback (most recent call last)  
<ipython-input-14-374a1a6f9f7e> in <module>  
----> 1 pandas.read_csv()  
  
TypeError: read_csv() missing 1 required positional argument: 'filepath_or_buffer'
```

We can also use an alias to give a library a nickname to make it easier to use. `pd` is the standard alias for `pandas`

```
import pandas as pd
```

We can read in from a local path or a url. Let's read in the course map page of our course website.

```
pd.read_html('https://rhodyprog4ds.github.io/BrownFall20/syllabus/course_map.html')
```

💡 Tip

note that `import` is a keyword and that in a Jupyter notebook, we can import anywhere and then the library can be used in any cell that is run after the import cell is run. It's good practice to put them at the top and make your notebook runnable in sequence, but Jupyter won't force you to.

💡 Tip

For example if you don't remember what kind of read functions there are in pandas, type `pandas.read` and then press tab to see options.

```

[ Unnamed: 0_level_0 topics \
  week
0      1      Unnamed: 1_level_1
1      2      [admin, python review]
2      3      Loading data, Python review
3      4      Exploratory Data Analysis
4      5      Data Cleaning
5      6      Databases, Merging DataFrames
6      7      Modeling, Naive Bayes, classification performa...
7      8      decision trees, cross validation
8      9      Regression
9     10      Clustering
10    11      SVM, parameter tuning
11    12      KNN, Model comparison
12    13      Text Analysis
13    14      Topic Modeling
14      Deep Learning

          skills
Unnamed: 2_level_1
0      process
1      [access, prepare, summarize]
2      [summarize, visualize]
3      [prepare, summarize, visualize]
4      [access, construct, summarize]
5      [classification, evaluate]
6      [classification, evaluate]
7      [regression, evaluate]
8      [clustering, evaluate]
9      [optimize, tools]
10    [compare, tools]
11    [unstructured]
12    [unstructured, tools]
13    [tools, compare] , skill \
  Unnamed: 0_level_0
  keyword      Unnamed: 1_level_1
0      python      pythonic code writing
1      process      describe data science as a process
2      access       access data in multiple formats
3      construct    construct datasets from multiple sources
4      summarize    Summarize and describe data
5      visualize    Visualize data
6      prepare      prepare data for analysis
7      classification      Apply classification
8      regression   Apply Regression
9      clustering   Clustering
10    evaluate     Evaluate model performance
11    optimize     Optimize model parameters
12    compare      compare models
13    unstructured model unstructured data
14    tools        use industry standard data science tools and w...
          Level 1 \
  Unnamed: 2_level_1
0  python code that mostly runs, occasional pep8 ...
1      Identify basic components of data science
2  load data from at least one format; identify t...
3  identify what should happen to merge datasets ...
4  Describe the shape and structure of a dataset ...
5  identify plot types, generate basic plots from...
6  identify if data is or is not ready for analys...
7  identify and describe what classification is, ...
8  identify what data that can be used for regres...
9      describe what clustering is
10 Explain basic performance metrics for differen...
11 Identify when model parameters need to be opti...
12      Qualitatively compare model classes
13 Identify options for representing text data an...
14 Solve well strucutred problems with a single t...

          Level 2 \
  Unnamed: 3_level_1
0  python code that reliably runs, frequent pep8 ...
1  Describe and define each stage of the data sci...
2  Load data for processing from the most common ...
3      apply basic merges
4  compute summary statandard statistics of a whol...
5  generate multiple plot types with complete lab...
6  apply data reshaping, cleaning, and filtering ...
7  fit preselected classification model to a dataset
8      can fit linear regression models
9      apply basic clustering
10 Apply basic model evaluation metrics to a held...
11 Manually optimize basic model parameters such ...
12 Compare model classes in specific terms and fi...
13 Apply at least one representation to transform...
14 Solve semi-strucutred, completely specified pr...

```

	Level 3		
	Unnamed: 4_level_1		
0	reliable, efficient, pythonic code that consis...		
1	Compare different ways that data science can f...		
2	access data from both common and uncommon form...		
3	merge data that is not automatically aligned		
4	Compute and interpret various summary statisti...		
5	generate complex plots with pandas and plottin...		
6	apply data reshaping, cleaning, and filtering ...		
7	fit and apply classification models and select...		
8	can fit and explain nonlinear regression		
9	apply multiple clustering techniques, and inte...		
10	Evaluate a model with multiple metrics and cro...		
11	Select optimal parameters based of mutiple qua...		
12	Evaluate tradeoffs between different model com...		
13	apply multilple representations and compare and...		
14	Scope, choose an appropriate tool pipeline and... ,		
	Unnamed: 0_level_0	A1	A2 \
	keyword	Unnamed: 1_level_1	Unnamed: 2_level_1
0	python	1	1
1	process	1	1
2	access	0	1
3	construct	0	0
4	summarize	0	0
5	visualize	0	0
6	prepare	0	0
7	classification	0	0
8	regression	0	0
9	clustering	0	0
10	evaluate	0	0
11	optimize	0	0
12	compare	0	0
13	unstructured	0	0
14	tools	0	0
	A3	A4	A5 \
	Unnamed: 3_level_1	Unnamed: 4_level_1	Unnamed: 5_level_1
0	1	1	0
1	0	0	0
2	1	1	0
3	0	0	1
4	1	1	1
5	1	1	0
6	0	1	1
7	0	0	0
8	0	0	0
9	0	0	0
10	0	0	0
11	0	0	0
12	0	0	0
13	0	0	0
14	0	0	0
	A6	A7	A8 \
	Unnamed: 6_level_1	Unnamed: 7_level_1	Unnamed: 8_level_1
0	0	0	0
1	0	0	0
2	0	0	0
3	1	0	0
4	1	1	1
5	1	1	1
6	0	0	0
7	1	1	0
8	0	0	1
9	0	0	0
10	0	0	0
11	0	0	0
12	0	0	0
13	0	0	0
14	0	0	0
	A9	A10	A11 \
	Unnamed: 9_level_1	Unnamed: 10_level_1	Unnamed: 11_level_1
0	0	0	0
1	0	0	0
2	0	0	0
3	0	0	0
4	1	1	1
5	1	1	1
6	0	0	0
7	0	1	0
8	0	0	1
9	1	0	1
10	0	1	1
11	0	1	1
12	0	0	1
13	0	0	0

14	0	1	1	
	A12	A13	# Assignments	
0	Unnamed: 12_level_1	Unnamed: 13_level_1	Unnamed: 14_level_1	
1	0	0	4	
2	0	0	2	
3	0	0	3	
4	1	1	2	
5	1	1	11	
6	0	0	10	
7	0	0	2	
8	0	0	3	
9	0	0	2	
10	0	0	2	
11	0	0	2	
12	0	1	2	
13	1	1	2	
14	1	1	4	,
	Unnamed: 0_level_0		Level 3 \	
	keyword		Unnamed: 1_level_1	
0	python	reliable, efficient, pythonic code that consis...		
1	process	Compare different ways that data science can f...		
2	access	access data from both common and uncommon form...		
3	construct	merge data that is not automatically aligned		
4	summarize	Compute and interpret various summary statisti...		
5	visualize	generate complex plots with pandas and plottin...		
6	prepare	apply data reshaping, cleaning, and filtering ...		
7	classification	fit and apply classification models and select...		
8	regression	can fit and explain nonlinear regression		
9	clustering	apply multiple clustering techniques, and inte...		
10	evaluate	Evaluate a model with multiple metrics and cro...		
11	optimize	Select optimal parameters based of mutiple qua...		
12	compare	Evaluate tradeoffs between different model com...		
13	unstructured	apply mulitple representations and compare and...		
14	tools	Scope, choose an appropriate tool pipeline and...		
	P1	P2	P3	P4
	Unnamed: 2_level_1	Unnamed: 3_level_1	Unnamed: 4_level_1	Unnamed: 5_level_1
0	1	1	0	0
1	0	1	1	0
2	1	1	0	0
3	1	1	0	0
4	1	1	0	0
5	1	1	0	0
6	1	1	0	0
7	0	1	1	0
8	0	1	1	0
9	0	1	1	0
10	0	1	1	0
11	0	0	1	1
12	0	0	1	1
13	0	0	1	1
14	0	0	1	1]

This makes a `list` of `pandas.DataFrame` objects. We can check that with the following

⚠️ Warning

This cell was added after class, but the explanation was given in class

```
type(pd.read_html('https://rhodyprog4ds.github.io/BrownFall20/syllabus/course_map.html'))
```

list

To work with it though, we should save to a variable, then we can index into that list.

```
df_list =
pd.read_html('https://rhodyprog4ds.github.io/BrownFall20/syllabus/course_map.html')
df_list[0]
```

Unnamed: 0_level_0	topics	skills
	week	
0	1	[admin, python review]
1	2	Loading data, Python review
2	3	Exploratory Data Analysis
3	4	Data Cleaning
4	5	Databases, Merging DataFrames
5	6	Modeling, Naive Bayes, classification performa...
6	7	decision trees, cross validation
7	8	Regression
8	9	Clustering
9	10	SVM, parameter tuning
10	11	KNN, Model comparison
11	12	Text Analysis
12	13	Topic Modeling
13	14	Deep Learning

When you display `DataFrames` in jupyter, they get nice formatting.

Review & Further Reading

- [strings](#) are [iterable](#), more specifically [sequences](#)
- [for loops](#), [looping techniques](#) and [list comprehensions](#)
- [dictionaries](#)
- imported [pandas](#) and [read data from a website](#)

If you've made it this far, [let me know](#) how you found these notes.

Class 4: Pandas

Today we will:

Remember, Programming is a Practice

- if you're curious about something try it
- you don't need me to give you answers about how code works, the interpreter will tell you
- if you don't remember details, remember you can get help from Jupyter

with a `?` after the function name without `()`

```
print?
```

or using the `tab` key inside the `()` for a function

```
print()
```

or from the core python, with the `help` function

```
help(print)
```

```
Help on built-in function print in module builtins:
```

```
print(...)  
    print(value, ..., sep=' ', end='\n', file=sys.stdout, flush=False)  
  
    Prints the values to a stream, or to sys.stdout by default.  
    Optional keyword arguments:  
        file: a file-like object (stream); defaults to the current sys.stdout.  
        sep:   string inserted between values, default a space.  
        end:   string appended after the last value, default a newline.  
        flush: whether to forcibly flush the stream.
```

Data in Pandas

We can import `pandas` again as before

```
import pandas as pd
```

and we can read in data.

```
pd.read_csv('https://raw.githubusercontent.com/brownsarahm/python-socialsci-  
files/master/data/SAFI_clean.csv')
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms
0	1	God	2016-11- 17T00:00:00Z	3	4	muddaub	1
1	1	God	2016-11- 17T00:00:00Z	7	9	muddaub	1
2	3	God	2016-11- 17T00:00:00Z	10	15	burntbricks	1
3	4	God	2016-11- 17T00:00:00Z	7	6	burntbricks	1

to be able to use this, we need to save it to a variable.

```
safi_df = pd.read_csv('https://raw.githubusercontent.com/brownsarahm/python-socialsci-  
files/master/data/SAFI_clean.csv')
```

This is an excerpt from the [SAFI dataset](#).

Another important thing to do is to check datatypes, this is how we know what things we can do with a variable.

```
type(safi_df)
```

```
pandas.core.frame.DataFrame
```

An important thing to check is the size of the dataset.

```
safi_df.shape
```

```
(131, 14)
```



we can also see the size when we print the whole dataset as in the first time we read the data in.

Recall that you can also tab complete

```
safi_df.shape
```

```
(131, 14)
```

To see the first 5 rows of the dataset, use the `head()` function

```
safi_df.head()
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	memr
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	
2	3	God	2016-11-17T00:00:00Z	10	15	burntbricks	1	
3	4	God	2016-11-17T00:00:00Z	7	6	burntbricks	1	
4	5	God	2016-11-17T00:00:00Z	7	40	burntbricks	1	

We can call this function with a value to change how many rows are returned

```
safi_df.head(3)
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	memr
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	
2	3	God	2016-11-17T00:00:00Z	10	15	burntbricks	1	

To know how this works, we can view the documentation for the function

```
help(safi_df.head)
```

⚠️ Warning

this was changed from using `?` for help in class so that the help is displayed in the rendered website, but the pop up was fine in real time

```

Help on method head in module pandas.core.generic:

head(n: int = 5) -> ~FrameOrSeries method of pandas.core.frame.DataFrame instance
Return the first `n` rows.

This function returns the first `n` rows for the object based
on position. It is useful for quickly testing if your object
has the right type of data in it.

For negative values of `n`, this function returns all rows except
the last `n` rows, equivalent to ``df[:-n]``.

Parameters
-----
n : int, default 5
    Number of rows to select.

Returns
-----
same type as caller
    The first `n` rows of the caller object.

See Also
-----
DataFrame.tail: Returns the last `n` rows.

Examples
-----
>>> df = pd.DataFrame({'animal': ['alligator', 'bee', 'falcon', 'lion',
...                                'monkey', 'parrot', 'shark', 'whale', 'zebra']})
>>> df
      animal
0  alligator
1        bee
2     falcon
3       lion
4     monkey
5    parrot
6     shark
7     whale
8     zebra

Viewing the first 5 lines

>>> df.head()
      animal
0  alligator
1        bee
2     falcon
3       lion
4     monkey

Viewing the first `n` lines (three in this case)

>>> df.head(3)
      animal
0  alligator
1        bee
2     falcon

For negative values of `n`

>>> df.head(-3)
      animal
0  alligator
1        bee
2     falcon
3       lion
4     monkey
5    parrot

```

Since it says `n =5` we know that the default value of the parameter `n` is 5. When a function has a default value, we can call the function without a value.

To view the last few lines, we use `tail`

```
safi_df.tail()
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms
126	126	Ruaca	2017-05-18T00:00:00Z	3	7	burntbricks	1
127	193	Ruaca	2017-06-04T00:00:00Z	7	10	cement	3
128	194	Ruaca	2017-06-04T00:00:00Z	4	5	muddaub	1
129	199	Chirodzo	2017-06-04T00:00:00Z	7	17	burntbricks	2
130	200	Chirodzo	2017-06-04T00:00:00Z	8	20	burntbricks	2

We can also get an [Index](#) for the columns of the DataFrame.

```
safi_df.columns
```

[Index](#) is another data type that is defined by [pandas](#).

```
Index(['key_ID', 'village', 'interview_date', 'no_membrs', 'years_liv',
       'respondent_wall_type', 'rooms', 'memb_assoc', 'affect_conflicts',
       'liv_count', 'items_owned', 'no_meals', 'months_lack_food',
       'instanceID'],
      dtype='object')
```

an [Index](#) variable is iterable so we can index into it

Try it Yourself

How would you view the name of the 3rd column?

First the correct answer:

```
'interview_date'
```

Now some misconceptions:

```
safi_df['interview_date']
```

```
0    2016-11-17T00:00:00Z
1    2016-11-17T00:00:00Z
2    2016-11-17T00:00:00Z
3    2016-11-17T00:00:00Z
4    2016-11-17T00:00:00Z
      ...
126   2017-05-18T00:00:00Z
127   2017-06-04T00:00:00Z
128   2017-06-04T00:00:00Z
129   2017-06-04T00:00:00Z
130   2017-06-04T00:00:00Z
Name: interview_date, Length: 131, dtype: object
```

Indexing with the column name) will return the *values* in the column

```
safi_df.columns[2]
```

```
TypeError
<ipython-input-17-bd02c7e8a4a6> in <module>
----> 1 safi_df.columns(2)

TypeError: 'Index' object is not callable
```

Using `()` returns an error, because `columns` is an *attribute* which is referenced as is with no `()`. We get a type error because functions in python are objects of type `callable` and properties are values not functions.

```
pd.DataFrame.columns[2]
```

```
TypeError
<ipython-input-18-40e277f3074e> in <module>
----> 1 pd.DataFrame.columns[2]

TypeError: 'pandas._libs.properties.AxisProperty' object is not subscriptable
```

This doesn't work because `columns` is an attribute of an object of type `pandas.DataFrame` and `pd.DataFrame.columns` is not an object.

We can see what the type of `pd.DataFrame` is with the `type` function.

```
type(pd.DataFrame)
```

```
type
```

Knowing about types is helpful for the individual columns of a dataset as well.

```
safi_df.dtypes
```

```
key_ID          int64
village         object
interview_date  object
no_membrs       int64
years_liv       int64
respondent_wall_type  object
rooms           int64
memb_assoc      object
affect_conflicts object
liv_count        int64
items_owned     object
no_meals         int64
months_lack_food object
instanceID      object
dtype: object
```

Note that it uses `int64` and `object` as the types.

```
safi_df.head(2)
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	merr
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	

We might want to look at what villages were included in the data.

```
pd.unique(safi_df['village'])
```

```
array(['God', 'Chirodzo', 'Ruaca'], dtype=object)
```

We can also get count of the number of each value

```
safi_df['village'].value_counts()
```

```
Ruaca      49
God        43
Chirodzo   39
Name: village, dtype: int64
```

➊ Try it Yourself!

how many surveyed farms have all type `mauddaub`?

Review and Further reading

- [reading data with pandas](#)
- [Python built in functions](#) and in particular the [type function](#)
- [Pandas DataFrames](#)
- [value_counts](#)

If you've made it this far, [let me know](#) how you found these notes.

Class 5: Accessing Data, continued

Today's agenda:

- warm up/ review
- announcements
- working with dataframes
- the power of functions as objects
- (maybe) exploratory data analysis

➊ Try it out ->

Read the tables off of the syllabus course map page with `read_html` and make a list of the shapes of all of the tables on the page. Save the output to a variable and paste the `value` of that variable as your answer to the question.

```
import pandas as pd
[df.shape for df in
pd.read_html('https://rhodyprog4ds.github.io/BrownFall20/syllabus/course_map.html')]
```

```
[(14, 3), (15, 5), (15, 15), (15, 6)]
```

Announcements

- annotated notes are up
- beginning portfolio prompts and instructions are up
- Assignment due Sunday,
- office hours will remain Fridays
- TA office hours posted.

More Pandas

We'll go back to the SAFI dataset from Wednesday.

```
safi_df = pd.read_csv('https://raw.githubusercontent.com/brownsarahm/python-socialsci-files/master/data/SAFI_clean.csv')
```

We downloaded the data into memory, but we can also write it to disk.

```
safi_df.to_csv('safi_clean.csv')
```

It will go to the same folder as the notebook, but we can also use a relative path. If we make a `data` folder in the folder where we've saved the notebook, we can write the file there instead.

```
safi_df.to_csv('data/safi_clean.csv')
```

Now we can read it in using the same path

```
safi_df2 = pd.read_csv('data/safi_clean.csv')
```

Note that now it has an extra column

```
safi_df2.head(2)
```

	Unnamed: 0	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	merr
0	0	1	God	2016-11-17T00:00:00Z	3	4	muddaub		
1	1	1	God	2016-11-17T00:00:00Z	7	9	muddaub		

```
safi_df.head(2)
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	merr
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	

We can prevent this by writing it out with the `index` parameter set to `False`

```
safi_df.to_csv('data/safi_clean.csv', index=False)
```

Now when we read it in, there's no extra column.

```
safi_df3 = pd.read_csv('data/safi_clean.csv')
safi_df3.head(3)
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	merr
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	
2	3	God	2016-11-17T00:00:00Z	10	15	burntbricks	1	

Tip

In class we used the Jupyter GUI to create a new folder. You could also use your computer's default file management tool (Windows Explorer, Mac Finder, etc). Here, since the notebooks have to run completely automatically for this website, we use a ipython `bash magic` cell to make the folder. Jupyter notebooks use an ipython kernel.

Tip

`False` must be with a capital letter to be a boolean variable in python, as with `True`. You'll know if you did it right in your jupyter notebook, if the word terms bold and green.

Recall, we indexed a column with the name in square brackets

```
safi_df['village']
```

```
0      God  
1      God  
2      God  
3      God  
4      God  
     ...  
126    Ruaca  
127    Ruaca  
128    Ruaca  
129  Chirodzo  
130  Chirodzo  
Name: village, Length: 131, dtype: object
```

To index rows, we can use `loc`

```
safi_df.loc[3]
```

```
key_ID          4  
village        God  
interview_date 2016-11-17T00:00:00Z  
no_membrs       7  
years_liv        6  
respondent_wall_type burntbricks  
rooms            1  
memb_assoc      NaN  
affect_conflicts NaN  
liv_count         2  
items_owned     bicycle;radio;cow_plough;solar_panel;mobile_phone  
no_meals           2  
months_lack_food Sept;Oct;Nov;Dec  
instanceID      uuid:148d1105-778a-4755-aa71-281eadd4a973  
Name: 3, dtype: object
```

To select a range, use :

```
safi_df.loc[3:5]
```

		key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	memr
3	4	God	2016-11-17T00:00:00Z	7	6		burntbricks	1	
4	5	God	2016-11-17T00:00:00Z	7	40		burntbricks	1	
5	6	God	2016-11-17T00:00:00Z	3	3		muddaub	1	

You only have to have a number on one side of the colon, it will go from the beginnig up to that number like this:

```
safi_df.loc[:4]
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	mem
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	
2	3	God	2016-11-17T00:00:00Z	10	15	burntbricks	1	
3	4	God	2016-11-17T00:00:00Z	7	6	burntbricks	1	
4	5	God	2016-11-17T00:00:00Z	7	40	burntbricks	1	

With two `:` we can also set an increment

```
safi_df.loc[::5]
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1
5	6	God	2016-11-17T00:00:00Z	3	3	muddaub	1
10	11	God	2016-11-21T00:00:00Z	6	20	sunbricks	1
15	16	God	2016-11-24T00:00:00Z	6	47	muddaub	1
20	21	God	2016-11-21T00:00:00Z	8	20	burntbricks	1
25	26	Ruaca	2016-11-21T00:00:00Z	3	20	burntbricks	2
30	31	Ruaca	2016-11-21T00:00:00Z	3	2	muddaub	1
35	36	Chirodzo	2016-11-17T00:00:00Z	6	23	sunbricks	1
40	41	God	2016-11-17T00:00:00Z	7	22	muddaub	1
45	46	Chirodzo	2016-11-17T00:00:00Z	10	42	burntbricks	2
50	51	Chirodzo	2016-11-16T00:00:00Z	5	30	muddaub	1
55	56	Chirodzo	2016-11-16T00:00:00Z	12	23	burntbricks	2
60	61	Chirodzo	2016-11-16T00:00:00Z	10	14	muddaub	1
65	66	Chirodzo	2016-11-16T00:00:00Z	10	37	burntbricks	3
70	71	Ruaca	2016-11-18T00:00:00Z	6	14	burntbricks	1
75	155	God	2016-11-24T00:00:00Z	4	4	burntbricks	1

key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms
80	182	God 2016-11-25T00:00:00Z	7	21	muddaub	3
85	197	God 2016-11-28T00:00:00Z	5	19	burntbricks	2
90	73	Ruaca 2017-04-26T00:00:00Z	7	9	burntbricks	2
95	101	God 2017-04-27T00:00:00Z	3	4	muddaub	1
100	104	Ruaca 2017-04-28T00:00:00Z	14	52	sunbricks	1
105	113	Ruaca 2017-05-03T00:00:00Z	11	26	burntbricks	3
110	108	God 2017-05-11T00:00:00Z	15	22	burntbricks	2
115	150	Ruaca 2017-05-18T00:00:00Z	7	8	muddaub	1
120	167	Ruaca 2017-06-03T00:00:00Z	8	24	muddaub	1
125	192	Chirodzo 2017-06-03T00:00:00Z	9	20	burntbricks	1
130	200	Chirodzo 2017-06-04T00:00:00Z	8	20	burntbricks	2

These can be combined to index a subset at an increment.

We can index columns in two ways, as we did on Wednesday

```
safi_df['village'].head(2)
```

```
0    God
1    God
Name: village, dtype: object
```

Or using a .

```
safi_df.village.head(2)
```

```
0    God
1    God
Name: village, dtype: object
```

We can select multiple columns, using a [list](#) of column names. We can define the list inline.

```
safi_df[['village', 'no_membrs', 'years_liv']].head(2)
```

```
village no_membrs years_liv  
0 God 3 4  
1 God 7 9
```

or in a separate variable

```
columns_of_interest = ['village','no_membrs','years_liv']  
safi_df[columns_of_interest].head(2)
```

```
village no_membrs years_liv  
0 God 3 4  
1 God 7 9
```

Functions are objects

```
syllabus_df_list =  
pd.read_html('https://rhodyprog4ds.github.io/BrownFall20/syllabus/course_map.html')
```

And we can put them in a dictionary. `lambda` functions are special functions defined in a single line.

```
greetingl = lambda name: 'hello ' + name  
greetingl('sarah')
```

```
'hello sarah'
```

is the same as

```
def greetingf(name):  
    return 'hello ' + name  
greetingf('sarah')
```

```
'hello sarah'
```

So, we can define a function in a dictionary like this:

```
view_rows = {0: lambda df: print(df.head()),  
            1: lambda df: print(df.tail())}
```

The `len` function works on all iterables

```
for df in syllabus_df_list:  
    num_row = len(df)  
    view_rows[num_row%2](df)
```

💡 Tip

this is how to do the equivalent of a switch or case in other languages in python

```

    Unnamed: 0_level_0                      topics \
        week          Unnamed: 1_level_1
0           1      [admin, python review]
1           2      Loading data, Python review
2           3      Exploratory Data Analysis
3           4      Data Cleaning
4           5      Databases, Merging DataFrames

        skills
        Unnamed: 2_level_1
0           process
1   [access, prepare, summarize]
2   [summarize, visualize]
3 [prepare, summarize, visualize]
4 [access, construct, summarize]

    Unnamed: 0_level_0                     skill \
        keyword          Unnamed: 1_level_1
10      evaluate      Evaluate model performance
11      optimize       Optimize model parameters
12      compare        compare models
13      unstructured   model unstructured data
14      tools          use industry standard data science tools and w...

          Level 1 \
        Unnamed: 2_level_1
10 Explain basic performance metrics for differen...
11 Identify when model parameters need to be opti...
12 Qualitatively compare model classes
13 Identify options for representing text data an...
14 Solve well strucutred problems with a single t...

          Level 2 \
        Unnamed: 3_level_1
10 Apply basic model evaluation metrics to a held...
11 Manually optimize basic model parameters such ...
12 Compare model classes in specific terms and fi...
13 Apply at least one representation to transform...
14 Solve semi-strucutred, completely specified pr...

          Level 3
        Unnamed: 4_level_1
10 Evaluate a model with multiple metrics and cro...
11 Select optimal parameters based of mutiple qua...
12 Evaluate tradeoffs between different model com...
13 apply mulitple representations and compare and...
14 Scope, choose an appropriate tool pipeline and...

    Unnamed: 0_level_0      A1      A2 \
        keyword Unnamed: 1_level_1 Unnamed: 2_level_1
10      evaluate      0      0
11      optimize       0      0
12      compare        0      0
13      unstructured   0      0
14      tools          0      0

          A3      A4      A5 \
        Unnamed: 3_level_1 Unnamed: 4_level_1 Unnamed: 5_level_1
10      0      0      0
11      0      0      0
12      0      0      0
13      0      0      0
14      0      0      0

          A6      A7      A8 \
        Unnamed: 6_level_1 Unnamed: 7_level_1 Unnamed: 8_level_1
10      0      0      0
11      0      0      0
12      0      0      0
13      0      0      0
14      0      0      0

          A9      A10     A11 \
        Unnamed: 9_level_1 Unnamed: 10_level_1 Unnamed: 11_level_1
10      0      1      1
11      0      1      1
12      0      0      1
13      0      0      0
14      0      1      1

          A12     A13 # Assignments
        Unnamed: 12_level_1 Unnamed: 13_level_1 Unnamed: 14_level_1
10      0      0      2
11      0      0      2
12      0      1      2
13      1      1      2
14      1      1      4

    Unnamed: 0_level_0                     Level 3 \
        keyword          Unnamed: 1_level_1

```

```

10      evaluate Evaluate a model with multiple metrics and cro...
11      optimize Select optimal parameters based of mutiple qua...
12      compare Evaluate tradeoffs between different model com...
13      unstructured apply mulitple representations and compare and...
14      tools Scope, choose an appropriate tool pipeline and...

```

	P1	P2	P3	P4
	Unnamed: 2_level_1	Unnamed: 3_level_1	Unnamed: 4_level_1	Unnamed: 5_level_1
10	0	1	1	0
11	0	0	1	1
12	0	0	1	1
13	0	0	1	1
14	0	0	1	1

The beginning of Exploratory Data Analysis

Pandas will give us descriptive statistics

```
safi_df.describe()
```

	key_ID	no_membrs	years_liv	rooms	liv_count	no_meals
count	131.000000	131.000000	131.000000	131.000000	131.000000	131.000000
mean	85.473282	7.19084	23.053435	1.740458	2.366412	2.603053
std	63.151628	3.17227	16.913041	1.092547	1.082775	0.491143
min	1.000000	2.000000	1.000000	1.000000	1.000000	2.000000
25%	32.500000	5.000000	12.000000	1.000000	1.000000	2.000000
50%	66.000000	7.000000	20.000000	1.000000	2.000000	3.000000
75%	138.000000	9.000000	27.500000	2.000000	3.000000	3.000000
max	202.000000	19.000000	96.000000	8.000000	5.000000	3.000000

The statistics of the `key_ID` column don't make a lot of sense. We can avoid that by making it the index

```
safi_df.head()
```

	key_ID	village	interview_date	no_membrs	years_liv	respondent_wall_type	rooms	mem
0	1	God	2016-11-17T00:00:00Z	3	4	muddaub	1	
1	1	God	2016-11-17T00:00:00Z	7	9	muddaub	1	
2	3	God	2016-11-17T00:00:00Z	10	15	burntbricks	1	
3	4	God	2016-11-17T00:00:00Z	7	6	burntbricks	1	
4	5	God	2016-11-17T00:00:00Z	7	40	burntbricks	1	

the `inplace` parameter of a `pandas` functions applies the operation to the DataFrame in memory, but then the function returns nothing, but if we display after that, we see that now the `key_ID` column is now the index.

```
safi_df.set_index('key_ID', inplace=True)
safi_df.head(2)
```

```
village interview_date no_membrs years_liv respondent_wall_type rooms memb_e
```

key_ID

```
1 God 2016-11-17T00:00:00Z 3 4 muddaub 1
```

```
1 God 2016-11-17T00:00:00Z 7 9 muddaub 1
```

and if we describe again, we see it doesn't compute on that column

```
safi_df.describe()
```

	no_membrs	years_liv	rooms	liv_count	no_meals
count	131.000000	131.000000	131.000000	131.000000	131.000000
mean	7.19084	23.053435	1.740458	2.366412	2.603053
std	3.17227	16.913041	1.092547	1.082775	0.491143
min	2.00000	1.000000	1.000000	1.000000	2.000000
25%	5.00000	12.000000	1.000000	1.000000	2.000000
50%	7.00000	20.000000	1.000000	2.000000	3.000000
75%	9.00000	27.500000	2.000000	3.000000	3.000000
max	19.00000	96.000000	8.000000	5.000000	3.000000

We can also call any of those on one column or one statistic.

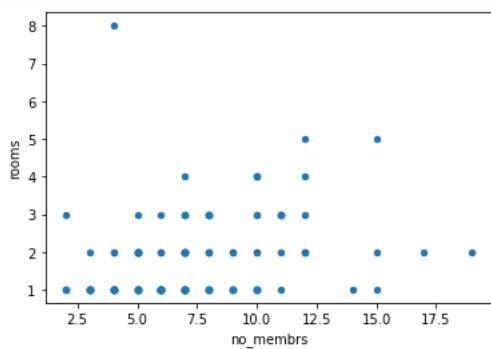
```
safi_df['rooms'].mean()
```

```
1.7404580152671756
```

Pandas also has some built in plotting functions.

```
safi_df.plot.scatter('no_membrs', 'rooms')
```

```
<AxesSubplot:xlabel='no_membrs', ylabel='rooms'>
```



After Class Questions

How can we clean data? what other topics will we cover in this class?



How does the syntax on the question from prismia today work?



Where I can go to find a list of all the syntax.



More Practice

These additional questions are for if you want more practice with things we've done this week, before class next week.

Which of the following is a dictionary?



What type is the shape of a pandas DataFrame?



What does indexing with -1 do?



Further Reading

If you've made it this far, [let me know](#) how you found these notes.

Class 6: Exploratory Data Analysis

Warmup

```
topics = ['what is data science', 'jupyter', 'conditional','functions', 'lists',
'dictionaries','pandas' ]
topics
```

```
['what is data science',
 'jupyter',
 'conditional',
 'functions',
 'lists',
 'dictionaries',
 'pandas']
```

What happens when we index with `-1`?

```
topics[-1]
```

```
'pandas'
```

We get the last value.

Recall last class we used `:` to index `DataFrames` with `.loc`, we can do that with lists too:

```
topics[-2:]
```

```
['dictionaries', 'pandas']
```

Announcements

- next assignment will go up today
- try practicing throughout the week
- Check Brightspace for TA office hours

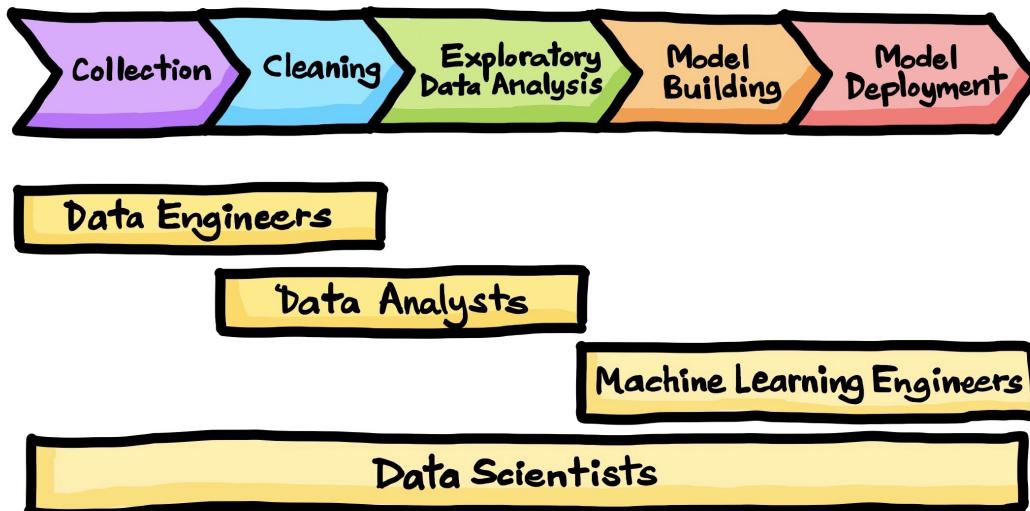
Why Exploratory Data Analysis (EDA) before cleaning?

Typically a Data process looks like one of these figures:

Data Science Process



THE DATA SCIENCE PROCESS



Data cleaning, which some of you asked about on Friday, would typically, *happen* before EDA, but it's hard to know what good data looks like, until you're used to manipulating it. Also, it's hard to check if data is good (and your cleaning worked) without some EDA. So, I'm choosing to *teach* EDA first. We'll do data cleaning next, with these tools from EDA available as options for checking and determining what cleaning to do.

EDA

```
# a version of the SAFI data with more columns
data_url = 'https://raw.githubusercontent.com/brownsarahm/python-socialsci-
files/master/data/SAFI_full_shortname.csv'
```

As usual, we import `pandas`

```
import pandas as pd
```

and pull in the data.

```
safi_df = pd.read_csv(data_url)
safi_df.head(2)
```

	key_id	interview_date	quest_no	start	end	province	district
0	1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica
1	2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica

2 rows × 65 columns

Recall on Friday, we can use `describe` to see several common descriptive statistics

```
safi_df.describe()
```

	key_id	quest_no	years_farm	note2	no_membrs	members_count	year
count	131.000000	131.000000	131.000000	0.0	131.000000	131.000000	131.000000
mean	66.000000	85.473282	15.832061	NaN	7.19084	7.19084	23.053
std	37.960506	63.151628	10.903883	NaN	3.17227	3.17227	16.913
min	1.000000	1.000000	1.000000	NaN	2.00000	2.00000	1.000
25%	33.500000	32.500000	8.000000	NaN	5.00000	5.00000	12.000
50%	66.000000	66.000000	15.000000	NaN	7.00000	7.00000	20.000
75%	98.500000	138.000000	20.500000	NaN	9.00000	9.00000	27.500
max	131.000000	202.000000	60.000000	NaN	19.00000	19.00000	96.000

8 rows × 25 columns

Then we remember that this includes `key_id` as a variable tht we don't actually want to treat as a variable, it's an index. We can change that with `set_index` and we can do it in memory (without assignment) using the `inplace` keyword.

```
safi_df.set_index('key_id', inplace=True)
safi_df.describe()
```

	quest_no	years_farm	note2	no_membrs	members_count	years_liv	respond
count	131.000000	131.000000	0.0	131.000000	131.000000	131.000000	131.000000
mean	85.473282	15.832061	NaN	7.19084	7.19084	23.053435	
std	63.151628	10.903883	NaN	3.17227	3.17227	16.913041	
min	1.000000	1.000000	NaN	2.00000	2.00000	1.000000	
25%	32.500000	8.000000	NaN	5.00000	5.00000	12.000000	
50%	66.000000	15.000000	NaN	7.00000	7.00000	20.000000	
75%	138.000000	20.500000	NaN	9.00000	9.00000	27.500000	
max	202.000000	60.000000	NaN	19.00000	19.00000	96.000000	

8 rows × 24 columns

Try it yourself!

You can use settings on `pd.read_csv` to set the index when the data is read in instead of doing this after the fact

Note

In class, we used `%load http://drsmmb.co/310` to pull in the url for the data. `Load` is an ipython (ther kernel tha Jupyter uses) magic function. It allows us to read in data from another place. I've set that short url to link to the download url for [this hackmd](#) as markdown(plain text). That way I can paste things and you can pull them directly into your notebook. We'll use it like a shared copy-paste.

We can also use the descriptive statistics on a single variable.

```
safi_df['years_farm'].describe()
```

```
count    131.000000
mean     15.832061
std      10.903883
min      1.000000
25%      8.000000
50%      15.000000
75%      20.500000
max      60.000000
Name: years_farm, dtype: float64
```

Note however that this is not well formatted, that's because it's a **Series** instead of a DataFrame

```
type(safi_df['years_farm'].describe())
```

```
pandas.core.series.Series
```

We can use `reset_index()` to make it back into a dataframe

```
safi_df['years_farm'].describe().reset_index()
```

```
index  years_farm
0   count  131.000000
1   mean   15.832061
2   std    10.903883
3   min    1.000000
4   25%    8.000000
5   50%    15.000000
6   75%    20.500000
7   max    60.000000
```

And we can drop the added index:

```
safi_df['years_farm'].describe().reset_index().set_index('index')
```

```
years_farm
index
count  131.000000
mean   15.832061
std    10.903883
min    1.000000
25%    8.000000
50%    15.000000
75%    20.500000
max    60.000000
```

➊ Note

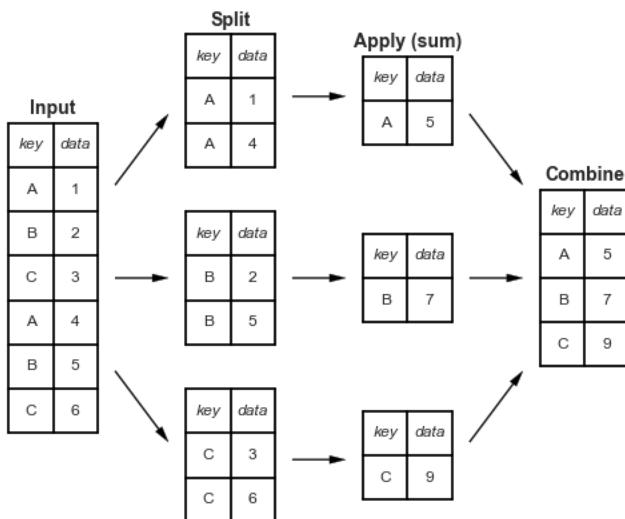
See that we can chain operations together. This is a helpful feature, but to be used with care. [Pep8](#), the style conventions for python, recommends no more than 79 characters per line. A [summary](#) and [another summary](#).

We can also use individual summary statistics on **DataFrame** or **Series** objects

```
safi_df['no_membrs'].max()
```

Split - Apply - Combine

A powerful tool in pandas (and data analysis in general) is the ability to apply functions to subsets of the data.



Tip

one column of a `pd.DataFrame` is a `pd.Series`

Note

In class, we used the `load` magic again here to get this url. Then split the cell, changed the second one to markdown with the code for the image

Tip

To include an image in a notebook, use:

```
![alt text here]
(url/or/path/to/image
)
```

in a markdown cell

For example, we can get descriptive statistics per village

```
safi_df.groupby('village').describe()
```

	quest_no									years_farm	
	count	mean	std	min	25%	50%	75%	max	count	mean	
village											
Chirodzo	39.0	62.487179	44.261705	8.0	44.5	55.0	64.5	200.0	39.0	17.2820	
God	43.0	81.720930	77.863839	1.0	14.5	40.0	168.5	202.0	43.0	14.8837	
Ruaca	49.0	107.061224	55.024013	23.0	72.0	113.0	152.0	194.0	49.0	15.5102	

3 rows × 192 columns

We can also rearrange this into a more usable format than this very wide format:

```
safi_df.groupby('village').describe().unstack().reset_index()
```

	level_0	level_1	village	0
0	quest_no	count	Chirodzo	39.000000
1	quest_no	count	God	43.000000
2	quest_no	count	Ruaca	49.000000
3	quest_no	mean	Chirodzo	62.487179
4	quest_no	mean	God	81.720930
...
571	gps_Accuracy	75%	God	13.500000
572	gps_Accuracy	75%	Ruaca	12.000000
573	gps_Accuracy	max	Chirodzo	39.000000
574	gps_Accuracy	max	God	30.000000
575	gps_Accuracy	max	Ruaca	2099.999000

576 rows × 4 columns

This, however, gives some funny variable names, to fix that, we first save it to a variable.

```
village_summary_df = safi_df.groupby('village').describe().unstack().reset_index()
```

Now we can use the rename function

```
help(village_summary_df.rename)
```

 Tip

Rename is also one thing you might do in cleaning a dataset.

```
Help on method rename in module pandas.core.frame:
```

```
rename(mapper=None, index=None, columns=None, axis=None, copy=True, inplace=False,
       level=None, errors='ignore') method of pandas.core.frame.DataFrame instance
```

```
    Alter axes labels.
```

```
    Function / dict values must be unique (1-to-1). Labels not contained in
    a dict / Series will be left as-is. Extra labels listed don't throw an
    error.
```

```
See the :ref:`user guide <basics.rename>` for more.
```

Parameters

```
-----
```

```
mapper : dict-like or function
    Dict-like or functions transformations to apply to
    that axis' values. Use either ``mapper`` and ``axis`` to
    specify the axis to target with ``mapper``, or ``index`` and
    ``columns``.
index : dict-like or function
    Alternative to specifying axis (``mapper, axis=0``
    is equivalent to ``index=mapper``).
columns : dict-like or function
    Alternative to specifying axis (``mapper, axis=1``
    is equivalent to ``columns=mapper``).
axis : {0 or 'index', 1 or 'columns'}, default 0
    Axis to target with ``mapper``. Can be either the axis name
    ('index', 'columns') or number (0, 1). The default is 'index'.
copy : bool, default True
    Also copy underlying data.
inplace : bool, default False
    Whether to return a new DataFrame. If True then value of copy is
    ignored.
level : int or level name, default None
    In case of a MultiIndex, only rename labels in the specified
    level.
errors : {'ignore', 'raise'}, default 'ignore'
    If 'raise', raise a `KeyError` when a dict-like `mapper`, `index`,
    or `columns` contains labels that are not present in the Index
    being transformed.
    If 'ignore', existing keys will be renamed and extra keys will be
    ignored.
```

Returns

```
-----
```

DataFrame

```
    DataFrame with the renamed axis labels.
```

Raises

```
-----
```

KeyError

```
    If any of the labels is not found in the selected axis and
    "errors='raise'".
```

See Also

```
-----
```

DataFrame.rename_axis

```
: Set the name of the axis.
```

Examples

```
-----
```

```
``DataFrame.rename`` supports two calling conventions
```

```
* ``(index=index_mapper, columns=columns_mapper, ...)``
* ``(mapper, axis={'index', 'columns'}, ...)``
```

```
We *highly* recommend using keyword arguments to clarify your
intent.
```

```
Rename columns using a mapping:
```

```
>>> df = pd.DataFrame({"A": [1, 2, 3], "B": [4, 5, 6]})
>>> df.rename(columns={"A": "a", "B": "c"})
   a   c
0  1   4
1  2   5
2  3   6
```

```
Rename index using a mapping:
```

```
>>> df.rename(index={0: "x", 1: "y", 2: "z"})
   A   B
x  1   4
y  2   5
z  3   6
```

```
Cast index labels to a different type:
```

```

>>> df.index
RangeIndex(start=0, stop=3, step=1)
>>> df.rename(index=str).index
Index(['0', '1', '2'], dtype='object')

>>> df.rename(columns={"A": "a", "B": "b", "C": "c"}, errors="raise")
Traceback (most recent call last):
KeyError: ['C'] not found in axis

Using axis-style parameters

>>> df.rename(str.lower, axis='columns')
   a   b
0  1  4
1  2  5
2  3  6

>>> df.rename({1: 2, 2: 4}, axis='index')
   A   B
0  1  4
2  2  5
4  3  6

```

Rename takes a dict-like or function that maps from what's there to what to change it to. We can either provide the mapper and an axis or pass the mapper as the columns parameter. The columns parameter makes it more human readable, and explicit what we're doing.

```

renaming_dict = {'level_0': 'variable',
                 'level_1': 'statistic',
                 0: 'value'}
village_summary_df.rename(columns= renaming_dict,inplace=True)

```

`inplace` again. See questions at the bottom for more on what it does

Now we have a much better summary table. This is in what's called tidy format. Each column

```
village_summary_df.head()
```

	variable	statistic	village	value
0	quest_no	count	Chirodzo	39.000000
1	quest_no	count	God	43.000000
2	quest_no	count	Ruaca	49.000000
3	quest_no	mean	Chirodzo	62.487179
4	quest_no	mean	God	81.720930

How could we use this to compute the average across villages for each statistic of each variable?

Try it yourself!

What would it look like to do this from the first result of `describe()` on the `groupby`, instead of with the `unstack` and `reset_index`?

We do need to `groupby 'statistic'` and use `mean`, but that's not enough:

```
village_summary_df.groupby('statistic').mean()
```

statistic	value
25%	44.807522
50%	48.428030
75%	54.062308
count	30.388889
max	105.468983
mean	49.868824
min	13.756913
std	24.588967

This averages across all of the variables, too. So instead we need to groupby two variables.

```
village_summary_df.groupby(['statistic','variable']).mean()
```

value		
statistic	variable	
25%	buildings_in_compound	1.000000
	gps_Accuracy	8.000000
	gps_Altitude	691.833333
	gps_Latitude	-19.112221
	gps_Longitude	33.480944
...
std	respondent_wall_type_other	NaN
	rooms	1.082409
	years_farm	10.832121
	years_liv	16.549343
	yes_group_count	1.074600

192 rows × 1 columns

```
safi_df.columns
```

```
Index(['interview_date', 'quest_no', 'start', 'end', 'province', 'district',
       'ward', 'village', 'years_farm', 'agr_assoc', 'note2', 'no_membrs',
       'members_count', 'remittance_money', 'years_liv', 'parents_liv',
       'sp_parents_liv', 'grand_liv', 'sp_grand_liv', 'respondent_roof_type',
       'respondent_wall_type', 'respondent_wall_type_other',
       'respondent_floor_type', 'window_type', 'buildings_in_compound',
       'rooms', 'other_buildings', 'no_plots', 'plots_count', 'water_use',
       'no_group_count', 'yes_group_count', 'no_enough_water',
       'months_no_water', 'period_use', 'exper_other', 'other_meth',
       'res_change', 'memb_assoc', 'resp_assoc', 'fees_water',
       'affect_conflicts', 'note', 'need_money', 'money_source',
       'money_source_other', 'crops_contr', 'empty_lab', 'du_labour',
       'liv_owned', 'liv_owned_other', 'liv_count', 'poultry',
       'du_look_aftr_cows', 'items_owned', 'items_owned_other', 'no_meals',
       'months_lack_food', 'no_food_mitigation', 'gps_Latitude',
       'gps_Longitude', 'gps_Altitude', 'gps_Accuracy', 'instanceID'],
      dtype='object')
```

What is the most common combination of `respondent_wall_type` and `respondent_floor_type`?

```
safi_df.groupby(['respondent_wall_type','respondent_floor_type'])
['instanceID'].count().reset_index()
```

	respondent_wall_type	respondent_floor_type	instanceID
0	burntbricks	cement	30
1	burntbricks	earth	37
2	cement	cement	1
3	muddaub	cement	3
4	muddaub	earth	43
5	sunbricks	cement	4
6	sunbricks	earth	13

We can read this table to see the result. Or we might want it in a different way

```
safi_df.groupby(['respondent_wall_type','respondent_floor_type'])
['instanceID'].count().unstack()
```

```

respondent_floor_type cement earth
respondent_wall_type
    burntbricks    30.0   37.0
        cement      1.0    NaN
    muddaub       3.0   43.0
    sunbricks     4.0   13.0

```

Questions After Class

What does the `inplace` parameter do?

We used `inplace` for the first time today right after we read in the data. Let's make a new DataFrame and compare what happens with and without.

```
safi_df_noinplace = pd.read_csv(data_url)
safi_df_inplace = pd.read_csv(data_url)
```

```
type(safi_df_noinplace.set_index('key_id'))
```

```
pandas.core.frame.DataFrame
```

```
safi_df_noinplace.head()
```

	key_id	interview_date	quest_no	start	end	province	district
0	1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica
1	2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica
2	3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Manica
3	4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Manica
4	5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Manica

5 rows × 65 columns

```
type(safi_df_inplace.set_index('key_id', inplace=True))
```

```
NoneType
```

```
safi_df_inplace.head()
```

Tip

I'm surprised I've remembered it repeatedly in class. I usually forget `inplace` don't get the output I want and then go back and add it. That's a normal part of programming.

	interview_date	quest_no	start	end	province	district	v
key_id							
1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica	E
2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica	E
3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Manica	E
4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Manica	E
5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Manica	E

5 rows × 64 columns

Without `inplace`, `set_index` returns a dataframe, with the index changed, with `inplace` it returns `None`. But, without `inplace`, we don't see the desired effect when we do our next step, but with `inplace` the index is changed.

Without the `inplace`, to save the changes you need to use an assignment. The following is equivalent to using `inplace`.

```
safi_df_noinplace= safi_df_noinplace.set_index('key_id')
safi_df_noinplace.head()
```

```
File "<ipython-input-30-1a84c65392b3>", line 1
    safi_df_noinplace= safi_df_noinplace.set_index('key_id') ^
SyntaxError: invalid syntax
```

Why is a `for` loop slower than a `pandas` operation?

TL;DR: for this example, it's actually possible to relatively easily write a faster loop, but for other operations, it's unlikely.

Basically, the advantage of using the library functions is that someone has already put a lot of thought into the optimal way to implement things and it leverages core features of the data structure. Here is a [blog post](#) about iterating and applying functions in pandas. It covers topics we haven't yet seen, but will.

The `pandas` implementation is also fewer lines and can scale for large datasets easily.

To test, we can do an experiment. Let's take the last thing we covered in class today, finding the most common combination of floor and wall types.

First we'll look at two solutions and verify that they're the same.

First, how to do it with `pandas`

```
wall_floor_df = safi_df.groupby(['respondent_wall_type','respondent_floor_type'])
['instanceID'].count()
wall_floor_df
```

respondent_wall_type	respondent_floor_type	
burntbricks	cement	30
	earth	37
cement	cement	1
muddaub	cement	3
	earth	43
sunbricks	cement	4
	earth	13

Name: instanceID, dtype: int64

Note

For example using `modin` that allows you to change 1 line of code (`import pandas as pd` to `import modin.pandas as pd`) to leverage parallelized computation. Base python loops won't do that.

We can read the answer off of that table, but let's get it programmatically:

```
wall_floor_df.idxmax()
```

```
('muddaub', 'earth')
```

Next, how to do it with a for loop (if you have a better for loop, make a PR so share it).

```
wall_floor_dict = {}

for wall,floor in zip(safi_df['respondent_wall_type'],safi_df['respondent_floor_type']):
    wf_key = '_'.join([wall,floor])
    if wf_key in wall_floor_dict.keys():
        wall_floor_dict[wf_key] +=1
    else:
        wall_floor_dict[wf_key] =1

wall_floor_dict
```

```
{'muddaub_earth': 43,
 'burntbricks_cement': 30,
 'burntbricks_earth': 37,
 'sunbricks_earth': 13,
 'muddaub_cement': 3,
 'sunbricks_cement': 4,
 'cement_cement': 1}
```

Again, we can read from the dict, but lets find it

```
max(wall_floor_dict, key=wall_floor_dict.get).split('_')
```

```
['muddaub', 'earth']
```

Now we can use a special feature of Jupyter notebooks to time them and see which is faster, called the [%timeit](#) magic. Displaying visual things is always slow, so we'll take the line that would display out of both options.

Now timing the pandas way, with a human deciding

```
%timeit -o
wall_floor_df = safi_df.groupby(['respondent_wall_type','respondent_floor_type'])
['instanceID'].count().reset_index()
```

```
2.81 ms ± 37 µs per loop (mean ± std. dev. of 7 runs, 100 loops each)
```

```
<TimeitResult : 2.81 ms ± 37 µs per loop (mean ± std. dev. of 7 runs, 100 loops each)>
```

```
t_pandas = _
```

And capture the time by using the `_` it gets the std out from the previous cell.

For loop with required interpretation

```
%timeit -o
wall_floor_dict = {}

for wall,floor in zip(safi_df['respondent_wall_type'],safi_df['respondent_floor_type']):
    wf_key = '_'.join([wall,floor])
    if wf_key in wall_floor_dict.keys():
        wall_floor_dict[wf_key] +=1
    else:
        wall_floor_dict[wf_key] =1

max(wall_floor_dict, key=wall_floor_dict.get).split('_')
```

```
88.4 µs ± 1.36 µs per loop (mean ± std. dev. of 7 runs, 10000 loops each)
```

💡 Tip

This timing can be good when you have a large dataset, you can load a small part and compare two ways of doing something you want to do with that. Then use only the faster on the whole dataset.

More detail on timing and profiling from the [text book](#)

```
<TimeitResult : 88.4 µs ± 1.36 µs per loop (mean ± std. dev. of 7 runs, 10000 loops each)>
```

```
t_loop = _
```

On my local computer it was about 12x faster with that loop. Here, we'll calculate it for the GitHub servers that host the course manual.

```
t_pandas.average/t_loop.average
```

```
31.751263002305738
```

For a more complex question (say we wanted to know a statistic more complicated than the count) the loop gets harder (and probably slower), but the pandas operation stays about the same.

If you've made it this far, [let me know](#) how you found these notes.

Class 7: Visualization for EDA

Announcements

Syllabus updated

1. [rubric](#) for summarize and visualize are slightly changed
2. [Please accept assignments](#) if you plan to not complete for any reason

Assignment updated to [clarify continuous and categorical variables](#)

Loading Data

Importing the libraries for today. We'll continue plotting with pandas and we'll use [seaborn](#) as well. Seaborn provides higher level plotting functions and [better formatting](#).

```
import pandas as pd
import seaborn as sns
```

Loading the data as usual.

```
data_url = 'https://raw.githubusercontent.com/brownsarahm/python-socialsci-
files/master/data/SAFI_full_shortname.csv'
```

We know that the `key_id` column should be used as an index, not as data, so we'll use the `index_col` parameter to do that from the beginning.

```
safi_df = pd.read_csv(data_url, index_col='key_id')
safi_df.head()
```

 Tip

You can also try it on your computer and see how it compares. Or try other loops/ways of doing it in pandas.

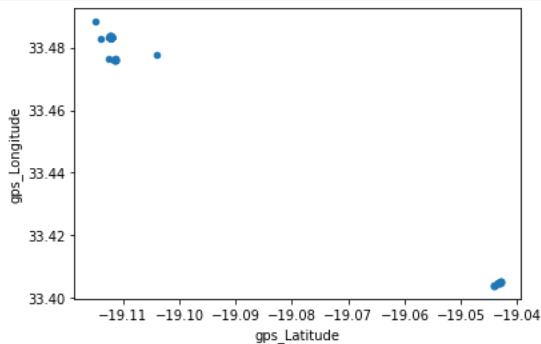
	interview_date	quest_no	start	end	province	district	v
key_id							
1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica	E
2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica	E
3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Manica	E
4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Manica	E
5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Manica	E

5 rows × 64 columns

We can make scatter plots as we saw Friday.

```
safi_df.plot.scatter('gps_Latitude','gps_Longitude')
```

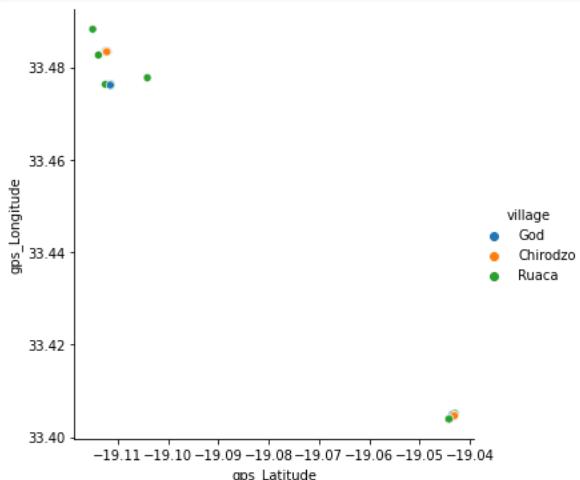
```
<AxesSubplot:xlabel='gps_Latitude', ylabel='gps_Longitude'>
```



With seaborn, however, we can control it more, changing the color of the points based on a column of the data.

```
sns.relplot(x= 'gps_Latitude',y='gps_Longitude',
            data=safi_df, hue='village')
```

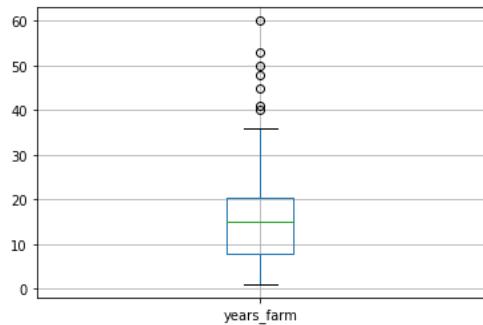
```
<seaborn.axisgrid.FacetGrid at 0x7f288c021310>
```



We can also plot a single variable to see the quantiles (the box is 25%-50%) and see if there are outliers (the points outside the box).

```
safi_df.boxplot('years_farm')
```

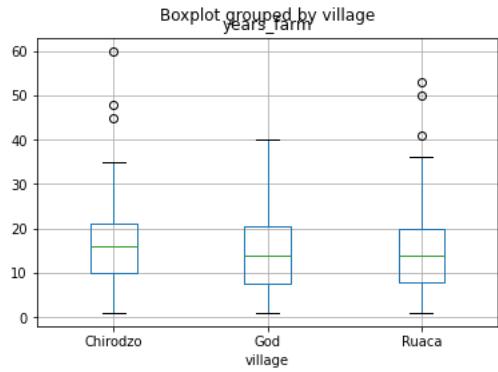
```
<AxesSubplot:>
```



We can do some more conditioning, even with only pandas. Using the `by` parameter will do a `groupby` operation first and then make the plot.

```
safi_df.boxplot('years_farm', by='village')
```

```
<AxesSubplot:title={'center':'years_farm'}, xlabel='village'>
```

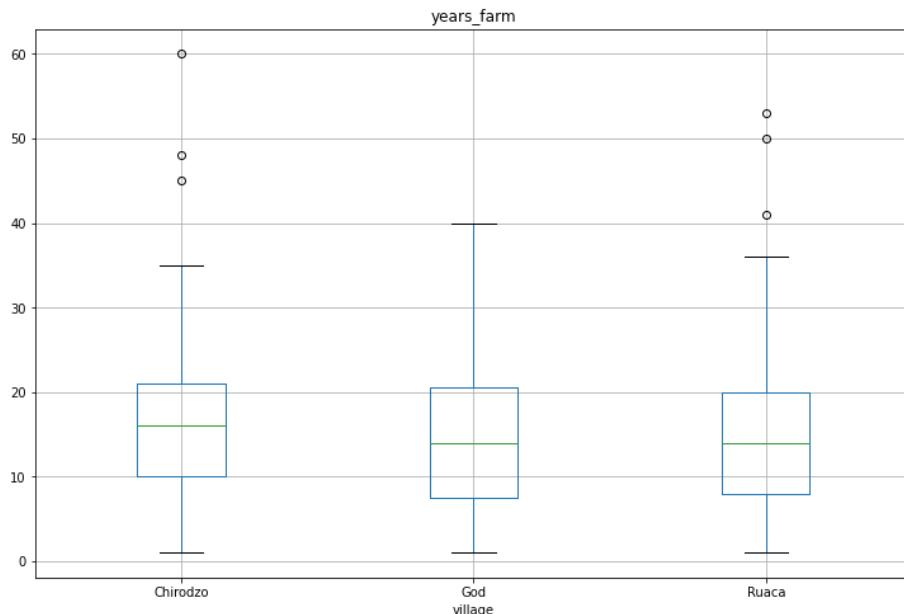


We can also make the figure larger

```
safi_df.boxplot('years_farm', by='village', figsize=(12,8))
```

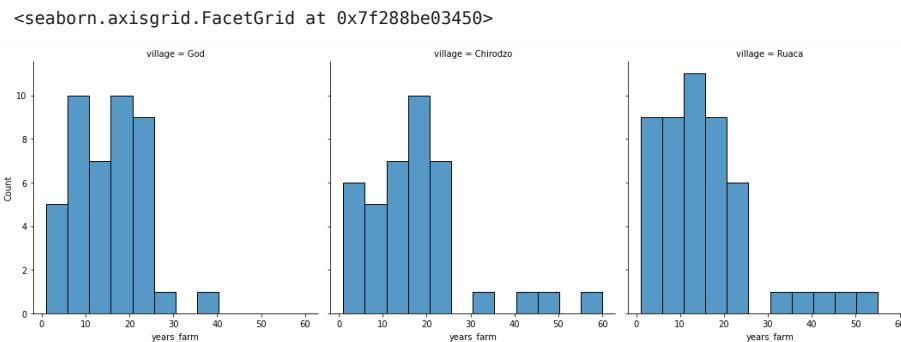
```
<AxesSubplot:title={'center':'years_farm'}, xlabel='village'>
```

Boxplot grouped by village

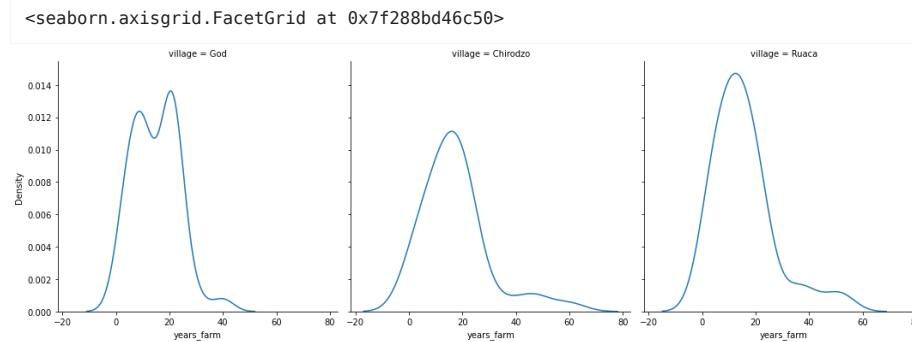


We can see how a single variable is distributed in more detail with the [seaborn displot](#).

```
sns.displot(data=safi_df, x='years_farm', col='village')
```



```
sns.displot(data=safi_df, x='years_farm', col='village', kind='kde')
```



Updating Seaborn

the `displot` is new in seaborn 0.11, on your terminal (mac, linux) or anaconda prompt(Windows):

```
pip install update seaborn
```

Then restart your notebook's kernel and re-run all cells

or, in a notebook, you can update with

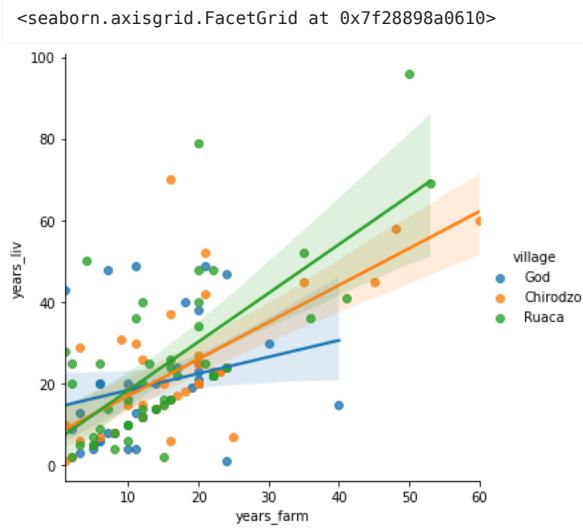
This is set to not run on the served site

```
!pip install seaborn==0.11.0
```

Regression plots

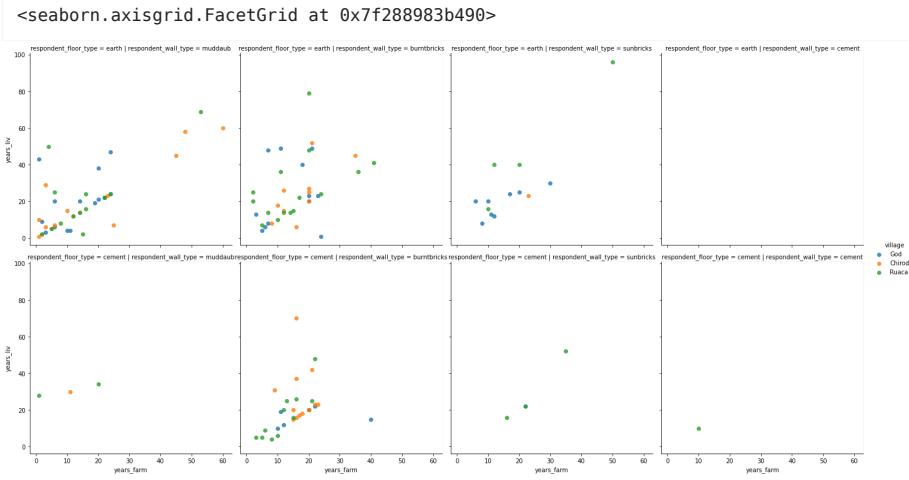
We can also plot with some calculations done for us already.

```
sns.lmplot(x='years_farm',y='years_liv', data=safi_df,  
hue='village')
```



And we can make grids of plots with the `row` and `col` parameters. We can turn off the regression lines with the `fit_reg` parameter.

```
sns.lmplot(x='years_farm',y='years_liv', data=safi_df,  
hue='village', col='respondent_wall_type',row='respondent_floor_type',  
fit_reg=False)
```



```
safi_df.groupby('poultry')['members_count'].mean()
```

```
poultry  
no      5.894737  
yes     7.720430  
Name: members_count, dtype: float64
```

Questions after class

More practice

Test out the parameters of the plotting functions to see what they do.

Further Reading

- [Pandas Plotting](#)
- [Seaborn Gallery](#)
- [Seaborn Tutorial](#)

If you've made it this far, [let me know](#) how you found these notes.

Class 8: Visualization and Starting to Clean Data

Announcements

1. closing notebooks
 - recall that they don't stop when you close the tabs
 - you need to stop it at the terminal it launched from
 - with: ctr + c as it says when you first [launch a notebook](#)
2. restart and rerun notebooks
 - notebooks are a continuous REPL as long as you have it open
 - if you change code, you could have, for example a variable that is no longer defined in the notebook as written, but that still exists in memory, so code that depends on it will still run, for now, but not after you restart next (eg if we run it while grading)
 - to check, [restart and rerun](#) your notebook
3. say hello on zoom for attendance

Setup

First we `import` packages and load data as normal.

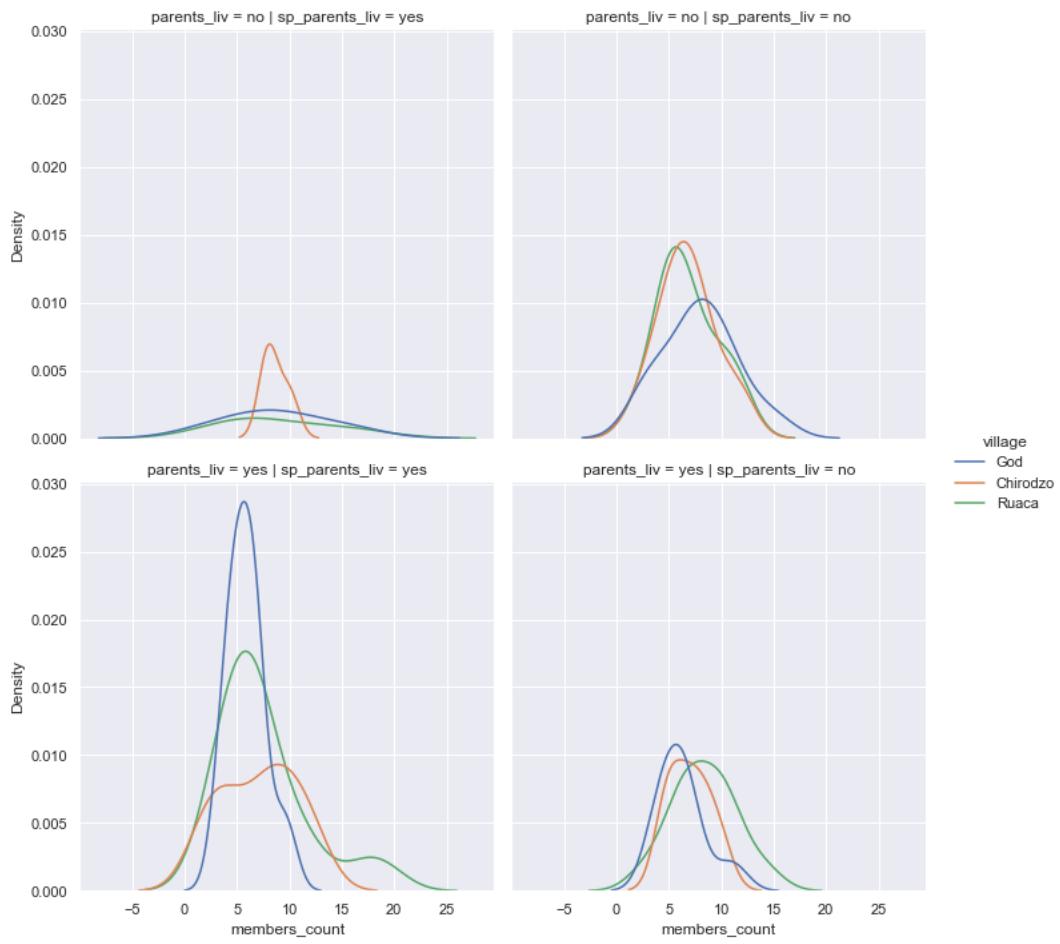
```
import pandas as pd
import seaborn as sns

data_url = 'https://raw.githubusercontent.com/brownsarahm/python-socialsci-
files/master/data/SAFI_full_shortname.csv'
sns.set(font_scale=1.25)
```

I've added one new command here, Seaborn's `[]set` function](<https://seaborn.pydata.org/generated/seaborn.set.html>). It sets a bunch of theme aspects for plotting, here I used it to increase the font size.

Warmup Activity

How recreate this plot?



First, load the data

```
safi_df = pd.read_csv(data_url)
```

This is a [displot](#).

```
sns.displot(data=safi_df, x='members_count', row='parents_liv',
            col = 'sp_parents_liv',hue='village',kind='kde')
```

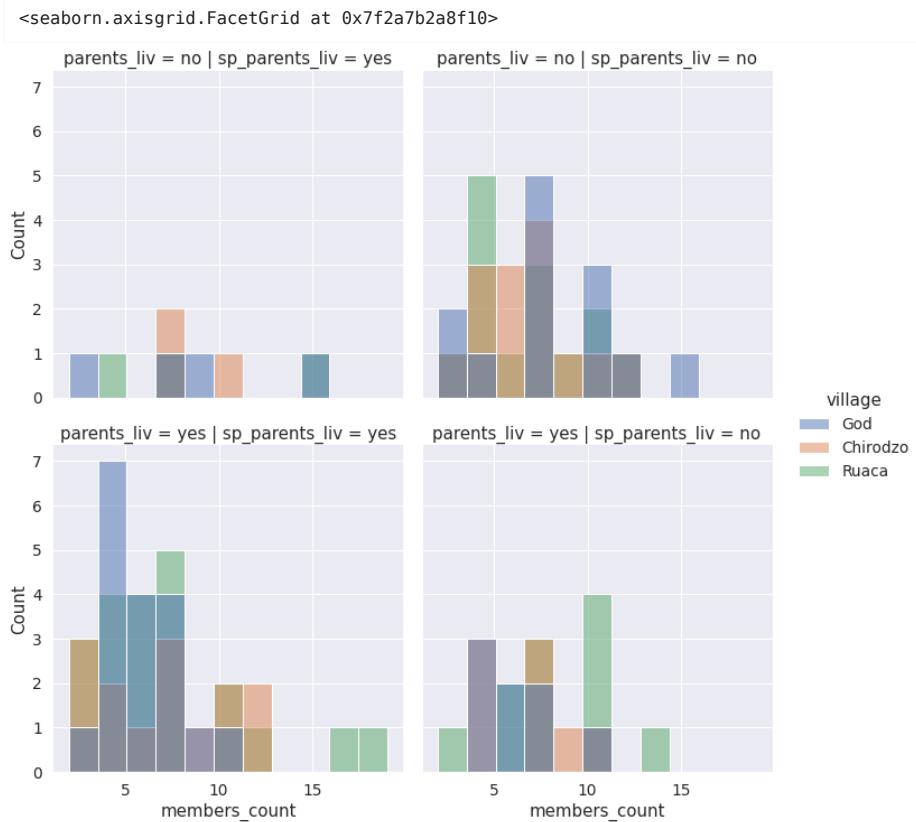
```
<seaborn.axisgrid.FacetGrid at 0x7f2a8094cc90>
```



This allows comparisons of how the village, respondent's parents living (`parents_liv`), and the spouse's (`sp_parents_liv`) influence the number of members of the household (`members_count`). The `kind='kde'` parameter uses the underlying [kdeplot](#) to apply [kernel density estimation](#)

We can understand better what this does, by comparing what happens when we take it out.

```
sns.displot(data=safi_df, x='members_count', row='parents_liv',
            col = 'sp_parents_liv',hue='village')
```



💡 Tip

This is a general strategy. Testing out parameters with different values is a valuable way to learn what they do, and to build intuition about what they do. The documentation also has demos of a lot of values.

What about the rest of the columns?

We've used this SAIFI dataset a lot, but we've only used a few of the many columns. Let's look at all of them.

```
safi_df.columns
```

```
Index(['key_id', 'interview_date', 'quest_no', 'start', 'end', 'province',
       'district', 'ward', 'village', 'years_farm', 'agr_assoc', 'note2',
       'no_membrs', 'members_count', 'remittance_money', 'years_liv',
       'parents_liv', 'sp_parents_liv', 'grand_liv', 'sp_grand_liv',
       'respondent_roof_type', 'respondent_wall_type',
       'respondent_wall_type_other', 'respondent_floor_type', 'window_type',
       'buildings_in_compound', 'rooms', 'other_buildings', 'no_plots',
       'plots_count', 'water_use', 'no_group_count', 'yes_group_count',
       'no_enough_water', 'months_no_water', 'period_use', 'exper_other',
       'other_meth', 'res_change', 'memb_assoc', 'resp_assoc', 'fees_water',
       'affect_conflicts', 'note', 'need_money', 'money_source',
       'money_source_other', 'crops_contr', 'empty_lab', 'du_labour',
       'liv_owned', 'liv_owned_other', 'liv_count', 'poultry',
       'du_look_aftr_cows', 'items_owned', 'items_owned_other', 'no_meals',
       'months_lack_food', 'no_food_mitigation', 'gps_Latitude',
       'gps_Longitude', 'gps_Altitude', 'gps_Accuracy', 'instanceID'],
      dtype='object')
```

We can look at the first few rows to recall what sort of data are available in each column.

```
safi_df.head()
```

	key_id	interview_date	quest_no	start	end	province	district
0	1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica
1	2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica
2	3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Manica
3	4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Manica
4	5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Manica

5 rows × 65 columns

It might be interesting to use the `items_owned` column to compare find out if there are differences between farms based on what items they have.

Maybe we'd want to see how many farms own each item, maybe we want to know how many farms own bicycle. Let's try `value_counts` out like we used before.

```
safi_df['items_owned'].value_counts()
```

```
['radio']
4
['mobile_phone']
4
['bicycle' ; 'radio' ; 'cow_plough' ; 'solar_panel' ; 'table' ; 'mobile_phone']
3
['bicycle' ; 'radio' ; 'cow_plough' ; 'solar_panel' ; 'solar_torch' ;
'mobile_phone']                                3
['bicycle' ; 'radio' ; 'mobile_phone']
3

..
['radio' ; 'solar_torch']
1
['cow_cart' ; 'cow_plough' ; 'solar_torch' ; 'mobile_phone']
1
['cow_cart' ; 'motorcycle' ; 'bicycle' ; 'radio' ; 'sterio' ; 'cow_plough' ;
'solar_panel' ; 'solar_torch' ; 'table' ; 'mobile_phone']    1
['motorcycle' ; 'television' ; 'radio' ; 'cow_plough' ; 'solar_panel' ;
'mobile_phone']                                     1
['cow_cart' ; 'motorcycle' ; 'bicycle' ; 'television' ; 'radio' ; 'cow_plough' ;
'solar_panel' ; 'mobile_phone']                     1
Name: items_owned, Length: 95, dtype: int64
```

The problem is this gives the count of each *list* of items instead of each item. What we'd want instead is one column for each item, where the values in the column are 1 when the farm owns that item and 0 when they don't. This representation is sometimes called 1 hot encoding and other times (including in pandas) it's called dummy variables.

Using Dummy Variables

Let's try this function out, first on the `village` column.

```
pd.get_dummies(data= safi_df,columns=['village'])
```

			key_id	interview_date	quest_no		start		end	province	district
0	1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Mar				
1	2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Mar				
2	3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Mar				
3	4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Mar				
4	5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Mar				
...
126	127	18 May 2017	126	2017-05-18T04:13:37.000Z	2017-05-18T04:35:47.000Z	Manica	Mar				
127	128	04 June 2017	193	2017-06-04T09:36:20.000Z	2017-06-04T10:13:32.000Z	Manica	Mar				
128	129	04 June 2017	194	2017-06-04T10:13:36.000Z	2017-06-04T10:32:06.000Z	Manica	Mar				
129	130	04 June 2017	199	2017-06-04T10:33:55.000Z	2017-06-04T10:52:22.000Z	Manica	Mar				
130	131	04 June 2017	200	2017-06-04T10:52:46.000Z	2017-06-04T11:08:13.000Z	Manica	Mar				

131 rows × 67 columns

Instead of scrolling, we can isolate the new columns we just created.

```
safi_df_villages = pd.get_dummies(data= safi_df,columns=['village'])
keep_cols = [col for col in safi_df_villages.columns if 'village_' in col]
safi_df_villages[keep_cols]
```

	village_Chirodzo	village_God	village_Ruaca
0	0	1	0
1	0	1	0
2	0	1	0
3	0	1	0
4	0	1	0
...
126	0	0	1
127	0	0	1
128	0	0	1
129	1	0	0
130	1	0	0

131 rows × 3 columns

Note that now we have 3 columns, one for each village. This column was already usable, but it was well formatted and useful to illustrate what `get_dummies` does.

Note

try this out and compare the two, test out `get_dummies` on another column to be sure you know what it does.

Now, we can try it on `items_owned`. This time we'll filter the columns for display and inspection right away

```
safi_df_items = pd.get_dummies(data= safi_df,columns=['items_owned'])
keep_cols = [col for col in safi_df_items.columns if 'village_' in col]
safi_df_items[keep_cols]
```

```
0
1
2
3
4
...
126
127
128
129
130
```

131 rows × 0 columns

This still isn't quite what we want, because each value in the `items_owned` column looks like a list but it's actually a string, we can check that with the `type` function.

```
type(safi_df['items_owned'][0])
```

```
str
```

Cleaning One Cell of Data

So, let's clean the value from one cell of that column and see what else is going on. First, we save it to a variable.

```
farm1_items = safi_df['items_owned'][0]
```

Let's look at it first.

```
farm1_items
```

```
["'bicycle' ; 'television' ; 'solar_panel' ; 'table'"]
```

We could try first casting it to a list.

```
list(farm1_items)
```

```
[[],  
 "",  
 'b',  
 'i',  
 'c',  
 'y',  
 'c',  
 'l',  
 'e',  
 "",  
 "",  
 ";",  
 "",  
 "",  
 "",  
 't',  
 'e',  
 'l',  
 'e',  
 'v',  
 'i',  
 's',  
 'i',  
 'o',  
 'n',  
 "",  
 "",  
 ";",  
 "",  
 "",  
 "",  
 's',  
 'o',  
 'l',  
 'a',  
 'r',  
 "-",  
 'p',  
 'a',  
 'n',  
 'e',  
 'l',  
 "",  
 "",  
 ";",  
 "",  
 "",  
 "",  
 't',  
 'a',  
 'b',  
 'l',  
 'e',  
 "",  
 ']' ]
```

That doesn't quite do it though, let's try separating it with the split method at the ';'.

```
farm1_items.split(';')
```

```
["['bicycle' ", " 'television' ", " 'solar_panel' ", " 'table']"]
```

This still has some extra characters, in it though. We should remove those, probably before we do the split so that we don't end up with empty items in the list.

```
farm1_items.replace('[', '').replace(']', '').replace('"', "")
```

```
'bicycle ; television ; solar_panel ; table'
```

That is now ready to split.

```
farm1_items.replace('[', '').replace(']', '').replace('"', "").split(';')
```

```
['bicycle ', ' television ', ' solar_panel ', ' table']
```

There are some empty spaces left. Python has a string function `strip` that removes leading and trailing (on the ends) whitespace (spaces, tabs, new lines), we have to apply it to each individual item of that list above. We can use a list comprehension for this.

```
[i.strip() for i in  
farm1_items.replace('[','').replace(']','').replace('\"','\"').split(';')]
```

```
['bicycle', 'television', 'solar_panel', 'table']
```

Applying this to the rest of the data

We can put that in a function so that we can reuse it.

```
def separate_items(row):  
    '''  
    clean the items owned column of one row of the safi_df  
    '''  
  
    return [i.strip() for i in  
row['items_owned'].replace('[','').replace(']','').replace('\"','\"').split(';')]
```

Pandas provides us special function to apply a function to a dataframe along a given axis.

```
safi_df.apply(separate_items, axis=1)
```

```
-----  
AttributeError                                     Traceback (most recent call last)  
<ipython-input-20-26c2d810f773> in <module>  
----> 1 safi_df.apply(separate_items, axis=1)  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/frame.py in apply(self, func, axis, raw, result_type, args,  
**kwds)  
    7550         kwds=kwds,  
    7551     )  
-> 7552         return op.get_result()  
    7553  
7554     def applymap(self, func) -> "DataFrame":  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/apply.py in get_result(self)  
    178             return self.apply_raw()  
    179  
--> 180         return self.apply_standard()  
    181  
182     def apply_empty_result(self):  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/apply.py in apply_standard(self)  
    269  
    270     def apply_standard(self):  
--> 271         results, res_index = self.apply_series_generator()  
    272  
    273     # wrap results  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/apply.py in apply_series_generator(self)  
    298             for i, v in enumerate(series_gen):  
    299                 # ignore SettingWithCopy here in case the user mutates  
--> 300                 results[i] = self.f(v)  
    301                 if isinstance(results[i], ABCSeries):  
    302                     # If we have a view on v, we need to make a copy  
because  
  
<ipython-input-19-f7c175567037> in separate_items(row)  
    4     ''''  
    5  
----> 6     return [i.strip() for i in  
row['items_owned'].replace('[','').replace(']','').replace('\"','\"').split(';')]  
    7  
  
AttributeError: 'float' object has no attribute 'replace'
```

This error means that there are some elements of that column that is a float instead of a string. We should check what that might be. We can check with a list comprehension and look at the ones that are `float`

```
[item_list for item_list in safi_df['items_owned'] if type(item_list)==float]
```

```
[nan, nan, nan, nan, nan, nan, nan, nan, nan, nan]
```

Since the floats are `nan`, we know that our function more or less is the right thing to do, but we need to modify our function to accommodate those values. We can replace them with an empty list.

```
def separate_items(row):
    """
    ...
    if type(row['items_owned'])==str:
        return [i.strip() for i in
row['items_owned'].replace('[','').replace(']','').replace('\"','").split(';')]
    else:
        return []
```

We can try this again:

```
safi_df['items_owned_clean'] = safi_df.apply(separate_items, axis=1)
```

it runs successfully and we can look at the output.

```
safi_df[['items_owned_clean']].head()
```

items_owned_clean

```
0      [bicycle, television, solar_panel, table]
1  [cow_cart, bicycle, radio, cow_plough, solar_p...
2                  [solar_torch]
3  [bicycle, radio, cow_plough, solar_panel, mobi...
4  [motorcyle, radio, cow_plough, mobile_phone]
```

This looks all good and we'll pick up from here on Monday.

Tip

If this were a regular analysis and not a tutorial, we would probably just edit the cell above, but for the purpose of making this tutorial more readable, we'll copy the function from above, into another cell, edit that cell, and overwrite in python memory by leaving it the same function name.

If instead we wanted to be able to compare the two functions, we'd give the second one a different name.

Class 9: Preparing Data For Analysis

- Say hello in the zoom chat
- join Prismia

Checking types is an important part of cleaning data, we need to figure out what is wrong with data before we can fix it.

Remember that data cleaning is a lot of exploration and iteration in the data. Let's review a few data types we've seen.

Warmup: type review

Lists are enclosed by square brackets (`[]`), they're ordered and iterable.

```
type([char for char in 'abcde'])
```

```
list
```

Dictionaries are comprised of `key:value` pairs and enclosed in curly brackets (`{}`) they're iterable and indexable by the keys.

```
type({char:i for i, char in enumerate('abcde')})
```

```
dict
```

This is a tuple, this data type is used to tie together items, but not usually to iterate over. It's used for pairs of things often or groups to iterate over multiple things at once. For example with the zip function.

```
type(('a', 'b', 'c', 'd', 'e'))
```

```
tuple
```

Splitting a string makes a list.

```
type('a b c d e'.split(' '))
```

```
list
```

Loading Data

Loading the data and packages as normal.

```
# %load http://drsmb.co/310
import pandas as pd
import seaborn as sns
```

```
safi_url = 'https://raw.githubusercontent.com/brownsarahm/python-socialsci-
files/master/data/SAFI_full_shortname.csv'
safi_df = pd.read_csv(safi_url)
```

Exploring Types

On Friday we figured out that the following function would clean one cell of the `safi_df['items_owned']` column.

```
def separate_items(row):
    """
    return [i.strip() for i in
    row['items_owned'].replace('[','').replace(']','').replace('"","").split(';')]
```

But when we applied it, we got an error.

```
safi_df.apply(separate_items, axis=1)
```

```

-----
AttributeError                                     Traceback (most recent call last)
<ipython-input-8-26c2d810f773> in <module>
----> 1 safi_df.apply(separate_items, axis=1)

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/pandas/core/frame.py in apply(self, func, axis, raw, result_type, args,
**kwds)
    7550         kwds=kwds,
    7551     )
-> 7552     return op.get_result()
    7553
7554 def applymap(self, func) -> "DataFrame":


/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/pandas/core/apply.py in get_result(self)
    178         return self.apply_raw()
    179
--> 180     return self.apply_standard()
    181
182     def apply_empty_result(self):


/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/pandas/core/apply.py in apply_standard(self)
    269
    270     def apply_standard(self):
--> 271         results, res_index = self.apply_series_generator()
    272
    273     # wrap results

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/pandas/core/apply.py in apply_series_generator(self)
    298             for i, v in enumerate(series_gen):
    299                 # ignore SettingWithCopy here in case the user mutates
--> 300                 results[i] = self.f(v)
    301                 if isinstance(results[i], ABCSeries):
    302                     # If we have a view on v, we need to make a copy
because

<ipython-input-7-1663c693d570> in separate_items(row)
    3     ''
    4
--> 5     return [i.strip() for i in
row['items_owned'].replace('[','').replace(']', '').replace("'", "").split(';')]

AttributeError: 'float' object has no attribute 'replace'

```

Let's figure out why. First we'll look at the data again to remember what we expected.

```
safi_df['items_owned']
```

```

0      ['bicycle' ;  'television' ;  'solar_panel' ; ...
1      ['cow_cart' ;  'bicycle' ;  'radio' ;  'cow_pl...
2          ...                                ['solar_torch']
3      ['bicycle' ;  'radio' ;  'cow_plough' ;  'sola...
4      ['motorcyle' ;  'radio' ;  'cow_plough' ;  'mo...
        ...
126     ['motorcyle' ;  'radio' ;  'solar_panel']
127     ['car' ;  'lorry' ;  'television' ;  'radio' ; ...
128     ['radio' ;  'solar_panel' ;  'solar_torch' ; ...
129     ['cow_cart' ;  'lorry' ;  'motorcyle' ;  'comp...
130     ['radio' ;  'cow_plough' ;  'solar_panel' ;  ...
Name: items_owned, Length: 131, dtype: object

```

The error says that some item is a float, let's look at the value of only the ones that are `float`

```
[item_list for item_list in safi_df['items_owned'] if type(item_list)==float]
```

```
[nan, nan, nan, nan, nan, nan, nan, nan, nan]
```

Now we can modify the cleaning function

```

def separate_items(row):
    ...
    if type(row['items_owned'])==str:
        return [i.strip() for i in
row['items_owned'].replace('[','').replace(']','').replace('"','"').split(';')]
    else:
        return []

```

and apply it again, to see if it works now.

```
safi_df.apply(separate_items, axis=1)
```

```

0      [bicycle, television, solar_panel, table]
1      [cow_cart, bicycle, radio, cow_plough, solar_p...
2          [solar_torch]
3      [bicycle, radio, cow_plough, solar_panel, mobi...
4          [motorcycle, radio, cow_plough, mobile_phone]
...
126     [motorcycle, radio, solar_panel]
127     [car, lorry, television, radio, sterio, cow_pl...
128     [radio, solar_panel, solar_torch, mobile_phone]
129     [cow_cart, lorry, motorcycle, computer, televis...
130     [radio, cow_plough, solar_panel, solar_torch, ...
Length: 131, dtype: object

```

Dummies From Lists

```
safi_df['items_owned'] = safi_df.apply(separate_items, axis=1)
```

```
pd.get_dummies(safi_df['items_owned'].apply(pd.Series).stack())
```

	bicycle	car	computer	cow_cart	cow_plough	electricity	fridge	lorry	mobile_ph
0	0	1	0	0	0	0	0	0	0
1	0	0	0	0	0	0	0	0	0
2	0	0	0	0	0	0	0	0	0
3	0	0	0	0	0	0	0	0	0
1 0	0	0	0	1	0	0	0	0	0
...
130 1	0	0	0	0	1	0	0	0	0
2	0	0	0	0	0	0	0	0	0
3	0	0	0	0	0	0	0	0	0
4	0	0	0	0	0	0	0	0	0
5	0	0	0	0	0	0	0	0	0

621 rows × 17 columns

```
safi_df['items_owned'].shape
```

```
(131,)
```

```
pd.get_dummies(safi_df['items_owned'].apply(pd.Series).stack()).index
```

```
MultiIndex([( 0,  0),
           ( 0,  1),
           ( 0,  2),
           ( 0,  3),
           ( 1,  0),
           ( 1,  1),
           ( 1,  2),
           ( 1,  3),
           ( 1,  4),
           ( 1,  5),
           ...
           (129,  8),
           (129,  9),
           (129, 10),
           (129, 11),
           (130,  0),
           (130,  1),
           (130,  2),
           (130,  3),
           (130,  4),
           (130,  5)],
          length=621)
```

```
pd.get_dummies(safi_df['items_owned'].apply(pd.Series).stack()).sum(level=0)
```

	bicycle	car	computer	cow_cart	cow_plough	electricity	fridge	lorry	mobile_phone
0	1	0	0	0	0	0	0	0	0
1	1	0	0	1	1	0	0	0	1
2	0	0	0	0	0	0	0	0	0
3	1	0	0	0	1	0	0	0	1
4	0	0	0	0	1	0	0	0	1
...
126	0	0	0	0	0	0	0	0	0
127	0	1	0	0	1	1	1	1	1
128	0	0	0	0	0	0	0	0	1
129	0	0	1	1	1	1	0	1	1
130	0	0	0	0	1	0	0	0	1

121 rows × 17 columns

```
safi_df['items_owned'].head()
```

```
0      [bicycle, television, solar_panel, table]
1      [cow_cart, bicycle, radio, cow_plough, solar_p...
2      [solar_torch]
3      [bicycle, radio, cow_plough, solar_panel, mobi...
4      [motorcycle, radio, cow_plough, mobile_phone]
Name: items_owned, dtype: object
```

```
safi_item_df = pd.get_dummies(safi_df['items_owned'].apply(pd.Series).stack()).sum(level=0)
```

Pandas concat

```
safi_df = pd.concat([safi_df, safi_item_df], axis=1)
```

```
safi_df.shape
```

```
(131, 82)
```

```
safi_df.head()
```

	key_id	interview_date	quest_no	start	end	province	district
0	1	17 November 2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	Manica	Manica
1	2	17 November 2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	Manica	Manica
2	3	17 November 2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	Manica	Manica
3	4	17 November 2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	Manica	Manica
4	5	17 November 2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	Manica	Manica

5 rows × 82 columns

Working with Dummy Variables

```
safi_df['bicycle']
```

0	1.0
1	1.0
2	0.0
3	1.0
4	0.0
	...
126	0.0
127	0.0
128	0.0
129	0.0
130	0.0

Name: bicycle, Length: 131, dtype: float64

```
sum(safi_df['bicycle'])
```

nan

```
safi_df['bicycle'].sum()
```

60.0

```
safi_df['bicycle'].count()
```

121

```
safi_df['bicycle'].shape
```

(131,)

```
pd.concat([safi_df, safi_item_df], axis=0).shape
```

(252, 82)

Questions after class

What does the `get_dummies` function do?

To illustrate, let's first make a small dataframe to look at it

```
ex_data = [['a', 2, 'x'], ['b', 5, 'o'], ['a', 4, 'o'], ['c', 2, 'x'], ['b', 3, 'x']]
ex_df = pd.DataFrame(data = ex_data, columns=['char', 'num', 'symbol'])
ex_df
```

```
char  num  symbol
0    a    2      x
1    b    5      o
2    a    4      o
3    c    2      x
4    b    3      x
```

When we apply `get_dummies` to one column (a `pd.Series`) it makes a column for each value of that column. First we look at that column to focus on what it looks like. Note that there are 3 different values.

```
ex_df['char']
```

```
0    a
1    b
2    a
3    c
4    b
Name: char, dtype: object
```

Now convert it to dummy variables, now we have 3 columns. For each row there is a 1 in the column corresponding to the value and zeros elsewhere.

```
pd.get_dummies(ex_df['char'])
```

```
a  b  c
0  1  0  0
1  0  1  0
2  1  0  0
3  0  0  1
4  0  1  0
```

We can confirm by summing across the rows, it's all ones.

```
pd.get_dummies(ex_df['char']).sum(axis=1)
```

```
0    1
1    1
2    1
3    1
4    1
dtype: int64
```

If we sum down the columns, we get the count of each value

```
pd.get_dummies(ex_df['char']).sum(axis=0)
```

```
a    2
b    2
c    1
dtype: int64
```

it's the same thing as `value_counts()`

```
ex_df['char'].value_counts()
```

```
a    2
b    2
c    1
Name: char, dtype: int64
```

We can also apply it one column, but append it to the whole dataset this way.

```
pd.get_dummies(ex_df,columns=['char'])
```

	num	symbol	char_a	char_b	char_c
0	2	x	1	0	0
1	5	o	0	1	0
2	4	o	1	0	0
3	2	x	0	0	1
4	3	x	0	1	0

This way, it prepends the column values with the original column name. This way concatenates on its own and we don't have to do it separately.

When we apply `get_dummies` on a whole DataFrame, without indicating a column it converts all of the columns of pandas type `object`

```
ex_df.dtypes
```

	char	object
num		int64
symbol		object
dtype:	object	

```
pd.get_dummies(ex_df,)
```

	num	char_a	char_b	char_c	symbol_o	symbol_x
0	2	1	0	0	0	1
1	5	0	1	0	1	0
2	4	1	0	0	1	0
3	2	0	0	1	0	1
4	3	0	1	0	0	1

Class 10: Cleaning review and Ray Summit Keynotes

- Say hello on zoom chat
- join prismia
- sign up so you can watch Ray Summit talks by Pandas and Scikit learn

```
import pandas as pd
```

```
# %load http://drsmb.co/310
data_url = 'https://github.com/rhodyprog4ds/inclass-data/raw/main/ca_dds_summary.xlsx'
```

Let's look at the data

```
pd.read_excel(data_url)
```

	Ethnicity	Asian	Unnamed: 2	Unnamed: 3	Unnamed: 4	Black	Unnamed: 6	Unnamed: 7	Unn	
0	Gender	Female		NaN	Male		NaN	Female	NaN	Male
1	NaN	count		mean	count		mean	count		
2	Age Cohort	Nan		Nan	Nan		Nan	Nan	Nan	
3	0 - 5	6	1544.33		NaN		NaN	NaN	NaN	
4	51 +	9	56577.7		NaN		NaN	NaN	NaN	
5	13-17	10	3385.7		10	3632.5	NaN	NaN	8 43	
6	18-21	18	10083.3		23	9218.52	NaN	NaN	7 91	
7	22-50	12	39470.9		17	39657.9	8	41284.1	9 :	
8	6-12	6	2165.33		12	2164.17	7	2566.57	NaN	

We can read multiple rows in as the header

```
pd.read_excel(data_url,header=list(range(4)))
```

Ethnicity	Asian				Black			
Gender	Female		Male		Female			
Unnamed: 0_level_2	count	mean	count	mean	count	mean		
Age Cohort	Unnamed: 1_level_3	Unnamed: 2_level_3	Unnamed: 3_level_3	Unnamed: 4_level_3	Unnamed: 5_level_3	Unnamed: 6_level_3		
0	0 - 5	6	1544.333333		NaN		NaN	NaN
1	51 +	9	56577.666667		NaN		NaN	NaN
2	13-17	10	3385.700000		10.0	3632.500000		NaN
3	18-21	18	10083.277778		23.0	9218.521739		NaN
4	22-50	12	39470.916667		17.0	39657.882353		8.0 41284.1250
5	6-12	6	2165.333333		12.0	2164.166667		7.0 2566.5714

Looks good, let's save this to a DataFrame

```
ca_dds_df = pd.read_excel(data_url,header=list(range(4)))
```

```
ca_dds_df.head()
```

Ethnicity	Asian				Black			
Gender	Female		Male		Female			
Unnamed: 0_level_2	count	mean	count	mean	count	mean		
Age Cohort	Unnamed: 1_level_3	Unnamed: 2_level_3	Unnamed: 3_level_3	Unnamed: 4_level_3	Unnamed: 5_level_3	Unnamed: 6_level_3		
0	0 - 5	6	1544.333333		NaN		NaN	NaN
1	51 +	9	56577.666667		NaN		NaN	NaN
2	13-17	10	3385.700000		10.0	3632.500000		NaN
3	18-21	18	10083.277778		23.0	9218.521739		NaN
4	22-50	12	39470.916667		17.0	39657.882353		8.0 41284.1250

```
ca_dds_df.columns
```

```

MultiIndex([(  'Ethnicity', 'Gender', 'Unnamed: 0_level_2', ...),
(  ('Asian', 'Female', 'count', ...),
(  ('Asian', 'Female', 'mean', ...),
(  ('Asian', 'Male', 'count', ...),
(  ('Asian', 'Male', 'mean', ...),
(  ('Black', 'Female', 'count', ...),
(  ('Black', 'Female', 'mean', ...),
(  ('Black', 'Male', 'count', ...),
(  ('Black', 'Male', 'mean', ...),
(  ('Hispanic', 'Female', 'count', ...),
(  ('Hispanic', 'Female', 'mean', ...),
(  ('Hispanic', 'Male', 'count', ...),
(  ('Hispanic', 'Male', 'mean', ...),
('White not Hispanic', 'Female', 'count', ...),
('White not Hispanic', 'Female', 'mean', ...),
('White not Hispanic', 'Male', 'count', ...),
('White not Hispanic', 'Male', 'mean', ...)],
)
)

```

Ray Summit Notes

contribute things you learned here

Pandas, by Wes

- Pandas was designed to do data science on your laptop
- It's designed to be coupled tightly to numpy, which is why it's not very fast, especially with strings
-

Scikit Learn

- Data science for the many not the mighty
- Machine learning for all

Class 11: Cleaning Data

```

import pandas as pd

# %load http://drsmr.co/310
data_url = 'https://github.com/rhodyprog4ds/inclass-data/raw/main/ca_dds_summary.xlsx'

```

Recall from Wednesday

```

ca_dds_df = pd.read_excel(data_url, header=list(range(3)))
ca_dds_df

```

	Ethnicity	Asian				Black				
		Gender	Female		Male		Female		Male	
			Unnamed: 0_level_2	count	mean	count	mean	count	mean	count
0	Age Cohort	NaN	NaN	NaN	NaN	NaN	NaN	NaN		
1	0 - 5	6.0	1544.333333	NaN	NaN	NaN	NaN	NaN		
2	51 +	9.0	56577.666667	NaN	NaN	NaN	NaN	NaN		
3	13-17	10.0	3385.700000	10.0	3632.500000	NaN	NaN	8.0		
4	18-21	18.0	10083.277778	23.0	9218.521739	NaN	NaN	7.0		
5	22-50	12.0	39470.916667	17.0	39657.882353	8.0	41284.125000	9.0		
6	6-12	6.0	2165.333333	12.0	2164.166667	7.0	2566.571429	NaN		

```

print(ca_dds_df)

```

	Ethnicity	Asian				Black				\
	Gender	Female		Male		Female		Male		
Unnamed: 0_level_2	count	mean	count	mean	count	mean	count	mean	count	
0	Age Cohort	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	
1	0 - 5	6.0	1544.333333	NaN	NaN	NaN	NaN	NaN	NaN	
2	51 +	9.0	56577.666667	NaN	NaN	NaN	NaN	NaN	NaN	
3	13-17	10.0	3385.700000	10.0	3632.500000	NaN	NaN	NaN	NaN	
4	18-21	18.0	10083.277778	23.0	9218.521739	NaN	NaN	NaN	NaN	
5	22-50	12.0	39470.916667	17.0	39657.882353	8.0	41284.125000	9.0	399	
6	6-12	6.0	2165.333333	12.0	2164.166667	7.0	2566.571429	NaN	NaN	

	Hispanic						\
	Male		Female		Male		
	mean	count	mean	count	mean	count	
0	NaN	NaN	NaN	NaN	NaN	NaN	
1	NaN	NaN	NaN	18.0	1431.333333	26.0	
2	NaN	NaN	NaN	NaN	NaN	12.0	
3	NaN	8.0	4367.250000	50.0	4043.720000	53.0	
4	NaN	7.0	9119.428571	41.0	10199.390244	37.0	
5	41284.125000	9.0	39941.000000	24.0	41005.541667	19.0	
6	2566.571429	NaN	NaN	54.0	2443.777778	37.0	

	White not Hispanic						
	Female		Male				
	mean	count	mean	count	mean		
0	NaN	NaN	NaN	NaN	NaN	NaN	
1	1366.807692	11.0	1316.818182	9.0	1428.111111	NaN	
2	53019.000000	37.0	53449.486486	29.0	51676.448276	NaN	
3	3871.849057	37.0	3818.810811	30.0	4009.866667	NaN	
4	9694.405405	36.0	10452.944444	33.0	9784.090909	NaN	
5	40821.263158	60.0	42113.816667	73.0	38604.452055	NaN	
6	2120.135135	24.0	2112.750000	22.0	1986.272727	NaN	

First we can look at the index

```
ca.dds_df.index
```

```
RangeIndex(start=0, stop=7, step=1)
```

then the first's name

```
ca.dds_df.columns[0]
```

```
('Ethnicity', 'Gender', 'Unnamed: 0_level_2')
```

For this we want to see that as the index

```
ca.dds_df.set_index(ca.dds_df.columns[0], inplace=True)
ca.dds_df
```

	Asian				Black				
	Female		Male		Female		Male		
	count	mean	count	mean	count	mean	count	mean	
(Ethnicity, Gender, Unnamed: 0_level_2)									
Age Cohort	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	
0 - 5	6.0	1544.333333	NaN	NaN	NaN	NaN	NaN	NaN	
51 +	9.0	56577.666667	NaN	NaN	NaN	NaN	NaN	NaN	
13-17	10.0	3385.700000	10.0	3632.500000	NaN	NaN	NaN	8.0	43
18-21	18.0	10083.277778	23.0	9218.521739	NaN	NaN	NaN	7.0	91
22-50	12.0	39470.916667	17.0	39657.882353	8.0	41284.125000	9.0	399	
6-12	6.0	2165.333333	12.0	2164.166667	7.0	2566.571429	NaN	NaN	

We can get rid of rows that are all `nan`, that have no data in them, using `dropna` with `how='all'`

```
ca_dds_df.dropna(how='all', inplace=True)
```

```
ca_dds_df.index
```

```
Index(['0 - 5', '51 +', '13-17', '18-21', '22-50', '6-12'], dtype='object', name= ('Ethnicity', 'Gender', 'Unnamed: 0_level_2'))
```

We want to still rename this, we'll fill that in manually

```
ca_dds_df.index.rename('Age Cohort', inplace=True)
```

Now we want to unstack,

```
ca_dds_df.unstack(level= [0,1])
```

```
-----  
IndexError Traceback (most recent call last)  
<ipython-input-11-06d6e6545889> in <module>  
----> 1 ca_dds_df.unstack(level= [0,1])  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/frame.py in unstack(self, level, fill_value)  
    7144         from pandas.core.reshape.reshape import unstack  
    7145  
-> 7146         return unstack(self, level, fill_value)  
    7147  
    7148     @Appender(  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/reshape/reshape.py in unstack(obj, level, fill_value)  
    406         # _unstack_multiple only handles MultiIndexes,  
    407         # and isn't needed for a single level  
--> 408         return _unstack_multiple(obj, level, fill_value=fill_value)  
    409     else:  
    410         level = level[0]  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/reshape/reshape.py in _unstack_multiple(data, clocs, fill_value)  
    327     if clocs in index.names:  
    328         clocs = [clocs]  
--> 329     clocs = [index._get_level_number(i) for i in clocs]  
    330  
    331     rlocs = [i for i in range(index.nlevels) if i not in clocs]  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/reshape/reshape.py in <listcomp>(.0)  
    327     if clocs in index.names:  
    328         clocs = [clocs]  
--> 329     clocs = [index._get_level_number(i) for i in clocs]  
    330  
    331     rlocs = [i for i in range(index.nlevels) if i not in clocs]  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/indexes/base.py in _get_level_number(self, level)  
    1413  
    1414     def _get_level_number(self, level) -> int:  
--> 1415         self._validate_index_level(level)  
    1416         return 0  
    1417  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/indexes/base.py in _validate_index_level(self, level)  
    1405             elif level > 0:  
    1406                 raise IndexError  
-> 1407                 f"Too many levels: Index has only 1 level, not {level +  
1}"  
    1408  
    1409             elif level != self.name:  
  
IndexError: Too many levels: Index has only 1 level, not 2
```

but this doesn't quite get what we want, instead we can

We can rotate (transpose `.T`) this, then unstack and rotate it back to get the outcome we want here.

```
ca_dds_df.T.unstack(level= [0,1]).T.reset_index()
```

	Age Cohort	level_1	level_2	count	mean
0	0 - 5	Asian	Female	6.0	1544.333333
1	0 - 5	Asian	Male	NaN	NaN
2	0 - 5	Black	Female	NaN	NaN
3	0 - 5	Black	Male	NaN	NaN
4	0 - 5	Hispanic	Female	18.0	1431.333333
5	0 - 5	Hispanic	Male	26.0	1366.807692
6	0 - 5	White not Hispanic	Female	11.0	1316.818182
7	0 - 5	White not Hispanic	Male	9.0	1428.111111
8	51 +	Asian	Female	9.0	56577.666667
9	51 +	Asian	Male	NaN	NaN
10	51 +	Black	Female	NaN	NaN
11	51 +	Black	Male	NaN	NaN
12	51 +	Hispanic	Female	NaN	NaN
13	51 +	Hispanic	Male	12.0	53019.000000
14	51 +	White not Hispanic	Female	37.0	53449.486486
15	51 +	White not Hispanic	Male	29.0	51676.448276
16	13-17	Asian	Female	10.0	3385.700000
17	13-17	Asian	Male	10.0	3632.500000
18	13-17	Black	Female	NaN	NaN
19	13-17	Black	Male	8.0	4367.250000
20	13-17	Hispanic	Female	50.0	4043.720000
21	13-17	Hispanic	Male	53.0	3871.849057
22	13-17	White not Hispanic	Female	37.0	3818.810811
23	13-17	White not Hispanic	Male	30.0	4009.866667
24	18-21	Asian	Female	18.0	10083.277778
25	18-21	Asian	Male	23.0	9218.521739
26	18-21	Black	Female	NaN	NaN
27	18-21	Black	Male	7.0	9119.428571
28	18-21	Hispanic	Female	41.0	10199.390244
29	18-21	Hispanic	Male	37.0	9694.405405
30	18-21	White not Hispanic	Female	36.0	10452.944444
31	18-21	White not Hispanic	Male	33.0	9784.090909
32	22-50	Asian	Female	12.0	39470.916667
33	22-50	Asian	Male	17.0	39657.882353
34	22-50	Black	Female	8.0	41284.125000
35	22-50	Black	Male	9.0	39941.000000
36	22-50	Hispanic	Female	24.0	41005.541667
37	22-50	Hispanic	Male	19.0	40821.263158
38	22-50	White not Hispanic	Female	60.0	42113.816667
39	22-50	White not Hispanic	Male	73.0	38604.452055
40	6-12	Asian	Female	6.0	2165.333333
41	6-12	Asian	Male	12.0	2164.166667
42	6-12	Black	Female	7.0	2566.571429
43	6-12	Black	Male	NaN	NaN
44	6-12	Hispanic	Female	54.0	2443.777778

Age Cohort		level_1	level_2	count	mean
45	6-12	Hispanic	Male	37.0	2120.135135
46	6-12	White not Hispanic	Female	24.0	2112.750000
47	6-12	White not Hispanic	Male	22.0	1986.272727

This looks good, so we'll save it to a new DataFrame

```
ca.dds_clean = ca.dds_df.T.unstack(level= [0,1]).T.reset_index()
ca.dds_clean.head()
```

Age Cohort		level_1	level_2	count	mean
0	0 - 5	Asian	Female	6.0	1544.333333
1	0 - 5	Asian	Male	NaN	NaN
2	0 - 5	Black	Female	NaN	NaN
3	0 - 5	Black	Male	NaN	NaN
4	0 - 5	Hispanic	Female	18.0	1431.333333

We want to rename the two with nondescript names, but the index is not mutable

```
ca.dds_clean.columns[1] = 'Race'
```

```
-----  
TypeError                                                 Traceback (most recent call last)  
<ipython-input-14-f771cccd0e865> in <module>  
----> 1 ca.dds_clean.columns[1] = 'Race'  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/pandas/core/indexes/base.py in __setitem__(self, key, value)  
    4079  
    4080      def __setitem__(self, key, value):  
-> 4081          raise TypeError("Index does not support mutable operations")  
    4082  
    4083      def __getitem__(self, key):  
  
TypeError: Index does not support mutable operations
```

```
type(ca.dds_clean.columns)
```

```
pandas.core.indexes.base.Index
```

Instead we can use rename with a dictionary, this way is good also because it's explicit and readable.

```
ca.dds_clean.rename(columns= {'level_1':'Race', 'level_2':'Gender'}, inplace=True)
ca.dds_clean.head()
```

Age Cohort	Race	Gender	count	mean
0	0 - 5	Asian	Female	6.0
1	0 - 5	Asian	Male	NaN
2	0 - 5	Black	Female	NaN
3	0 - 5	Black	Male	NaN
4	0 - 5	Hispanic	Female	18.0

```
ca.dds_clean['count'][0]
```

```
6.0
```

```
ca.dds_clean.Gender[0]
```

```
'Female'
```

```
ca_dds_clean.Gender[0].lower()
```

```
'female'
```

```
clean_cols = {c:c.lower().replace(' ','_') for c in ca_dds_clean.columns}
ca_dds_clean.rename(columns=clean_cols,inplace=True)
ca_dds_clean.head()
```

	age_cohort	race	gender	count	mean
0	0 - 5	Asian	Female	6.0	1544.333333
1	0 - 5	Asian	Male	NaN	NaN
2	0 - 5	Black	Female	NaN	NaN
3	0 - 5	Black	Male	NaN	NaN
4	0 - 5	Hispanic	Female	18.0	1431.333333

```
ca_dds_clean.fillna('*')
```

	age_cohort	race	gender	count	mean
0	0 - 5	Asian	Female	6	1544.33
1	0 - 5	Asian	Male	*	*
2	0 - 5	Black	Female	*	*
3	0 - 5	Black	Male	*	*
4	0 - 5	Hispanic	Female	18	1431.33
5	0 - 5	Hispanic	Male	26	1366.81
6	0 - 5	White not Hispanic	Female	11	1316.82
7	0 - 5	White not Hispanic	Male	9	1428.11
8	51 +	Asian	Female	9	56577.7
9	51 +	Asian	Male	*	*
10	51 +	Black	Female	*	*
11	51 +	Black	Male	*	*
12	51 +	Hispanic	Female	*	*
13	51 +	Hispanic	Male	12	53019
14	51 +	White not Hispanic	Female	37	53449.5
15	51 +	White not Hispanic	Male	29	51676.4
16	13-17	Asian	Female	10	3385.7
17	13-17	Asian	Male	10	3632.5
18	13-17	Black	Female	*	*
19	13-17	Black	Male	8	4367.25
20	13-17	Hispanic	Female	50	4043.72
21	13-17	Hispanic	Male	53	3871.85
22	13-17	White not Hispanic	Female	37	3818.81
23	13-17	White not Hispanic	Male	30	4009.87
24	18-21	Asian	Female	18	10083.3
25	18-21	Asian	Male	23	9218.52
26	18-21	Black	Female	*	*
27	18-21	Black	Male	7	9119.43
28	18-21	Hispanic	Female	41	10199.4
29	18-21	Hispanic	Male	37	9694.41
30	18-21	White not Hispanic	Female	36	10452.9
31	18-21	White not Hispanic	Male	33	9784.09
32	22-50	Asian	Female	12	39470.9
33	22-50	Asian	Male	17	39657.9
34	22-50	Black	Female	8	41284.1
35	22-50	Black	Male	9	39941
36	22-50	Hispanic	Female	24	41005.5
37	22-50	Hispanic	Male	19	40821.3
38	22-50	White not Hispanic	Female	60	42113.8
39	22-50	White not Hispanic	Male	73	38604.5
40	6-12	Asian	Female	6	2165.33
41	6-12	Asian	Male	12	2164.17
42	6-12	Black	Female	7	2566.57
43	6-12	Black	Male	*	*
44	6-12	Hispanic	Female	54	2443.78

	age_cohort	race	gender	count	mean
45	6-12	Hispanic	Male	37	2120.14
46	6-12	White not Hispanic	Female	24	2112.75
47	6-12	White not Hispanic	Male	22	1986.27

Questions after class

What happens when we use dropna?

We can see the [documentation](#) it drops the whole row or column.

```
import numpy as np #
test_df = pd.DataFrame(data = [[1,3,4,5],[2 ,6, np.nan]])
```

```
-----
AttributeError                                     Traceback (most recent call last)
<ipython-input-22-482bb49100aa> in <module>
      1 import numpy as np #
----> 2 test_df = pd.DataFrame(data = [[1,3,4,5],[2 ,6, np.nan]])

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-packages/numpy/__init__.py
in __getattr__(attr)
    213         else:
    214             raise AttributeError("module {!r} has no attribute "
--> 215                         "{!r}".format(__name__, attr))
    216     def __dir__():

AttributeError: module 'numpy' has no attribute 'nan'
```

Class 12: Constructing Datasets from Multiple Sources

```
import pandas as pd
```

Introduction

Sometimes we use data from multiple sources. For example data may be provided across a bunch of tables and we want to put them together. Or we may want to build our own dataset to answer the questions that we want to ask. For example we have a dataset of drivers who have been pulled over by the police, and need to compare distribution of them between day and night times. If we only have time of being pulled over without mentioning day/night, we can use another dataset including time of sunset for every day and build our own dataset out of these two datasets to find the answer. Later we will also learn about getting data out of databases with some small database operations through SQLite and some python libraries.

To use relative paths as in class (`data/2018-games.csv`) instead of a full url

<https://raw.githubusercontent.com/rhodyprog4ds/inclass-data/main/2018-games.csv>, download data from [this GitHub repo](#) by clicking on the green code button and choosing .zip. Then unzip the data and save it in a data folder in the same folder as the notebook. For the class notes, the urls make it so that the notebook can run without having to store the data in another place.

```
games_df18 = pd.read_csv('https://raw.githubusercontent.com/rhodyprog4ds/inclass-
data/main/2018-games.csv')
games_df19 = pd.read_csv('https://raw.githubusercontent.com/rhodyprog4ds/inclass-
data/main/2019-games.csv')
```

Stacking DataFrames

Let's look at the first couple of rows to see how the configuration of data is.

```
games_df18.head(2)
```

```

Unnamed: 0 GAME_DATE_EST GAME_ID GAME_STATUS_TEXT HOME_TEAM_ID VISITOR_TEAM_ID
0 16196 2019-06-13 41800406 Final 1610612744 1610612761
1 16197 2019-06-10 41800405 Final 1610612761 1610612744
2 rows × 22 columns

```

```
games_df18.head(2)
```

```

Unnamed: 0 GAME_DATE_EST GAME_ID GAME_STATUS_TEXT HOME_TEAM_ID VISITOR_TEAM_ID
0 0 2020-03-01 21900895 Final 1610612766 1610612750
1 1 2020-03-01 21900896 Final 1610612750 1610612766
2 rows × 22 columns

```

As we can see, both datasets have the same columns, and just for two different years.

```
games_df18.shape
```

```
(1378, 22)
```

```
games_df19.shape
```

```
(965, 22)
```

Let's concatenate two dataframes to make one dataframe out of them.

```
games_df = pd.concat([games_df18,games_df19])
games_df.shape
```

```
(2343, 22)
```

As we can see rows are added up but we have the same number of columns.

```
games_df.columns
```

```
Index(['Unnamed: 0', 'GAME_DATE_EST', 'GAME_ID', 'GAME_STATUS_TEXT',
       'HOME_TEAM_ID', 'VISITOR_TEAM_ID', 'SEASON', 'TEAM_ID_home', 'PTS_home',
       'FG_PCT_home', 'FT_PCT_home', 'FG3_PCT_home', 'AST_home', 'REB_home',
       'TEAM_ID_away', 'PTS_away', 'FG_PCT_away', 'FT_PCT_away',
       'FG3_PCT_away', 'AST_away', 'REB_away', 'HOME_TEAM_WINS'],
      dtype='object')
```

Let's drop the column we do not need.

```
games_df.drop(columns= 'Unnamed: 0',inplace=True, )
games_df.head(2)
```

```

GAME_DATE_EST GAME_ID GAME_STATUS_TEXT HOME_TEAM_ID VISITOR_TEAM_ID
0 2019-06-13 41800406 Final 1610612744 1610612761
1 2019-06-10 41800405 Final 1610612761 1610612744
2 rows × 21 columns

```

Merging Data Frames

Now we read another dataset which some of its columns are the same as dataframe "games_df" and some are different.

```
teams_df = pd.read_csv('https://raw.githubusercontent.com/rhodyprog4ds/inclass-data/main/teams.csv')
teams_df.head(2)
```

LEAGUE_ID	TEAM_ID	MIN_YEAR	MAX_YEAR	ABBREVIATION	NICKNAME	YEARF
0	0	1610612737	1949	2019	ATL	Hawks
1	0	1610612738	1946	2019	BOS	Celtics

We use left_on='TEAM_ID' and right_on = 'HOME_TEAM_ID' to match two dataframes.

```
merge1_df = pd.merge(teams_df,games_df, left_on='TEAM_ID', right_on = 'HOME_TEAM_ID')
merge1_df.head(2)
```

LEAGUE_ID	TEAM_ID	MIN_YEAR	MAX_YEAR	ABBREVIATION	NICKNAME	YEARF
0	0	1610612737	1949	2019	ATL	Hawks
1	0	1610612737	1949	2019	ATL	Hawks

2 rows × 35 columns

We want information for each game and append the team info onto that. So, lets try another settings in our merging.

```
merge1_df.shape
```

(2343, 35)

```
merge2_df = pd.merge(teams_df, games_df, left_on='TEAM_ID', right_on = 'HOME_TEAM_ID',
how='outer')
```

```
merge2_df.head(2)
```

LEAGUE_ID	TEAM_ID	MIN_YEAR	MAX_YEAR	ABBREVIATION	NICKNAME	YEARF
0	0	1610612737	1949	2019	ATL	Hawks
1	0	1610612737	1949	2019	ATL	Hawks

2 rows × 35 columns

```
merge2_df.shape
```

(2343, 35)

We can group by "ARENA" and then look at the "mean" statistics.

```
merge1_df.groupby('ARENA').mean()
```

	LEAGUE_ID	TEAM_ID	MIN_YEAR	MAX_YEAR	YEARFOUNDED	A
ARENA						
AT&T Center	0.0	1.610613e+09	1976.000000	2019.0	1976.000000	
American Airlines Center	0.0	1.610613e+09	1980.000000	2019.0	1980.000000	
AmericanAirlines Arena	0.0	1.610613e+09	1988.000000	2019.0	1988.000000	
Amway Center	0.0	1.610613e+09	1989.000000	2019.0	1989.000000	
Bankers Life Fieldhouse	0.0	1.610613e+09	1976.000000	2019.0	1976.000000	
Barclays Center	0.0	1.610613e+09	1976.000000	2019.0	1976.000000	
Capital One Arena	0.0	1.610613e+09	1961.000000	2019.0	1961.000000	
Chase Center	0.0	1.610613e+09	1946.000000	2019.0	1946.000000	
Chesapeake Energy Arena	0.0	1.610613e+09	1967.000000	2019.0	1967.000000	
FedExForum	0.0	1.610613e+09	1995.000000	2019.0	1995.000000	
Fiserv Forum	0.0	1.610613e+09	1968.000000	2019.0	1968.000000	
Golden 1 Center	0.0	1.610613e+09	1948.000000	2019.0	1948.000000	
Little Caesars Arena	0.0	1.610613e+09	1948.000000	2019.0	1948.000000	
Madison Square Garden	0.0	1.610613e+09	1946.000000	2019.0	1946.000000	
Moda Center	0.0	1.610613e+09	1970.000000	2019.0	1970.000000	
Pepsi Center	0.0	1.610613e+09	1976.000000	2019.0	1976.000000	
Quicken Loans Arena	0.0	1.610613e+09	1970.000000	2019.0	1970.000000	
Scotiabank Arena	0.0	1.610613e+09	1995.000000	2019.0	1995.000000	
Smoothie King Center	0.0	1.610613e+09	2002.000000	2019.0	2002.000000	
Spectrum Center	0.0	1.610613e+09	1988.000000	2019.0	1988.000000	
Staples Center	0.0	1.610613e+09	1959.210191	2019.0	1959.210191	
State Farm Arena	0.0	1.610613e+09	1949.000000	2019.0	1949.000000	
TD Garden	0.0	1.610613e+09	1946.000000	2019.0	1946.000000	
Talking Stick Resort Arena	0.0	1.610613e+09	1968.000000	2019.0	1968.000000	
Target Center	0.0	1.610613e+09	1989.000000	2019.0	1989.000000	
Toyota Center	0.0	1.610613e+09	1967.000000	2019.0	1967.000000	
United Center	0.0	1.610613e+09	1966.000000	2019.0	1966.000000	
Vivint Smart Home Arena	0.0	1.610613e+09	1974.000000	2019.0	1974.000000	
Wells Fargo Center	0.0	1.610613e+09	1949.000000	2019.0	1949.000000	

29 rows × 25 columns

How to combine the same data for different outcomes

We can combine datasets in different ways to learn different things about the data. Let's now read info about the players.

```
players18 = pd.read_csv('https://raw.githubusercontent.com/rhodyprog4ds/inclass-data/main/2018-players.csv')
players19 = pd.read_csv('https://raw.githubusercontent.com/rhodyprog4ds/inclass-data/main/2019-players.csv')
players18.head()
```

```

      Unnamed:0  PLAYER_NAME    TEAM_ID  PLAYER_ID  SEASON
0        626    Kawhi Leonard  1610612761     202695  2018
1        627    Pascal Siakam  1610612761     1627783  2018
2        628    Marc Gasol    1610612761     201188  2018
3        629    Danny Green   1610612761     201980  2018
4        630    Kyle Lowry    1610612761     200768  2018

```

First let's look at the shape of each of them to have a reference for what happens when we try different merges.

```
players18.shape, players19.shape
```

```
((748, 5), (626, 5))
```

One thing we might want to do is to put all of the information into one long DataFrame. We can do this with `concat`.

```
pd.concat([players18,players19])
```

```

      Unnamed:0  PLAYER_NAME    TEAM_ID  PLAYER_ID  SEASON
0        626    Kawhi Leonard  1610612761     202695  2018
1        627    Pascal Siakam  1610612761     1627783  2018
2        628    Marc Gasol    1610612761     201188  2018
3        629    Danny Green   1610612761     201980  2018
4        630    Kyle Lowry    1610612761     200768  2018
...
621      621  Anthony Bennett  1610612745     203461  2019
622      622    Ray Spalding  1610612737     1629034  2019
623      623    Devyn Marble  1610612744     203906  2019
624      624  Hassani Gravett  1610612753     1629755  2019
625      625  JaKeenan Gant  1610612754     1629721  2019

```

1374 rows × 5 columns

This allows us to see all the players for each year, we could do `groupby` and `count` to see how many players played each year for example.

We can check that this is the size we expected.

```
pd.concat([players18,players19]).shape
```

```
(1374, 5)
```

If we use the default merge settings we get an empty result because the two DataFrames have the same columns so pandas tries to merge `on` all of the columns, but there are no rows that have the same value in all of the columns, so there's nothing left.

```
pd.merge(players18,players19,)
```

```
      Unnamed:0  PLAYER_NAME  TEAM_ID  PLAYER_ID  SEASON
```

If we merge with `on='PLAYER_ID'`, it only requires that one column to be the same to match rows from the two DataFrames together. With the default value for `how` or explicitly setting `how='inner'` we get the info of players who played both seasons.

```
pd.merge(players18,players19,on='PLAYER_ID', how='inner')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAYE
0	626	Kawhi Leonard	1610612761	202695	2018	299	Kaw
1	627	Pascal Siakam	1610612761	1627783	2018	275	Pas
2	628	Marc Gasol	1610612761	201188	2018	276	
3	1157	Marc Gasol	1610612763	201188	2018	276	
4	629	Danny Green	1610612761	201980	2018	202	D
..
533	1339	Abdul Gaddy	1610612760	203583	2018	561	A
534	1342	Andre Roberson	1610612760	203460	2018	566	Andr
535	1348	Norvel Pelle	1610612755	203658	2018	24	N
536	1350	Denzel Valentine	1610612741	1627756	2018	58	Denze
537	1353	C.J. Wilcox	1610612754	203912	2018	573	

538 rows × 9 columns

When we use `outer` we get one row for each player who played in either season or both seasons. From this we can for example see who changed teams, who are the rookies in 2019 and who retired or was unsigned in 2019.

```
pd.merge(players18,players19,on='PLAYER_ID', how='outer')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAY
0	626.0	Kawhi Leonard	1.610613e+09	202695	2018.0	299.0	K:
1	627.0	Pascal Siakam	1.610613e+09	1627783	2018.0	275.0	P
2	628.0	Marc Gasol	1.610613e+09	201188	2018.0	276.0	
3	1157.0	Marc Gasol	1.610613e+09	201188	2018.0	276.0	
4	629.0	Danny Green	1.610613e+09	201980	2018.0	202.0	
..
922	NaN	NaN	NaN	1629097	NaN	619.0	
923	NaN	NaN	NaN	203461	NaN	621.0	Ant
924	NaN	NaN	NaN	203906	NaN	623.0	D
925	NaN	NaN	NaN	1629755	NaN	624.0	Has
926	NaN	NaN	NaN	1629721	NaN	625.0	Ja

927 rows × 9 columns

Using `left` gives us the '`PLAYER_ID`' that are in the `left(players18)` DataFrame, including those that are in both DataFrame `right` would give players in the `players19` DataFrame or both DataFrames. With this result, we can see who retired, but not the 2019 rookies.

```
pd.merge(players18,players19,on='PLAYER_ID', how='left')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAYE
0	626	Kawhi Leonard	1610612761	202695	2018	299.0	Kav
1	627	Pascal Siakam	1610612761	1627783	2018	275.0	Pa
2	628	Marc Gasol	1610612761	201188	2018	276.0	
3	629	Danny Green	1610612761	201980	2018	202.0	Di
4	630	Kyle Lowry	1610612761	200768	2018	451.0	
..
749	1369	Tyrius Walker	1610612752	1629246	2018	NaN	
750	1370	Marcus Lee	1610612748	1629159	2018	NaN	
751	1371	Trey Lewis	1610612762	1629163	2018	NaN	
752	1372	Emanuel Terry	1610612743	1629150	2018	NaN	
753	1373	Justin Bibbs	1610612738	1629167	2018	NaN	

754 rows × 9 columns

Try it yourself

Try different merges and inspect them:

- how many rows & columns?
- Where are NaN values inserted?
- What rows from the original datasets are not included?
- describe each type of merge in your own words

Split a DataFrame into separate data frames by subsetting the columns and indexing the rows with `loc`, then use concat to put it back together. Programmatically check that it's back together correctly.

Class 13: Data from multiple sources and Databases

Welcome:

1. Say hello on zoom
2. Download data from Mondays' notes if you don't have it already
3. log onto Prismia.chat

```
import pandas as pd
```

Merging review

```
df18_players = pd.read_csv('data/2018-players.csv')
df19_players = pd.read_csv('data/2019-players.csv')
```

```
df18_players.shape, df19_players.shape
```

```
((748, 5), (626, 5))
```

```
pd.merge(df18_players,df19_players,how='inner',on='PLAYER_ID')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAYE
0	626	Kawhi Leonard	1610612761	202695	2018	299	Kav
1	627	Pascal Siakam	1610612761	1627783	2018	275	Pa
2	628	Marc Gasol	1610612761	201188	2018	276	
3	1157	Marc Gasol	1610612763	201188	2018	276	
4	629	Danny Green	1610612761	201980	2018	202	Di
..
533	1339	Abdul Gaddy	1610612760	203583	2018	561	A
534	1342	Andre Roberson	1610612760	203460	2018	566	Andr
535	1348	Norvel Pelle	1610612755	203658	2018	24	N
536	1350	Denzel Valentine	1610612741	1627756	2018	58	Denze
537	1353	C.J. Wilcox	1610612754	203912	2018	573	

538 rows × 9 columns

```
pd.merge(df18_players,df19_players,how='outer',on='PLAYER_ID')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAY
0	626.0	Kawhi Leonard	1.610613e+09	202695	2018.0	299.0	Ka
1	627.0	Pascal Siakam	1.610613e+09	1627783	2018.0	275.0	Pa
2	628.0	Marc Gasol	1.610613e+09	201188	2018.0	276.0	
3	1157.0	Marc Gasol	1.610613e+09	201188	2018.0	276.0	
4	629.0	Danny Green	1.610613e+09	201980	2018.0	202.0	Di
..
922	NaN	NaN	NaN	NaN	1629097	NaN	619.0
923	NaN	NaN	NaN	NaN	203461	NaN	621.0
924	NaN	NaN	NaN	NaN	203906	NaN	623.0
925	NaN	NaN	NaN	NaN	1629755	NaN	624.0
926	NaN	NaN	NaN	NaN	1629721	NaN	625.0

927 rows × 9 columns

```
pd.merge(df18_players,df19_players,how='left',on='PLAYER_ID')
```

	Unnamed: 0_x	PLAYER_NAME_x	TEAM_ID_x	PLAYER_ID	SEASON_x	Unnamed: 0_y	PLAYE
0	626	Kawhi Leonard	1610612761	202695	2018	299.0	Kav
1	627	Pascal Siakam	1610612761	1627783	2018	275.0	Pa
2	628	Marc Gasol	1610612761	201188	2018	276.0	
3	629	Danny Green	1610612761	201980	2018	202.0	Di
4	630	Kyle Lowry	1610612761	200768	2018	451.0	
..
749	1369	Tyrius Walker	1610612752	1629246	2018	NaN	
750	1370	Marcus Lee	1610612748	1629159	2018	NaN	
751	1371	Trey Lewis	1610612762	1629163	2018	NaN	
752	1372	Emanuel Terry	1610612743	1629150	2018	NaN	
753	1373	Justin Bibbs	1610612738	1629167	2018	NaN	

754 rows × 9 columns

```
df18_games = pd.read_csv('data/2018-games.csv')
df19_games = pd.read_csv('data/2019-games.csv')
```

```
games_df = pd.concat([df18_games, df19_games])
games_df.head()
```

```
Unnamed: 0 GAME_DATE_EST GAME_ID GAME_STATUS_TEXT HOME_TEAM_ID VISITOR
0 16196 2019-06-13 41800406 Final 1610612744 1€
1 16197 2019-06-10 41800405 Final 1610612761 1€
2 16198 2019-06-07 41800404 Final 1610612744 1€
3 16199 2019-06-05 41800403 Final 1610612744 1€
4 16200 2019-06-02 41800402 Final 1610612761 1€
```

5 rows × 22 columns

Data

In order to work with large dataset, one option is to use a computer that has more memory, such as an HPC system. An easier option is to load data from a database instead of loading all data into memory. One way of working with databases is to use a Python library named `sqlite3` which allows us to create a connection between the database and Python so that we can pull data into Python to work with it.

⚠️ Warning

To work with a database it needs to be downloaded, we can't load it via url like we did with the `.csv` files that we've been using, because when we need to make a connection to the database that can dynamically interact with the database.

Today, we'll work with a [database version](#) of the same SAFI dataset we've been working with. Download it and put it in a `data` folder that's in the same folder as your notebook.

First we'll import the needed library, this one doesn't have a standard alias, so we'll use it by referring to the whole name.

```
import sqlite3
```

First we'll connect to the database

```
con = sqlite3.connect('data/SQL_SAFI.sqlite')
```

Next we establish a cursor with the connection and then run a query.

```
cur = con.cursor()
cur.execute("SELECT * FROM Farms")
```

```
<sqlite3.Cursor at 0x7fbf450e30a0>
```

Here, "SELECT * FROM Farms" is a query we made into the cursor. It does not return the data, it just says what data to get ready, this is another opportunity to get the data in smaller chunks, we can choose in the next step to bring all or only some of the data from the query's result into Python.

This query says to get all (*) the data from the `Farms` table.

Now, we can use this cursor to pull all of the results from that query into our notebook.

```
rows = cur.fetchall()
rows
```



```
'no',
3,
3,
'yes',
None,
3.0,
'yes',
"[ 'Aug', 'Sept' ]",
2.0,
'yes',
'no',
None,
'yes',
'no',
'no',
'once',
'no',
None,
None,
'yes',
'no',
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None,
3,
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'table', 'mobile_phone' ]",
2,
"[ 'Jan', 'Sept', 'Oct', 'Nov', 'Dec' ]",
"[ 'na', 'reduce_meals', 'restrict_adults', 'borrow_food', 'seek_government' ]",
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19.0,
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'Bandula',
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10,
10,
'no',
15,
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'no',
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'no',
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1,
'no',
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"[ 'none' ]",
None,
1,
'yes',
```

```
'no',
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'Bandula',
'God',
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7,
'no',
6,
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'no',
'no',
'no',
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'burntbricks',
'earth',
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1,
1,
'no',
3,
3,
'no',
3.0,
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'yes',
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2,
'yes',
'no',
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'2017-04-02T17:27:35.000Z',
'Manica',
'Manica',
'Bandula',
'God',
18,
'no',
7,
7,
'no',
40,
```


None,
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None,
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'earth',
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1,
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4.0,
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None,
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```
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2,
1,
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3,
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```

```
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'no',
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"[ 'Jan', 'Dec' ]",
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3,
"[ 'Jan', 'Oct', 'Nov', 'Dec' ]",
```



```
'mabatisloping',
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'yes',
```

```
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'God',
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10,
10,
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20,
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'no',
'yes',
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'burntbricks',
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"[ 'bicycle', 'radio', 'cow_plough', 'solar_panel', 'table', 'mobile_phone']",
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```

```
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30,
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'yes',
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'sunbricks',
'earth',
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'no',
"[ 'bicycle', 'radio', 'cow_plough', 'solar_panel', 'table']",
2,
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(350,
```

We can see it returns the data without much structure

```
type(rows)
```

list

and examine the type of one element of it as well.

```
type(rows[0])
```

tuple

we can also combine use of this library (to access databases) with **Pandas** functions to pull data from databases directly into a datagrame

```
df = pd.read_sql_query("SELECT * FROM Farms",con)
df.head()
```

	Id	Country	A01_interview_date	A03_quest_no	A04_start	A05_end	A
0	1	Moz	17/11/2016		2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z	
1	2	Moz	17/11/2016		2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z	
2	3	Moz	17/11/2016		2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z	
3	4	Moz	17/11/2016		2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z	
4	5	Moz	17/11/2016		2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z	

5 rows × 61 columns

This is the familiar SAFI data that we've been working with

More Specific Queries

There are several ways that we can make our queries return only subsets of the data, this is where the database can start to provide real advantages, we can do sophisticated subsetting of our dataset in the database, which is optimized for performance at exactly these tasks, instead of in Python which is a general purpose programming language.

Structured Query Language (SQL) is a powerful way of working with databases. We'll use it only through pandas going forward so that we can also use the nice visualizations from the notebook, but there are many other ways to use SQL.

For example we might only be interested in the location of the farms that were surveyed, so we can choose only those columns, by putting the column names in place of the * in the query.

```
location_df = pd.read_sql_query("SELECT Country, A06_Province, A07_district, A08_ward,
A09_village FROM Farms",con)
location_df.head()
```

	Country	A06_province	A07_district	A08_ward	A09_village
0	Moz	Manica	Manica	Bandula	God
1	Moz	Manica	Manica	Bandula	God
2	Moz	Manica	Manica	Bandula	God
3	Moz	Manica	Manica	Bandula	God
4	Moz	Manica	Manica	Bandula	God

We can limit for the number of rows we get by using "LIMIT" keyword in the query.

```
pd.read_sql_query("SELECT * FROM Farms LIMIT 10",con)
```

	Id	Country	A01_interview_date	A03_quest_no	A04_start	A05_end
0	1	Moz	17/11/2016	1	2017-03-23T09:49:57.000Z	2017-04-02T17:29:08.000Z
1	2	Moz	17/11/2016	1	2017-04-02T09:48:16.000Z	2017-04-02T17:26:19.000Z
2	3	Moz	17/11/2016	3	2017-04-02T14:35:26.000Z	2017-04-02T17:26:53.000Z
3	4	Moz	17/11/2016	4	2017-04-02T14:55:18.000Z	2017-04-02T17:27:16.000Z
4	5	Moz	17/11/2016	5	2017-04-02T15:10:35.000Z	2017-04-02T17:27:35.000Z
5	6	Moz	17/11/2016	6	2017-04-02T15:27:25.000Z	2017-04-02T17:28:02.000Z
6	7	Moz	17/11/2016	7	2017-04-02T15:38:01.000Z	2017-04-02T17:28:19.000Z
7	8	Moz	16/11/2016	8	2017-04-02T15:59:52.000Z	2017-04-02T17:28:39.000Z
8	9	Moz	16/11/2016	9	2017-04-02T16:23:36.000Z	2017-04-02T16:42:08.000Z
9	10	Moz	16/12/2016	10	2017-04-02T17:03:28.000Z	2017-04-02T17:25:11.000Z

10 rows × 61 columns

➊ Try it Yourself!

How would you return only the first 5 rows of the columns about food?

We can use a “where” keyword to filter and only choose a subset of the rows conditionally, based on *values* of the data. For example, we can look only at the farms that have living parents.

```
parents_df = pd.read_sql_query("SELECT Id, B17_parents_liv FROM Farms WHERE
B17_parents_liv = 'yes'",con)
parents_df.head()
```

	Id	B17_parents_liv
0	2	yes
1	5	yes
2	7	yes
3	8	yes
4	9	yes

Try it yourself

1. Write a queries to read from the database, the Id column and two subsets of columns to 2 separate dataframes. Then use pandas to merge the dataframes into one.
- 2.

Class 15: Intro to ML & Modeling

[handwritten notes from class](#)

What is a Model?

A model is a simplified representation of some part of the world. A famous quote about models is:

All models are wrong, but some are useful –[George Box](#)[^wiki]

You might have seen models in chemistry class, for example about an atom:

read more in the [Model Based Machine Learning Book](#)



Fig. 1 Brighterorange / [CC BY-SA](#)

An atom doesn't actually look like this, but this is a useful representation to help people learn how they function.

In machine learning, we use models, that are generally *statistical* models.

A statistical model is a mathematical model that embodies a set of statistical assumptions concerning the generation of sample data (and similar data from a larger population). A statistical model represents, often in considerably idealized form, the data-generating process

–[wikipedia](#)

Types of Models in Machine Learning

Starting from a dataset, we first make an additional designation about how we will use the different variables (columns). We will call most of them the *features*, which we denote mathematically with \mathbf{X} and we'll choose one to be the *target* or *labels*, denoted by y .

The core assumption for just about all machine learning is that there exists some function f so that for the i th sample

$$y_i = f(\mathbf{x}_i)$$

Then with different additional assumptions we get different types of machine learning:

- if both features (\mathbf{X}) and target (y) are observed (contained in our dataset) it's [supervised learning code](#)
- if only the features (\mathbf{X}) are observed, it's [unsupervised learning code](#)

There are more types, that we won't cover much in class but will mention here for completeness:

- if we have many samples for the features (\mathbf{X}) but only labels for some it's [semi-supervised learning](#)
- if we have only features (\mathbf{X}) and an oracle (a person or some other labeler) that we can ask for labels from with a budget, so we can't get all of them, it's [active learning](#).
- [reinforcement learning](#) involves taking actions and getting rewards.

Tip

a common convention in mathematical notation is to denote matrices (tables) by capitalized bold characters , vectors (lists) by lowercase bold characters and scalars (single values) by lowercase regular characters.

For example:

- all of the features would be: \mathbf{X}
- one variable for all samples or all variables for one sample would be \mathbf{x}
- one variable for one sample would be x

Supervised Learning

we'll focus on supervised learning first. we can take that same core assumption and use it with additional information about our target variable to determine learning **task** we are working to do.

$$y_i = f(\mathbf{x}_i)$$

- if y_i are discrete (eg flower species) we are doing **classification**
- if y_i are continuous (eg height) we are doing **regression**

Machine Learning Pipeline

To do machine learning we start with **training data** which we put as input to the **learning algorithm**. A learning algorithm might be a generic optimization procedure or a specialized procedure for a specific model. The learning algorithm outputs a trained **model** or the parameters of the model. When we deploy a model we pair the **fit model** with a **prediction algorithm** or **decision** algorithm to evaluate a new sample in the world.

In experimenting and design, we need **testing data** to evaluate how well our learning algorithm understood the world. We need to use previously unseen data, because if we don't we can't tell if the prediction algorithm is using a rule that the learning algorithm produced or just looking up from a lookup table the result. This can be thought of like the difference between memorization and understanding.

When the model does well on the training data, but not on test data, we say that it does not generalize well.

Try it yourself

1. List different machine learning applications you've interacted with and try to figure out what the features would be, what the target would be, and then what type of learning it is

Glossary

💡 Tip

This week we've learned a lot of new terms. Contribute definition below to form a glossary.

Term	Definition
Model	a mathematical representation of assumptions about the world

Class 16: Naive Bayes Classification

To learn a classifier, we need labeled data (features and target)

We split our data twice:

- sample-wise: test and train
- variable-wise: features and target

Naive Bayes with Sci-kit Learn

We will use a new package today, [Scikit-Learn](#). Its package name for importing is `sklearn` but we don't import it with an alias, in general. It's a large module and we most often import just the parts we need.

To do that we use a new Python keyword `from`. We can identify a package and then import a submodule or a package and submodule with `.` and then import specific functions or classes.

```
import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
```

💡 Tip

Recall when a word turns green & bold in a notebook, it's a python keyword, or reserved word.

We can tell from this code that `test_train_split` is probably a function because it's in lowercase and `sklearn` follows [PEP 8](#) the Python Style Guide pretty strictly. We can also check with type

```
type(train_test_split)
```

```
function
```

We can tell `GaussianNB` is probably a class because it's in [CapWords](#), also known as [camel case](#).

Again we can check.

```
type(GaussianNB)
```

```
abc.ABCMeta
```

That's an abstract base class.

Today we'll work with the `iris` dataset, which has been used for demonstrating statistical analyses since 1936. It contains 4 measurements of flowers from 3 different species.

```
iris_df = sns.load_dataset('iris')
```

As usual, we look at the structure.

```
iris_df.head()
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

Next we examine the data little further to consider what the structure is like for classification purposes.

```
sns.pairplot(data= iris_df, hue= 'species')
```

```
<seaborn.axisgrid.PairGrid at 0x7f0674d1eb10>
```



In order for classification to work, we're looking to see that the groups of samples of different classes (here, species) do not overlap too much. We're also looking at the shape of the groups of samples of each class.

Naive Bayes

Naive Bayes assumes that the features are uncorrelated (Naive) and that we can pick the most probable class. It can assume different distributions of the features conditioned on the class, though. We'll use Gaussian Naive Bayes. Gaussian distributed samples in the plot above would be roughly ovals with more points at the center and uncorrelated Gaussian data would be distributed in circles, not ovals. We have some ovals and some overlap, but enough we can expect this classifier to work pretty well.

First we instantiate the classifier object with the constructor method.

```
gnb = GaussianNB()
```

We just made it exist so far, nothing more, so we can check its type.

```
type(gnb)
```

```
sklearn.naive_bayes.GaussianNB
```

We can also use the `get_params` method to see what it looks like.

```
gnb.get_params()
```

```
{'priors': None, 'var_smoothing': 1e-09}
```

Training a Model

Before we train the model, we have to get our data out of the DataFrame, because our `gnb.fit` method takes arrays. We can use the `values` attribute

```
iris_df.values
```

```
array([[5.1, 3.5, 1.4, 0.2, 'setosa'],
       [4.9, 3.0, 1.4, 0.2, 'setosa'],
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       [6.7, 3.1, 4.4, 1.4, 'versicolor'],
       [5.6, 3.0, 4.5, 1.5, 'versicolor'],
       [5.8, 2.7, 4.1, 1.0, 'versicolor'],
       [6.2, 2.2, 4.5, 1.5, 'versicolor'],
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       [6.1, 2.8, 4.0, 1.3, 'versicolor'],
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       [5.4, 3.0, 4.5, 1.5, 'versicolor'],
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       [6.7, 3.1, 4.7, 1.5, 'versicolor']],
```

```
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[5.7, 2.8, 4.1, 1.3, 'versicolor'],
[6.3, 3.3, 6.0, 2.5, 'virginica'],
[5.8, 2.7, 5.1, 1.9, 'virginica'],
[7.1, 3.0, 5.9, 2.1, 'virginica'],
[6.3, 2.9, 5.6, 1.8, 'virginica'],
[6.5, 3.0, 5.8, 2.2, 'virginica'],
[7.6, 3.0, 6.6, 2.1, 'virginica'],
[4.9, 2.5, 4.5, 1.7, 'virginica'],
[7.3, 2.9, 6.3, 1.8, 'virginica'],
[6.7, 2.5, 5.8, 1.8, 'virginica'],
[7.2, 3.6, 6.1, 2.5, 'virginica'],
[6.5, 3.2, 5.1, 2.0, 'virginica'],
[6.4, 2.7, 5.3, 1.9, 'virginica'],
[6.8, 3.0, 5.5, 2.1, 'virginica'],
[5.7, 2.5, 5.0, 2.0, 'virginica'],
[5.8, 2.8, 5.1, 2.4, 'virginica'],
[6.4, 3.2, 5.3, 2.3, 'virginica'],
[6.5, 3.0, 5.5, 1.8, 'virginica'],
[7.7, 3.8, 6.7, 2.2, 'virginica'],
[7.7, 2.6, 6.9, 2.3, 'virginica'],
[6.0, 2.2, 5.0, 1.5, 'virginica'],
[6.9, 3.2, 5.7, 2.3, 'virginica'],
[5.6, 2.8, 4.9, 2.0, 'virginica'],
[7.7, 2.8, 6.7, 2.0, 'virginica'],
[6.3, 2.7, 4.9, 1.8, 'virginica'],
[6.7, 3.3, 5.7, 2.1, 'virginica'],
[7.2, 3.2, 6.0, 1.8, 'virginica'],
[6.2, 2.8, 4.8, 1.8, 'virginica'],
[6.1, 3.0, 4.9, 1.8, 'virginica'],
[6.4, 2.8, 5.6, 2.1, 'virginica'],
[7.2, 3.0, 5.8, 1.6, 'virginica'],
[7.4, 2.8, 6.1, 1.9, 'virginica'],
[7.9, 3.8, 6.4, 2.0, 'virginica'],
[6.4, 2.8, 5.6, 2.2, 'virginica'],
[6.3, 2.8, 5.1, 1.5, 'virginica'],
[6.1, 2.6, 5.6, 1.4, 'virginica'],
[7.7, 3.0, 6.1, 2.3, 'virginica'],
[6.3, 3.4, 5.6, 2.4, 'virginica'],
[6.4, 3.1, 5.5, 1.8, 'virginica'],
[6.0, 3.0, 4.8, 1.8, 'virginica'],
[6.9, 3.1, 5.4, 2.1, 'virginica'],
[6.7, 3.1, 5.6, 2.4, 'virginica'],
[6.9, 3.1, 5.1, 2.3, 'virginica'],
[5.8, 2.7, 5.1, 1.9, 'virginica'],
[6.8, 3.2, 5.9, 2.3, 'virginica'],
[6.7, 3.3, 5.7, 2.5, 'virginica'],
[6.7, 3.0, 5.2, 2.3, 'virginica'],
[6.3, 2.5, 5.0, 1.9, 'virginica'],
[6.5, 3.0, 5.2, 2.0, 'virginica'],
[6.2, 3.4, 5.4, 2.3, 'virginica'],
[5.9, 3.0, 5.1, 1.8, 'virginica]], dtype=object)
```

Then we create test and train splits of our data. We'll set equal parts (`test_size = .5`) and set the random state so that we can get the same result.

```
X_train, X_test, y_train, y_test = train_test_split(iris_df.values[:, :4],
                                                    iris_df.values[:, -1],
                                                    test_size = .5,
                                                    random_state = 0)
```

try it yourself!

1. rerun the test `train_test_split` without random state, to see how it's different
2. change the `test_size` to different sizes and see what happens.

Now we use the *training* data to fit our model.

```
gnb.fit(X_train, y_train)
```

```
GaussianNB()
```

Now we can predict using the test data's features only.

```
y_pred = gnb.predict(X_test)  
y_pred
```

```
array(['virginica', 'versicolor', 'setosa', 'virginica', 'setosa',  
      'virginica', 'setosa', 'versicolor', 'versicolor', 'versicolor',  
      'versicolor', 'versicolor', 'versicolor', 'versicolor',  
      'versicolor', 'setosa', 'versicolor', 'versicolor', 'setosa',  
      'virginica', 'versicolor', 'setosa', 'setosa', 'setosa',  
      'virginica', 'setosa', 'versicolor', 'versicolor',  
      'setosa', 'virginica', 'versicolor', 'setosa', 'virginica',  
      'virginica', 'versicolor', 'setosa', 'versicolor', 'versicolor',  
      'versicolor', 'virginica', 'setosa', 'virginica', 'setosa',  
      'setosa', 'versicolor', 'virginica', 'virginica', 'versicolor',  
      'virginica', 'versicolor', 'versicolor', 'virginica', 'versicolor',  
      'virginica', 'versicolor', 'setosa', 'virginica', 'versicolor',  
      'versicolor', 'versicolor', 'versicolor', 'virginica', 'setosa',  
      'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',  
      'versicolor'], dtype='<U10')
```

We can compare this to the y_test to see how well our classifier works.

```
y_test
```

```
array(['virginica', 'versicolor', 'setosa', 'virginica', 'setosa',  
      'virginica', 'setosa', 'versicolor', 'versicolor', 'versicolor',  
      'virginica', 'versicolor', 'versicolor', 'versicolor',  
      'versicolor', 'setosa', 'versicolor', 'versicolor', 'setosa',  
      'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',  
      'virginica', 'setosa', 'setosa', 'versicolor', 'versicolor',  
      'setosa', 'virginica', 'versicolor', 'setosa', 'virginica',  
      'virginica', 'versicolor', 'setosa', 'versicolor', 'versicolor',  
      'versicolor', 'virginica', 'setosa', 'virginica', 'setosa',  
      'setosa', 'versicolor', 'virginica', 'virginica', 'virginica',  
      'virginica', 'versicolor', 'virginica', 'versicolor', 'versicolor',  
      'virginica', 'virginica', 'virginica', 'virginica', 'versicolor',  
      'virginica', 'versicolor', 'setosa', 'virginica', 'versicolor',  
      'versicolor', 'versicolor', 'versicolor', 'virginica', 'setosa',  
      'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',  
      'versicolor'], dtype=object)
```

We can simply use base python to get the number correct with a boolean and sum since `False` turns to 0 and `True` to 1.

```
sum(y_pred == y_test)
```

```
71
```

and compare that with the total

```
len(y_pred)
```

```
75
```

or compute accuracy

```
sum(y_pred == y_test)/len(y_pred)
```

```
0.9466666666666667
```

Questions after class

I'm curious about other classifiers

Great! We'll see more Friday, next week, and next month. You're also encouraged to try out others and experiment with scikit learn.

How could this be related to NLP?

Naive Bayes isn't a good classifier for NLP most of the time because words are not independent, but for very coarse distinctions it can be applied. The *Gaussian* naive Bayes we used today doesn't apply to NLP without a lot of transformation because in general, words are more like categorical (or binary/dummies) variables than continuous valued.

We'll work with text data toward the end of the semester.

How we got the values from the DataFrame

A DataFrame is an object and one of the attributes is [values](#), we accessed that directly. However, there is a new, safer way through the [to_numpy method](#).

Class 17: Evaluating Classification and Midsemester Feedback

1. share your favorite rainy day activity (or just say hi) in the zoom chat for attendance
2. log onto prismia

Naive Bayes Review

Main assumptions:

- classification assumes that features will separate the groups
- NB: conditionally independent

```
# %load http://drsmby.co/310
import pandas as pd
import seaborn as sns
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
```

```
iris = sns.load_dataset("iris")
iris.head()
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

```
X_train, X_test, y_train, y_test = train_test_split(iris.values[:, :4],
                                                    iris.species.values,
                                                    test_size=0.5, random_state=0)
```

```
gnb = GaussianNB()
y_pred = gnb.fit(X_train, y_train).predict(X_test)
```

```
y_pred
```

```
array(['virginica', 'versicolor', 'setosa', 'virginica', 'setosa',
       'virginica', 'setosa', 'versicolor', 'versicolor', 'versicolor',
       'versicolor', 'versicolor', 'versicolor', 'versicolor',
       'versicolor', 'setosa', 'versicolor', 'versicolor', 'setosa',
       'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',
       'virginica', 'setosa', 'setosa', 'versicolor', 'versicolor',
       'setosa', 'virginica', 'versicolor', 'setosa', 'virginica',
       'virginica', 'versicolor', 'setosa', 'versicolor', 'versicolor',
       'versicolor', 'virginica', 'virginica', 'versicolor', 'versicolor',
       'virginica', 'versicolor', 'virginica', 'versicolor', 'versicolor',
       'virginica', 'versicolor', 'setosa', 'virginica', 'versicolor',
       'versicolor', 'versicolor', 'versicolor', 'virginica', 'setosa',
       'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',
       'versicolor'], dtype='<U10')
```

```
y_test
```

```
array(['virginica', 'versicolor', 'setosa', 'virginica', 'setosa',
       'virginica', 'setosa', 'versicolor', 'versicolor', 'versicolor',
       'versicolor', 'versicolor', 'versicolor', 'versicolor',
       'versicolor', 'setosa', 'versicolor', 'versicolor', 'setosa',
       'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',
       'virginica', 'setosa', 'setosa', 'versicolor', 'versicolor',
       'setosa', 'virginica', 'versicolor', 'setosa', 'virginica',
       'virginica', 'versicolor', 'setosa', 'versicolor', 'versicolor',
       'versicolor', 'virginica', 'setosa', 'virginica', 'setosa',
       'setosa', 'versicolor', 'virginica', 'virginica', 'virginica',
       'virginica', 'versicolor', 'virginica', 'versicolor', 'versicolor',
       'virginica', 'virginica', 'virginica', 'virginica', 'versicolor',
       'virginica', 'versicolor', 'setosa', 'virginica', 'versicolor',
       'versicolor', 'versicolor', 'versicolor', 'virginica', 'setosa',
       'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',
       'versicolor'], dtype=object)
```

```
sum(y_pred == y_test)
```

```
71
```

```
len(y_pred)
```

```
75
```

Evaluating Classification

We can use the `score` method to compute accuracy by providing both features and target for the test set.

```
gnb.score(X_test, y_test)
```

```
0.9466666666666667
```

Which is the same as the manual way we did before

```
sum(y_pred == y_test)/len(y_pred)
```

```
0.9466666666666667
```

Scikit learn also provides a whole metrics module. We'll look at two functions from it today.

```
from sklearn.metrics import confusion_matrix, classification_report
```

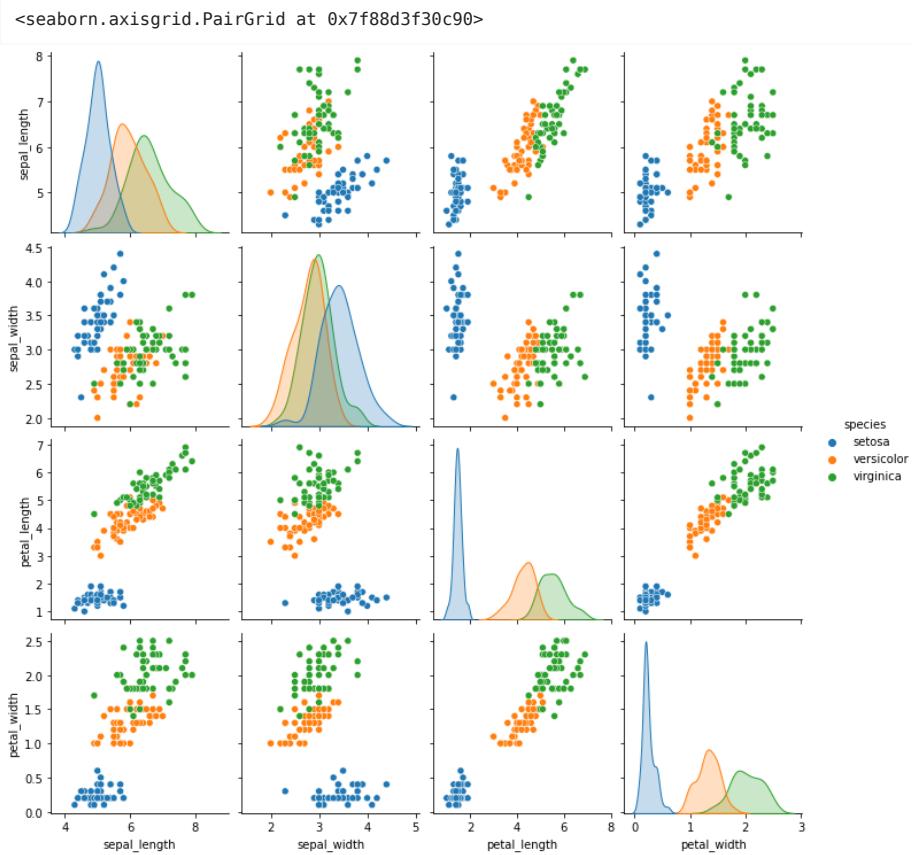
First a Confusion matrix, which is the basic table of how the decisions were made. It counts how many from each true class were predicted to be in each class. My favorite reference on these is the table on the [wikipedia article](#).

```
confusion_matrix(y_test,y_pred,)
```

```
array([[21,  0,  0],
       [ 0, 30,  0],
       [ 0,  4, 20]])
```

We can see this makes sense from looking at the data.

```
sns.pairplot(data =iris, hue='species')
```



We can also get a formatted report that uses the confusion matrix to compute multiple metrics

```
print(classification_report(y_test,y_pred))
```

	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	21
versicolor	0.88	1.00	0.94	30
virginica	1.00	0.83	0.91	24
accuracy			0.95	75
macro avg	0.96	0.94	0.95	75
weighted avg	0.95	0.95	0.95	75

Inspecting a classifier

We can serialize any python object to inspect it with `__dict__`

```
gnb.__dict__
```

```

{'priors': None,
 'var_smoothing': 1e-09,
 'n_features_in_': 4,
 'epsilon_': 3.6399040000000003e-09,
 'classes_': array(['setosa', 'versicolor', 'virginica'], dtype='<U10'),
 'theta_': array([[4.97586207, 3.35862069, 1.44827586, 0.23448276],
 [5.935 , 2.71 , 4.185 , 1.3 ],
 [6.77692308, 3.09230769, 5.73461538, 2.10769231]]),
 'sigma_': array([[0.10321047, 0.13208086, 0.01629013, 0.00846612],
 [0.256275 , 0.0829 , 0.255275 , 0.046 ],
 [0.38869823, 0.10147929, 0.31303255, 0.04763314]]),
 'class_count_': array([29., 20., 26.]),
 'class_prior_': array([0.38666667, 0.26666667, 0.34666667])}

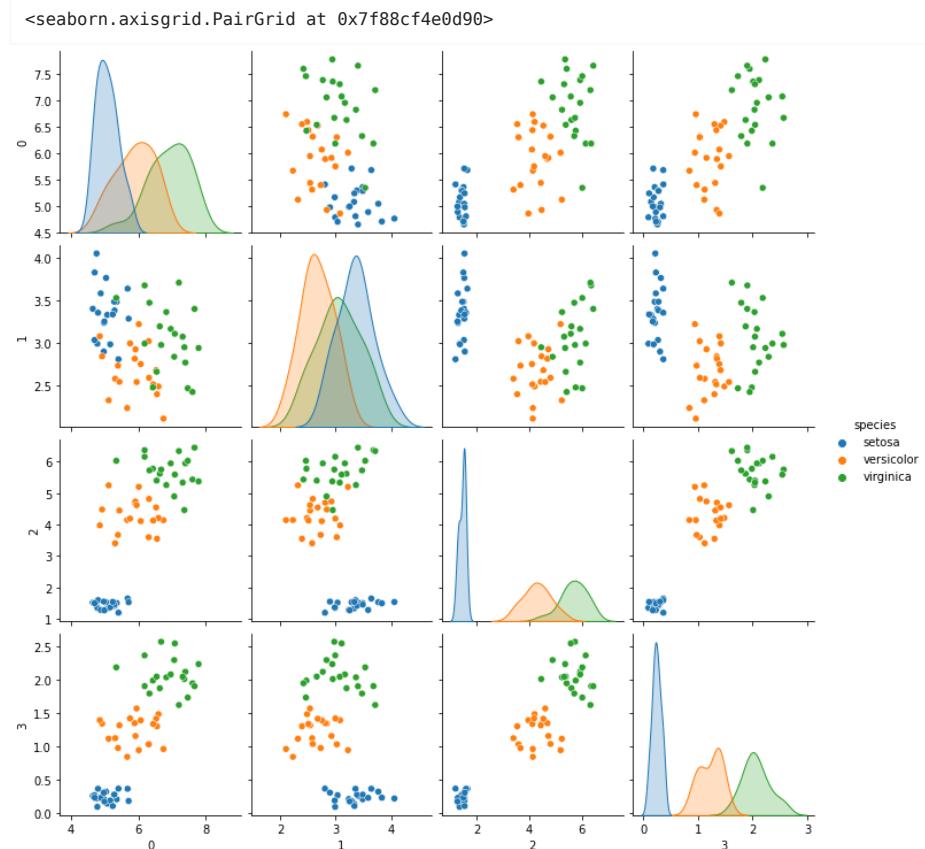
```

It includes attributes `theta_` and `sigma_` these characterize the mean and the variance of each class. Naive Bayes is a generative classifier, since we're using the Gaussian NB, it means it assumes that the data are Gaussian in each class. For a generative classifier, we can draw synthetic data from the model it learned. We can use this to decide if what it learned makes sense by comparing it qualitatively (or statistically) to the real data.

```

# %load http://drsmr.co/310
gnb_df = pd.DataFrame(np.concatenate([np.random.multivariate_normal(mu,
sig*np.eye(4), 20)
for mu, sig in zip(gnb.theta_, gnb.sigma_)]))
gnb_df['species'] = [ci for cl in [[c]*20 for c in gnb.classes_] for ci in cl]
sns.pairplot(data = gnb_df, hue='species')

```



Feedback

Please complete this [feedback survey](#) to provide input on how the semester is going for you so far.

More practice

1. Try breaking down all the steps that are in the last cell one by one to understand them.
2. Read the documentation for naive bayes, how does it use the model parameters to make a prediction?

Class 18: Mid Semester Checkin, Git, & How GNB makes decisions

- Share your favorite thing about fall (or just say hi) in the zoom chat for attendance
- log onto Prismia
- accept Assignment 6
- install git or GitBash if needed

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np
sns.set_theme(font_scale = 2)
```

Feedback Review

Thank you for the feedback. You can add more on Friday's notes if you did not give any yet.

```
feedback_df = pd.read_csv('data/feedback_structured.csv')
feedback_df.head()
```

			Rank the following as what you feel the grading (when you do/not earn achievements) so far actually reflects about your performance in the class [How well I follow instructions]	Rank the following as what you feel the grading (when you do/not earn achievements) so far actually reflects about your performance in the class [What I understand about the material]	Rank the following as what you feel the grading (when you do/not earn achievements) so far actually reflects about your performance in the class [How much effort I put into assignments]
0	4	3	I think they reflect my understanding well	Reflected moderately in the grading	Reflected strongly in the grading
1	4	4	I think the achievements underestimate what I ...	Reflected strongly in the grading	Reflected strongly in the grading
2	3	3	I think they reflect my understanding well	Reflected moderately in the grading	Reflected moderately in the grading
3	4	3	I think they reflect my understanding well	Reflected moderately in the grading	Reflected moderately in the grading
4	3	3	I think they reflect my understanding well	Reflected strongly in the grading	Reflected moderately in the grading

```
el_meaning = {1: 'much less',
 2: 'a little less',
 3: 'about right',
 4: 'a little more',
 5: 'much more'}

expected_learning = feedback_df[feedback_df.columns[0]]
el_counts,_ = np.histogram(expected_learning,bins = [i+.5 for i in range(6)])
el_df = pd.DataFrame(data = el_counts,index = el_meaning.values(),columns=[expected_learning._name],)
# el_df.reset_index(inplace=True)
# el_df = el_df.rename(columns= {'index':'amount'}).set_index('amount')
el_df.rename_axis(index='amount relative to expectations',inplace=True)
el_df.plot.bar(legend=False,title=expected_learning._name,);
```

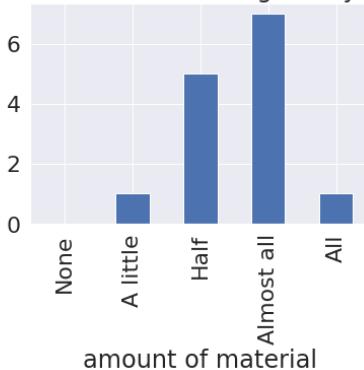
How much do you think you've learned so far this semester?



```
u_meaning = {0:'None', 1:'A little',3:'Half', 4:'Almost all',5:'All'}
u_col = feedback_df[feedback_df.columns[1]]
u_counts,_ = np.histogram(u_col,bins = [i+.5 for i in range(6)])
understanding_df = pd.DataFrame(data = u_counts,index = u_meaning.values(),columns=[u_col._name],)

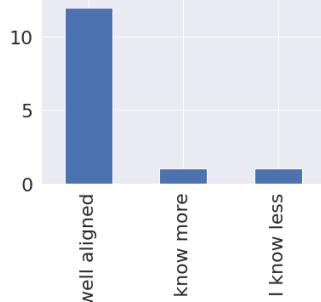
understanding_df.rename_axis(index='amount of material',inplace=True)
understanding_df.plot.bar(legend=False,title=u_col._name,);
```

How much of the material that's been taught do you feel you understand?



```
align = [c_name for c_name in feedback_df.columns if 'align' in c_name][0]
align_df = feedback_df[align].copy()
align_counts = align_df.value_counts()
align_kw = {'well':'well aligned','more':'I know more','less':'I know less'}
## kw_replace =
align_counts.rename(index={s: [align_kw[kw] for kw in align_kw.keys() if kw in s][0]
for s in align_counts.index},inplace=True)
align_counts.plot.bar(title=align_counts._name);
```

How do you think the achievements you've earned so far align with your understanding?



```

reflect_cols = [c_name for c_name in feedback_df.columns if 'reflects' in c_name]
reflect_df = feedback_df[reflect_cols].copy()
reflect_df.rename(columns = lambda c: c.split('[')[1].replace(']', ''), inplace=True)
reflect_df = reflect_df.melt(var_name='factor', value_name='grading reflection')
reflect_df['grading reflection'] = reflect_df['grading reflection'].str.replace(' in the grading', '')
reflect_df['grading reflection'] = reflect_df['grading reflection'].str.replace('Reflected', '')

factor_kw = ['follow instructions', 'understand', 'effort']
# kw_replace =
reflect_df.replace({s: [kw for kw in factor_kw if kw in s][0] for s in
reflect_df['factor']}, inplace=True)
g = sns.displot(data = reflect_df, x='grading reflection', col='factor', hue = 'grading
reflection',)
g.set(xticklabels=[]);


```



```

fair_cols = [c_name for c_name in feedback_df.columns if 'fair' in c_name]
fair_df = feedback_df[fair_cols].copy()
fair_df.rename(columns = lambda c: c.split('[')[1].replace(']', ''), inplace=True)
fair_df = fair_df.melt(var_name='factor', value_name='fairness')
fair_df['fairness'] = fair_df['fairness'].str.replace(' in the grading', '')
# reflect_df['grading reflection'] = reflect_df['grading
reflection'].str.replace('Reflected', '')

factor_kw = ['follow instructions', 'understand', 'effort']
# # kw_replace =
fair_df.replace({s: [kw for kw in factor_kw if kw in s][0] for s in
fair_df['factor']}, inplace=True)
g = sns.displot(data = reflect_df, x='grading reflection', col='factor', hue = 'grading
reflection',)
g.set(xticklabels=[]);
# fair_df

```



```

activity_col = [cname for cname in feedback_df.columns if 'outside' in cname][0]
activity_counts =
pd.get_dummies(feedback_df[activity_col].str.split(',').apply(pd.Series).stack()).sum()
activity_counts = activity_counts.rename_axis(index='activites outside of class')
# activity_counts.plot.bar()
ac_clean = activity_counts.reset_index().rename(columns={0:'students'})
sns.barplot(data=ac_clean,y='activites outside of class',x='students',);

```



Changes based on your feedback for the rest of the semester

- Assignments posted by Wed, due Tuesday (after this one, but I'm giving class time for it)
- Assignments will have a bit more detail or advice (this one is very structured because of the short time, but not all will be this closed).
- Office hours will stay the same, but deadlines moved

- Will collect descriptions of any [issued with Prismia.chat](#) to pass on to the developers

Assignment

- Accept
- Git demo
- Read & make notes of a plan (what do you need to do? what methods will you use?)
- Discuss your plan in a breakout room

Questions:

1. Errors means misclassifications, samples where the classifier's prediction is not correct.

More on Naive Bayes

From last week:

```
# %load http://drsmb.co/310
import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB

iris = sns.load_dataset("iris")
X_train, X_test, y_train, y_test = train_test_split(iris.values[:, :4],
                                                    iris.values[:, -1],
                                                    test_size=0.5, random_state=5)

gnb = GaussianNB()
y_pred = gnb.fit(X_train, y_train).predict(X_test)
```

Remember we can check accuracy with the score method

```
gnb.score(X_test, y_test)
```

```
0.9466666666666667
```

We can inspect the trained classifier

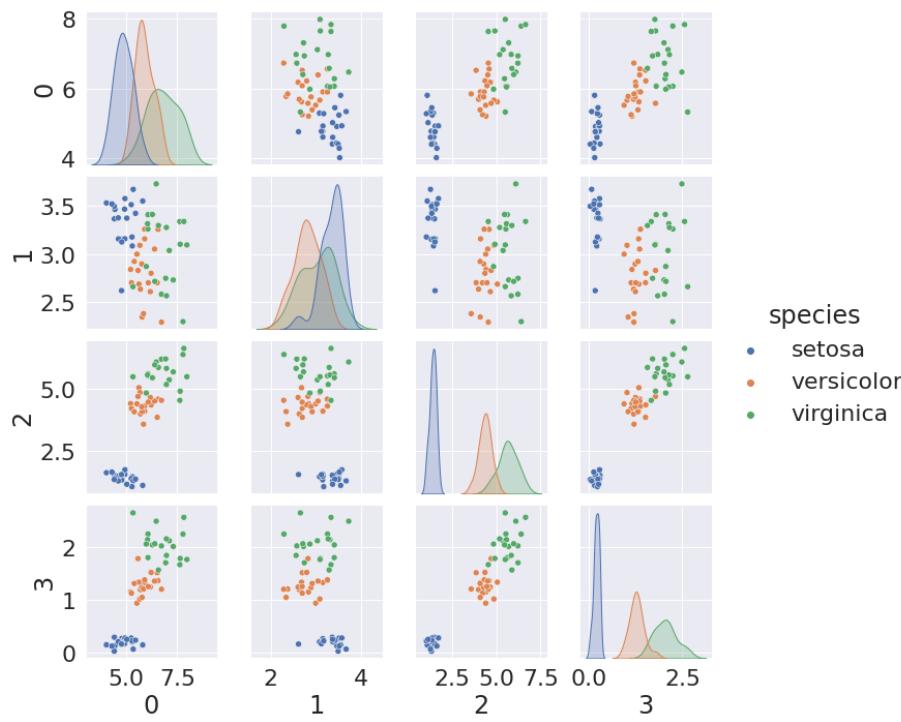
```
gnb.__dict__
```

```
{'priors': None,
 'var_smoothing': 1e-09,
 'n_features_in_': 4,
 'epsilon_': 3.3870506666666674e-09,
 'classes_': array(['setosa', 'versicolor', 'virginica'], dtype='<U10'),
 'theta_': array([[4.98571429, 3.40714286, 1.47142857, 0.23214286],
                 [5.96190476, 2.76666667, 4.34285714, 1.31904762],
                 [6.61923077, 2.95384615, 5.59230769, 2.01153846]]),
 'sigma_': array([[0.15551021, 0.17709184, 0.03275511, 0.00718113],
                  [0.23188209, 0.0984127 , 0.15482994, 0.03392291],
                  [0.48232249, 0.1071006 , 0.34686391, 0.06948225]]),
 'class_count_': array([28., 21., 26.]),
 'class_prior_': array([0.37333333, 0.28      , 0.34666667])}
```

Gaussian Naive Bayes learns the mean and variance for each feature in each class. We can use that to generate synthetic data.

```
df = pd.DataFrame(np.concatenate([np.random.multivariate_normal(mu, sig*np.eye(4), 20)
                                    for mu, sig in zip(gnb.theta_, gnb.sigma_)]))
df['species'] = [ci for cl in [[c]*20 for c in gnb.classes_] for ci in cl]
sns.pairplot(data=df, hue='species')
```

```
<seaborn.axisgrid.PairGrid at 0x7f8642c14a90>
```



It decides for its predictions by computing probabilities and predicting the highest probability one. We can have it return these probabilities directly.

```
gnb.predict_proba(X_test)
```

```
File "<ipython-input-14-e6a42f6fa178>", line 1
    gnb.predict_proba(X_test)
^
SyntaxError: invalid syntax
```

We can use these to see how confident it is on each point. First, we'll get the probability of the predicted class (the max in each row of the matrix).

```
pob_ypred = np.max(gnb.predict_proba(X_test), axis=1)
pob_ypred
```

```
array([0.9999498 , 0.70485761, 0.99999988, 1.        , 1.        ,
       0.99999528, 1.        , 0.88532403, 1.        , 0.98995685,
       0.98928212, 0.69406879, 0.99999999, 0.99999997, 1.        ,
       1.        , 0.90927501, 0.99999997, 1.        , 1.        ,
       0.99972898, 0.99999839, 1.        , 0.92449508, 0.99970908,
       0.99999997, 0.99857943, 0.99965594, 0.98844145, 0.99999983,
       1.        , 0.99993213, 0.9999939 , 1.        , 0.80885454,
       1.        , 1.        , 1.        , 1.        , 0.99999993,
       0.98915134, 0.99998516, 1.        , 1.        , 0.95513579,
       0.85016072, 0.99525954, 0.99999988, 1.        , 1.        ,
       0.98735446, 1.        , 1.        , 1.        , 1.        ,
       0.99906987, 0.67288225, 0.99999514, 0.9999897 , 0.96679497,
       0.99999729, 0.99986416, 0.99999998, 0.55849652, 1.        ,
       0.98840745, 1.        , 0.94307153, 0.99999991, 0.99990409,
       1.        , 0.99977424, 0.92205013, 0.99937796, 0.99994699])
```

Now, we'll put the test data into a `DataFrame` so we can use high level plotting functions from `pandas` and `seaborn`. We could use the `numpy` data structures that scikit learn uses directly with `matplotlib` plotting and `numpy` statistics, but this is a little more compact and readable.

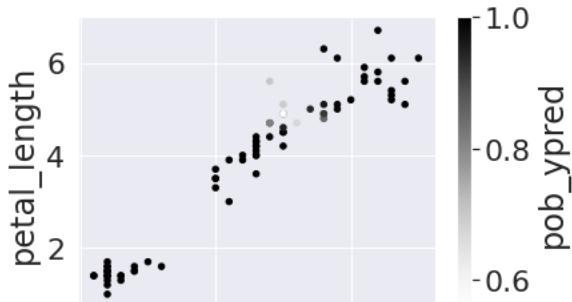
```
iris_test_df = pd.DataFrame(X_test, columns = iris.columns[:4])
iris_test_df['species'] = y_test
iris_test_df['species_pred'] = y_pred
iris_test_df['pob_ypred'] = pob_ypred
iris_test_df.head()
```

	sepal_length	sepal_width	petal_length	petal_width	species	species_pred	pob_ypred
0	5.8	2.7	3.9	1.2	versicolor	versicolor	0.999950
1	6.1	2.6	5.6	1.4	virginica	virginica	0.704858
2	5.8	2.8	5.1	2.4	virginica	virginica	1.000000
3	4.4	3.2	1.3	0.2	setosa	setosa	1.000000
4	7.2	3.6	6.1	2.5	virginica	virginica	1.000000

First we'll look at the one pairwise view of the data that was the most visually separate, 'petal_width' vs 'petal_length'

```
iris_test_df.plot.scatter(x='petal_width', y = 'petal_length',c='pob_ypred')
```

<AxesSubplot:xlabel='petal_width', ylabel='petal_length'>



Notice that the points near the boundary are the lowest probability. We can inspect further by adding a 'correct' column with boolean values

```
iris_test_df['correct'] = iris_test_df['species'] == iris_test_df['species_pred']
```

Now we can use EDA (exploratory data analysis) to further examine.

```
iris_test_df.groupby('correct')['pob_ypred'].describe()
```

	count	mean	std	min	25%	50%	75%	max
correct								
False	4.0	0.770240	0.176779	0.558497	0.660176	0.789696	0.899761	0.943072
True	71.0	0.980814	0.059866	0.672882	0.999224	1.000000	1.000000	1.000000

The incorrectly (`False`) predicted samples were much lower confidence and higher variance in confidence. This suggests that the classifier is a good fit for the data, though it's imperfect on the test set, and that we should trust how it's working.

Try it yourself

1. Can any of these things help with Assignment 6?
2. Review the datasets you've used for prior analyses, can they be used for any classification?
3. What kinds of data would be needed to train a classifier for some ML systems that you interact with?

Class 19: Decision Trees

1. log onto Prismia
2. say hello in the zoom chat

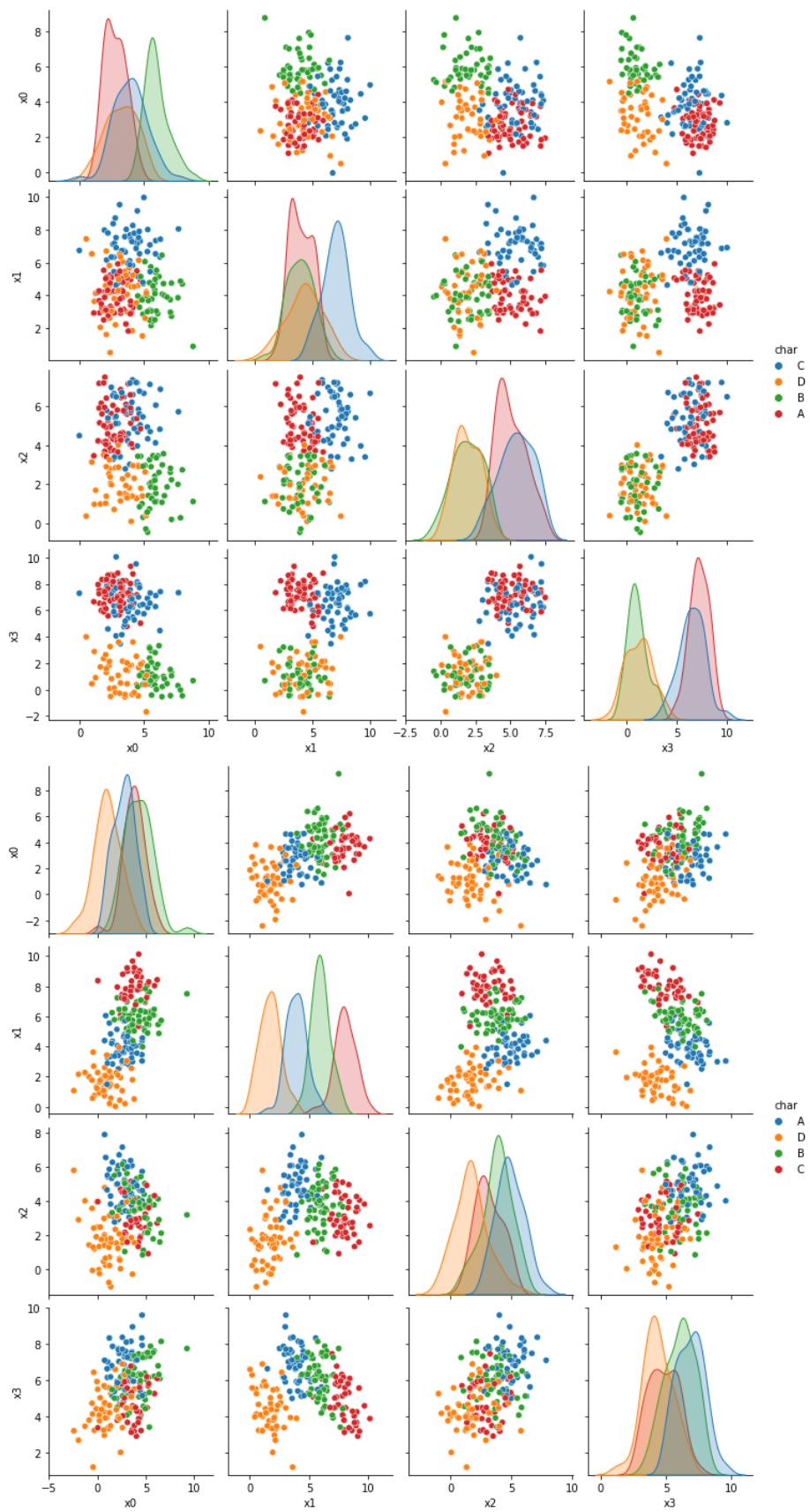
```
# %load http://drsmb.co/310
import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.datasets import load_iris
from sklearn import tree
```

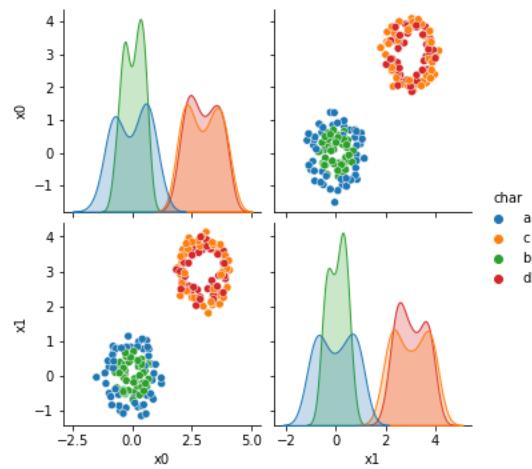
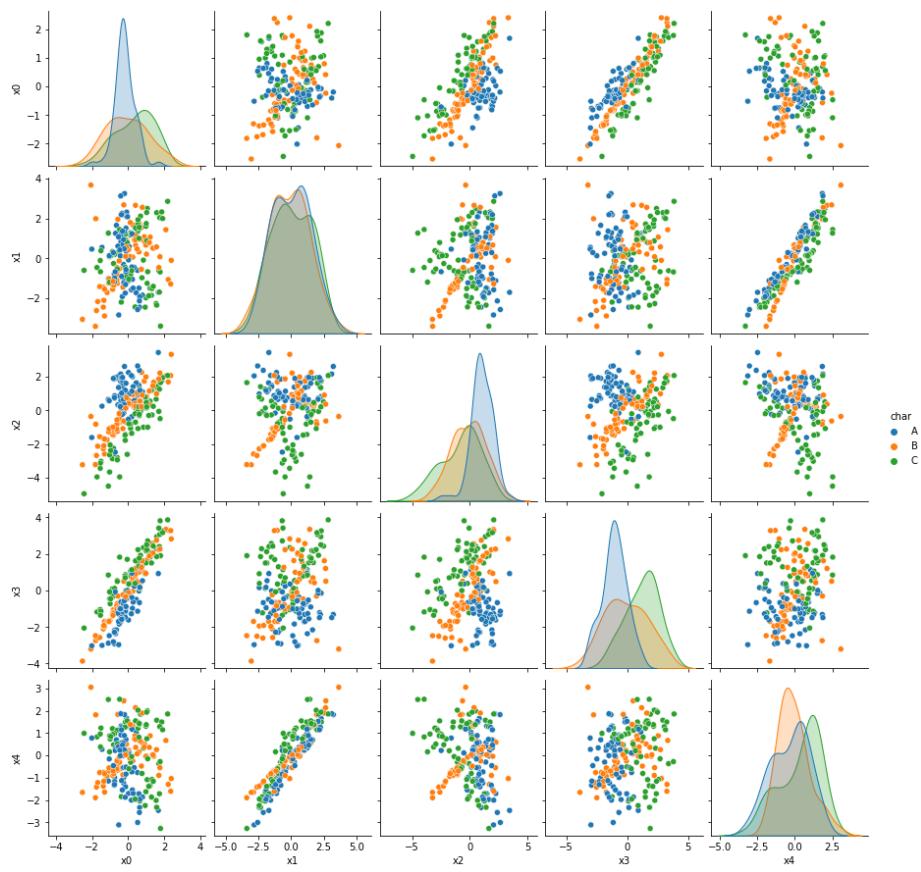
First we'll review the datasets from the assignment briefly. More details on them will be on the solutions repo.

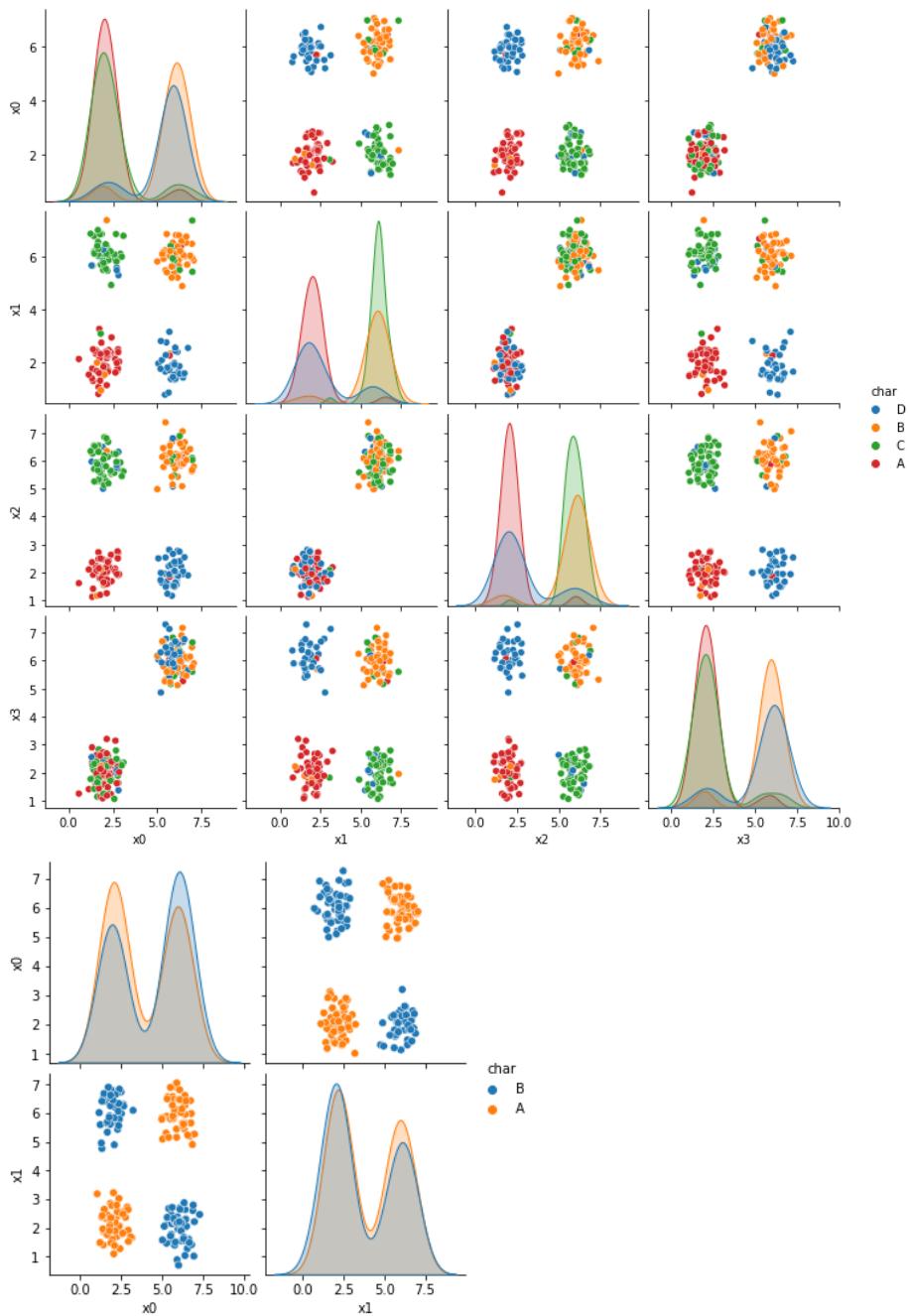
```
a6_data = 'https://raw.githubusercontent.com/rhodyprog4ds/06-naive-
bayes/main/data/dataset'
data_urls = [a6_data + str(i) + '.csv' for i in range(1,7)]

# read in only the columns with actual data
[sns.pairplot(data =pd.read_csv(url,usecols=lambda c: not('Unnamed' in c)),
hue='char') for url in data_urls]
```

```
[<seaborn.axisgrid.PairGrid at 0x7fc832b97090>,
<seaborn.axisgrid.PairGrid at 0x7fc82d0fce50>,
<seaborn.axisgrid.PairGrid at 0x7fc82c72d8d0>,
<seaborn.axisgrid.PairGrid at 0x7fc82b3c9f50>,
<seaborn.axisgrid.PairGrid at 0x7fc82ae62050>,
<seaborn.axisgrid.PairGrid at 0x7fc82aa59590>]
```







Now we're looking at dataset 6 more deeply.

```
df6= pd.read_csv(data_urls[-1],usecols=[1,2,3])
df6.head()
```

	x0	x1	char
0	6.14	2.10	B
1	2.22	2.39	A
2	2.27	5.44	B
3	1.03	3.19	A
4	2.25	1.71	A

This one Naive Bayes does poorly on, even though the classes are separable because each class is not defined by a single region. Since each class has two noncontinuous regions.

```
dt = DecisionTreeClassifier()
X_train, X_test, y_train, y_test = train_test_split(df6.values[:,2:],df6.values[:,2])
```

```
y_train
```

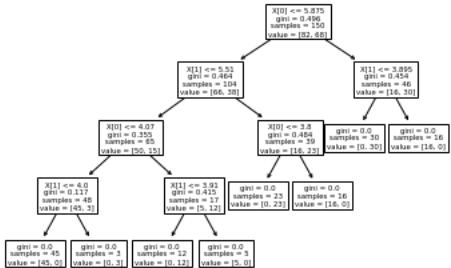
```
array(['A', 'A', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'B',
       'B', 'A', 'A', 'B', 'A', 'B', 'B', 'A', 'A', 'A', 'A', 'B', 'A',
       'A', 'A', 'B', 'A', 'B', 'B', 'A', 'A', 'B', 'A', 'B', 'B', 'A',
       'A', 'B', 'A', 'A', 'B', 'A', 'A', 'B', 'B', 'A', 'B', 'B', 'B',
       'A', 'B', 'A', 'A', 'B', 'A', 'A', 'B', 'B', 'A', 'B', 'B', 'A',
       'A', 'A', 'A', 'B', 'A', 'B', 'A', 'B', 'B', 'A', 'A', 'B', 'B',
       'B', 'A', 'B', 'A', 'B', 'B', 'A', 'A', 'B', 'A', 'A', 'B', 'B',
       'A', 'B', 'B', 'A', 'A', 'B', 'A', 'B', 'B', 'A', 'A', 'B', 'B',
       'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A',
       'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A',
       'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A',
       'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A', 'B', 'B', 'A'],
      dtype=object)
```

```
dt.fit(X_train,y_train)
```

```
DecisionTreeClassifier()
```

```
tree.plot_tree(dt)
```

```
[Text(221.20714285714286, 195.696, 'X[0] <= 5.875\n gini = 0.496\n samples = 150\n value = [82, 68]'),
 Text(155.44285714285715, 152.208, 'X[1] <= 5.51\n gini = 0.464\n samples = 104\n value = [66, 38]'),
 Text(95.65714285714286, 108.72, 'X[0] <= 4.07\n gini = 0.355\n samples = 65\n value = [50, 15]'),
 Text(47.82857142857143, 65.232, 'X[1] <= 4.0\n gini = 0.117\n samples = 48\n value = [45, 3]'),
 Text(23.914285714285715, 21.744, 'gini = 0.0\n samples = 45\n value = [45, 0]'),
 Text(71.74285714285715, 21.744, 'gini = 0.0\n samples = 3\n value = [0, 3]'),
 Text(143.4857142857143, 65.232, 'X[1] <= 3.91\n gini = 0.415\n samples = 17\n value = [5, 12]'),
 Text(119.57142857142857, 21.744, 'gini = 0.0\n samples = 12\n value = [0, 12]'),
 Text(167.4, 21.744, 'gini = 0.0\n samples = 5\n value = [5, 0]'),
 Text(215.22857142857143, 108.72, 'X[0] <= 3.8\n gini = 0.484\n samples = 39\n value = [16, 23]'),
 Text(191.31428571428572, 65.232, 'gini = 0.0\n samples = 23\n value = [0, 23]'),
 Text(239.14285714285714, 65.232, 'gini = 0.0\n samples = 16\n value = [16, 0]'),
 Text(286.9714285714286, 152.208, 'X[1] <= 3.895\n gini = 0.454\n samples = 46\n value = [16, 30]'),
 Text(263.0571428571429, 108.72, 'gini = 0.0\n samples = 30\n value = [0, 30]'),
 Text(310.8857142857143, 108.72, 'gini = 0.0\n samples = 16\n value = [16, 0]')]
```



```
print(tree.export_text(dt))
```

```
|--- feature_0 <= 5.88
|   |--- feature_1 <= 5.51
|   |   |--- feature_0 <= 4.07
|   |   |   |--- feature_1 <= 4.00
|   |   |   |   |--- class: A
|   |   |   |   |--- feature_1 >  4.00
|   |   |   |   |--- class: B
|   |   |--- feature_0 >  4.07
|   |   |   |--- feature_1 <= 3.91
|   |   |   |   |--- class: B
|   |   |   |   |--- feature_1 >  3.91
|   |   |   |   |--- class: A
|   |--- feature_1 >  5.51
|   |   |--- feature_0 <= 3.80
|   |   |   |--- class: B
|   |   |   |--- feature_0 >  3.80
|   |   |   |--- class: A
|--- feature_0 >  5.88
|   |--- feature_1 <= 3.89
|   |   |--- class: B
|   |--- feature_1 >  3.89
|   |   |--- class: A
```

```
dt.score(X_test,y_test)
```

```
1.0
```

```
dt.get_depth()
```

```
4
```

```
dt2 = DecisionTreeClassifier(max_depth=2)
```

```
dt2.fit(X_train, y_train,)
```

```
DecisionTreeClassifier(max_depth=2)
```

```
dt2.score(X_test,y_test)
```

```
0.72
```

```
print(tree.export_text(dt2))
```

```
|--- feature_0 <= 5.88
|   |--- feature_1 <= 5.51
|   |   |--- class: A
|   |   |--- feature_1 >  5.51
|   |   |   |--- class: B
|--- feature_0 >  5.88
|   |--- feature_1 <= 3.89
|   |   |--- class: B
|   |   |--- feature_1 >  3.89
|   |   |--- class: A
```

```
dt2.score(X_train,y_train)
```

```
0.7933333333333333
```

Class 20: Decision Trees and Cross Validation

1. Share your favorite beverage (or say hi) in the zoom chat
2. log onto prismia
3. Accept assignment 7

Assignment 7

Make a plan with a group:

- what methods do you need to use in part 1?
- try to outline with psuedocode what you'll do for part 2 & 3

Share any questions you have.

Followup:

1. assignment clarified to require 3 values for the parameter in part 2
2. more tips on finding data sets added to assignment text

Complexity of Decision Trees

```
# %load http://drsmr.co/310
import pandas as pd
import seaborn as sns
import numpy as np
from sklearn import tree
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import train_test_split
d6_url = 'https://raw.githubusercontent.com/rhodyprog4ds/06-naive-
bayes/main/data/dataset6.csv'
```

```
df6= pd.read_csv(d6_url,usecols=[1,2,3])
df6.head()
```

	x0	x1	char
0	6.14	2.10	B
1	2.22	2.39	A
2	2.27	5.44	B
3	1.03	3.19	A
4	2.25	1.71	A

```
X_train, X_test, y_train, y_test = train_test_split(df6.values[:, :2], df6.values[:, 2],
train_size=.8)
```

```
dt = tree.DecisionTreeClassifier(min_samples_leaf = 10)
dt.fit(X_train,y_train)
```

```
DecisionTreeClassifier(min_samples_leaf=10)
```

```
print(tree.export_text(dt))
```

```
--- feature_0 <= 5.51
|   --- feature_1 <= 4.00
|   |   --- feature_0 <= 2.88
|   |   |   class: A
|   |   --- feature_0 > 2.88
|   |   |   --- class: B
|   --- feature_1 > 4.00
|   |   --- feature_0 <= 2.63
|   |   |   class: B
|   |   --- feature_0 > 2.63
|   |   |   |   --- class: A
|   --- feature_0 > 5.51
|   |   --- feature_1 <= 3.89
|   |   |   class: B
|   |   --- feature_1 > 3.89
|   |   |   |   --- class: A
```

```
dt2 = tree.DecisionTreeClassifier(min_samples_leaf = 50)
dt2.fit(X_train,y_train)
```

```
DecisionTreeClassifier(min_samples_leaf=50)
```

Note

`df6.values` is a numpy array, which is a good datastructure for storing matrices of data. We can index into numpy arrays using [rows, columns]. Here, `df6.values[:, :2]` we take all the rows (:) and the columns up to, but not including index 2 for the features (X) :2 and use columns at index 2 for the target(y).

```
print(tree.export_text(dt2))
```

```
|--- feature_0 <= 5.51
|   |--- class: A
|--- feature_0 >  5.51
|   |--- class: B
```

```
dt2.score(X_test,y_test)
```

```
0.5
```

```
dt.score(X_test,y_test)
```

```
1.0
```

```
df6.shape
```

```
(200, 3)
```

Training, Test set size and Cross Validation

```
dt3 = tree.DecisionTreeClassifier()
dt3.fit(df6.values[:-1,:2],df6.values[:-1,2],)
```

```
DecisionTreeClassifier()
```

```
print(tree.export_text(dt3))
```

```
|--- feature_0 <= 5.88
|   |--- feature_1 <= 5.33
|   |   |--- feature_0 <= 4.07
|   |   |   |--- feature_1 <= 4.00
|   |   |   |   |--- class: A
|   |   |   |   |--- feature_1 >  4.00
|   |   |   |   |--- class: B
|   |   |--- feature_0 >  4.07
|   |   |   |--- feature_1 <= 3.91
|   |   |   |   |--- class: B
|   |   |   |   |--- feature_1 >  3.91
|   |   |   |   |--- class: A
|   |--- feature_1 >  5.33
|   |   |--- feature_0 <= 4.09
|   |   |   |--- class: B
|   |   |   |--- feature_0 >  4.09
|   |   |   |--- class: A
|--- feature_0 >  5.88
|   |--- feature_1 <= 3.89
|   |   |--- class: B
|   |--- feature_1 >  3.89
|   |   |--- class: A
```

```
dt4 = tree.DecisionTreeClassifier(max_depth=2)
cv_scores = cross_val_score(dt4,df6.values[:,2:],df6.values[:,2],cv=100 )
cv_scores
```

```
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/sklearn/model_selection/_split.py:672: UserWarning: The least populated
class in y has only 99 members, which is less than n_splits=100.
  % (min_groups, self.n_splits)), UserWarning)
```

```
array([1. , 1. , 0.5, 0.5, 1. , 0.5, 1. , 0.5, 0.5, 1. , 1. , 0.5, 0.5,
       1. , 1. , 0.5, 1. , 0.5, 1. , 1. , 0.5, 0.5, 1. , 0. , 1. ,
       1. , 1. , 0.5, 1. , 0.5, 0.5, 0.5, 1. , 0.5, 1. , 1. , 0.5,
       0.5, 1. , 0.5, 0.5, 0.5, 1. , 0.5, 0.5, 1. , 1. , 0.5, 1. , 1. ,
       1. , 1. , 0.5, 1. , 1. , 1. , 1. , 0. , 1. , 0.5, 0.5,
       1. , 0. , 1. , 0.5, 1. , 0.5, 1. , 1. , 1. , 1. , 0.5, 0.5,
       0.5, 1. , 1. , 1. , 0. , 1. , 1. , 1. , 1. , 1. , 0.5, 1. ,
```

```
np.mean(cv_scores)
```

```
0.755
```

Class 21: Regression

1. Log onto prismia
2. Share a topic you're most interested in applying data science to in the zoom chat

Regression Introduction

What is the difference in data that's well suited for regression vs classification

- [] regression is better for more features
- [] regression can work with categorical features
- [x] regression uses a continuous target variable

Explanation: the difference is that for regression we can use continuous target variables. Either classification or regression can work with high dimensional data (dimension refers to the number of features). Either can also work with a mixture of categorical or continuous valued features.

```
import numpy as np
import seaborn as sns
from sklearn import datasets, linear_model
from sklearn.metrics import mean_squared_error, r2_score
import pandas as pd
sns.set_theme(font_scale=2)
```

Today we will work with the [diabetes dataset](#) and we'll use the `sklearn.datasets` module to load it. When we load data this way, it gets loaded as a `numpy` array

```
diabetes_X, diabetes_y = datasets.load_diabetes(return_X_y = True)
```

Since it's not a `pandas.DataFrame` we don't have the `head` method, but we can index using square brackets.

```
diabetes_X[:5]
```

```
array([[ 0.03807591,  0.05068012,  0.06169621,  0.02187235, -0.0442235 ,
       -0.03482076, -0.04340085, -0.00259226,  0.01990842, -0.01764613],
      [-0.00188202, -0.04464164, -0.05147406, -0.02632783, -0.00844872,
       -0.01916334,  0.07441156, -0.03949338, -0.06832974, -0.09220405],
      [ 0.08529891,  0.05068012,  0.04445121, -0.00567061, -0.04559945,
       -0.03419447, -0.03235593, -0.00259226,  0.00286377, -0.02593034],
      [-0.08906294, -0.04464164, -0.01159501, -0.03665645,  0.01219057,
       0.02499059, -0.03603757,  0.03430886,  0.02269202, -0.00936191],
      [ 0.00538306, -0.04464164, -0.03638469,  0.02187235,  0.00393485,
       0.01559614,  0.00814208, -0.00259226, -0.03199144, -0.04664087]])
```

and the same for the labels.

```
diabetes_y[:5]
```

```
array([151.,  75., 141., 206., 135.])
```

Since this data is not in any sorted order, we can split into test and train using indexing instead of the `test_train_split` function. This way we all get the same split, without setting the `random_state` parameter. In most cases, it's best to use the function, but it's good to know different ways to do things.

We'll start by using only one feature, the one in column 8.

```
diabetes_X_train = diabetes_X[:-20,8]
diabetes_X_test = diabetes_X[-20:,8]
diabetes_y_train = diabetes_y[:-20]
diabetes_y_test = diabetes_y[-20:]
```

Now we can instantiate the object

```
regr = linear_model.LinearRegression()
```

All `sklearn` estimators have the same methods, and all take any specialized parameters in the constructor. We've used the default values here, but this is an important design feature of scikit learn, because it makes their pipeline infrastructure and functions for cross validation work.

```
regr.fit(diabetes_X_train,diabetes_y_train)
```

```
-----  
ValueError                                                 Traceback (most recent call last)  
<ipython-input-7-7563ebc7151a> in <module>  
----> 1 regr.fit(diabetes_X_train,diabetes_y_train)  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/sklearn/linear_model/_base.py in fit(self, X, y, sample_weight)  
  504         n_jobs_ = self.n_jobs  
  505         X, y = self._validate_data(X, y, accept_sparse=['csr', 'csc', 'coo'],  
--> 506                                         y_numeric=True, multi_output=True)  
  507  
  508         if sample_weight is not None:  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-packages/sklearn/base.py in  
_validate_data(self, X, y, reset, validate_separately, **check_params)  
  430             y = check_array(y, **check_y_params)  
  431         else:  
--> 432             X, y = check_X_y(X, y, **check_params)  
  433         out = X, y  
  434  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/sklearn/utils/validation.py in inner_f(*args, **kwargs)  
  70             FutureWarning)  
  71         kwargs.update({k: arg for k, arg in zip(sig.parameters, args)})  
--> 72     return f(**kwargs)  
  73     return inner_f  
  74  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/sklearn/utils/validation.py in check_X_y(X, y, accept_sparse,  
accept_large_sparse, dtype, order, copy, force_all_finite, ensure_2d, allow_nd,  
multi_output, ensure_min_samples, ensure_min_features, y_numeric, estimator)  
  800             ensure_min_samples=ensure_min_samples,  
  801             ensure_min_features=ensure_min_features,  
--> 802             estimator=estimator)  
  803         if multi_output:  
  804             y = check_array(y, accept_sparse='csr', force_all_finite=True,  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/sklearn/utils/validation.py in inner_f(*args, **kwargs)  
  70             FutureWarning)  
  71         kwargs.update({k: arg for k, arg in zip(sig.parameters, args)})  
--> 72     return f(**kwargs)  
  73     return inner_f  
  74  
  
/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/sklearn/utils/validation.py in check_array(array, accept_sparse,  
accept_large_sparse, dtype, order, copy, force_all_finite, ensure_2d, allow_nd,  
ensure_min_samples, ensure_min_features, estimator)  
  621             "Reshape your data either using array.reshape(-1, 1) if "  
  622             "your data has a single feature or array.reshape(1, -1) "  
--> 623             "if it contains a single sample.".format(array))  
  624  
  625     # in the future np.flexible dtypes will be handled like object dtypes  
  
ValueError: Expected 2D array, got 1D array instead:  
array=[ 0.01990842 -0.06832974  0.00286377  0.02269202 -0.03199144 -0.04118039  
-0.06291295 -0.03581673 -0.01495648  0.06773633 -0.06291295 -0.09643322  
-0.03075121  0.03839325 -0.03199144  0.03605579  0.05228   0.02736771  
-0.01811827 -0.00894402 -0.01190068 -0.07212845 -0.0611766  0.13359898  
-0.02595242  0.01919903 -0.0425721 -0.01599827 -0.00060925  0.0594238  
-0.02712865 -0.03712835  0.00027149 -0.01811827 -0.0594727  0.02131085  
 0.054724  0.01703713  0.07142403  0.01919903  0.01255315 -0.04986847  
-0.009911896 -0.0425721  0.03243323 -0.01495648 -0.0079794 -0.01811827  
-0.0594727  0.03546194 -0.02139368 -0.00239668 -0.01811827 -0.03324879  
-0.04118039 -0.06648815  0.03365681 -0.05615757 -0.00060925  0.03839325  
-0.05140054  0.0366458 -0.08238148 -0.03075121 -0.01919705  0.0423449  
 0.00027149 -0.01190068 -0.01290794 -0.05140054 -0.06291295  0.02671426  
 0.08449528 -0.00514531  0.00371174  0.00620932 -0.03075121 -0.07408887  
 0.04289569 -0.06832974  0.02539313 -0.00608025 -0.07020931 -0.01599827  
-0.06832974 -0.04118039 -0.09643322  0.04289569 -0.03452372 -0.02479119  
-0.07212845  0.03723201  0.01255315 -0.03324879 -0.04118039  0.0011438  
 0.06078775  0.00286377 -0.00608025  0.0117839  0.03723201 -0.03075121  
-0.00514531 -0.01090444 -0.0611766 -0.01090444 -0.0594727 -0.05295879  
 0.04560081 -0.02595242 -0.12609739 -0.02364456 -0.02364456  0.04666077  
 0.09924023  0.08379677  0.02200405  0.07912108  0.02061233  0.02736771  
-0.03980959  0.01990842  0.03243323  0.01556684 -0.01599827  0.08058546  
-0.09393565  0.00455189 -0.03075121  0.03896837  0.02269202 -0.04687948  
-0.01290794 -0.07212845 -0.02712865  0.05078151 -0.06648815  0.01703713  
 0.02801651  0.054724 -0.03324879  0.04506617  0.04560081 -0.04118039  
 0.00620932 -0.01599827  0.01482271  0.02605609 -0.03075121  0.06123791  
-0.01919705 -0.00894402  0.02671426 -0.03452372  0.00864028  0.03781448  
-0.02139368  0.01022564 -0.03075121  0.0423449 -0.03581673  0.07573759  
-0.03075121 -0.02364456  0.01776348 -0.05615757 -0.07814091  0.07763279  
 0.06898221  0.13359898 -0.09643322 -0.05140054  0.07341008 -0.01811827
```

```

0.01703713 -0.02712865 -0.00330371 0.07222365 -0.04836172 0.01407245
0.01630495 -0.02595242 0.01556684 0.01556684 0.03723201 0.06731722
0.04718617 -0.08682899 -0.00149859 -0.02028875 0.02337484 0.01022564
-0.02952762 0.03953988 0.00200784 0.00783714 -0.03324879 0.01331597
-0.07408887 0.08989869 -0.02832024 -0.05295879 0.08449528 0.02539313
0.00200784 0.06345592 0.00943641 0.04613233 -0.00149859 -0.01190068
-0.03324879 -0.02251217 -0.00060925 -0.02251217 -0.03980959 0.0366458
0.04560081 0.03365681 0.00702686 0.0011438 -0.10164355 0.0011438
0.0011438 -0.01811827 -0.03581673 0.06301662 -0.03980959 -0.03845911
-0.02712865 -0.02364456 0.05027649 0.00943641 0.02405258 -0.02595242
0.06432823 -0.00514531 0.04067226 -0.04836172 0.0110081 0.06604821
0.03365681 -0.02952762 -0.07814091 -0.0439854 -0.0439854 -0.04986847
0.04506617 -0.01090444 0.06389312 0.0702113 0.07573759 0.08449528
-0.00421986 0.03486419 0.12005338 -0.00330371 0.02801651 -0.00514531
-0.03712835 -0.0439854 -0.08023654 -0.01290794 0.03486419 -0.07212845
-0.0594727 0.01482271 -0.08682899 0.02337484 0.02930041 -0.01190068
0.00864028 -0.01811827 -0.01811827 0.10635428 0.01556684 0.0110081
0.05757286 -0.02139368 0.0011438 -0.0159827 0.01630495 0.02405258
-0.00894402 -0.06832974 0.00286377 0.05988072 -0.01919705 0.03243323
-0.07020931 0.06123791 0.02993565 0.05803913 -0.06291295 -0.05454415
-0.07408887 0.01630495 -0.05615757 -0.06468302 -0.03075121 -0.02595242
0.04289569 -0.02139368 0.03119299 0.07380215 -0.02251217 0.01776348
-0.03980959 0.02337484 -0.0611766 0.02671426 -0.01811827 0.01255315
-0.03845911 0.05520504 -0.01190068 -0.01090444 0.07496834 0.02472532
0.00943641 0.02866072 0.03304707 0.09864637 0.13339573 0.12901941
0.06604821 0.08449528 0.02993565 -0.00060925 -0.00149859 -0.0170521
-0.02364456 -0.04118039 0.05710419 0.02930041 -0.06291295 -0.05140054
0.08094791 0.00538437 -0.02251217 0.04344317 -0.03581673 -0.02479119
-0.02364456 -0.05780007 0.00702686 -0.00060925 0.06257518 0.00371174
-0.01090444 -0.06111766 0.0702113 -0.05780007 -0.02028875 0.13237265
0.07419254 -0.03324879 -0.02832024 0.04976866 -0.08238148 0.01482271
0.01331597 -0.07212845 0.00371174 0.01919903 0.01482271 0.03119299
0.07102158 0.03365681 0.08553312 -0.00514531 -0.0611766 0.11934399
-0.01090444 -0.0425721 -0.00060925 0.02801651 0.04613233 0.00943641
0.02269202 -0.04687948 -0.05140054 -0.01290794 0.10413761 -0.01811827
0.02405258 -0.02595242 -0.03075121 -0.10436482 0.02671426 -0.04118039
0.10329226 -0.08913686 -0.04118039 -0.03980959 0.13008061 0.03181522
-0.09643322 0.03605579 0.04506617 0.03953988 -0.03452372 -0.0425721
0.01556684 0.06168585 -0.02712865 0.0366458 -0.05615757 -0.04836172
0.05803913 0.03056649 0.06604821 -0.05140054 0.00620932 -0.03581673
0.054724 0.01482271 -0.02952762 -0.04687948 -0.02139368 -0.07408887
-0.03324879 0.04976866].

```

Reshape your data either using `array.reshape(-1, 1)` if your data has a single feature
or `array.reshape(1, -1)` if it contains a single sample.

we get an error because the training data (`diabetes_X_train`) is only one dimensional. Scikit learn's `fit` methods require a 2d-array even if fitting with only one feature.

```
diabetes_X_train.shape
```

```
(422, )
```

we can use `np.newaxis` to fix this problem.

```
diabetes_X_train = diabetes_X[:-20,np.newaxis,8]
diabetes_X_test = diabetes_X[-20:,:,np.newaxis,8]
```

and check the shape

```
diabetes_X_train.shape
```

```
(422, 1)
```

It's the same shape, in a mathematical sense, but in terms of the underlying data structure this one has a second dimension (it's just only 1 long) in that dimension. This is tricky, but an important thing to know how to fix.

Now we can fit our model

```
regr.fit(diabetes_X_train,diabetes_y_train)
```

```
LinearRegression()
```

and examine the coefficient

```
regr.coef_
```

```
array([900.39171612])
```

and make predictions with the model

```
diabetes_y_pred = regr.predict(diabetes_X_test)
```

we can evaluate with `mean_squared_error` (see below for more detail), which is as it's named. It computes the error in each prediction ($y_{\text{pred}} - y_{\text{true}}$), squares it, then averages them all together.

```
mean_squared_error(diabetes_y_test,diabetes_y_pred)
```

```
2923.342534244987
```

We can also use the r^2 score, which is more normalized, it's best value is 1.

```
r2_score(diabetes_y_test,diabetes_y_pred)
```

```
0.3948984231023219
```

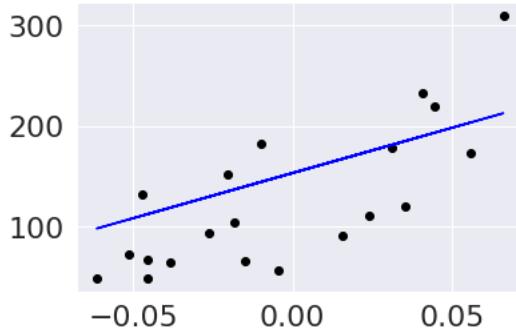
To get an even better idea, we can plot. We'll use `matplotlib` to plot, since our data is not in a DataFrame. `seaborn` and `pandas` use `matplotlib` under the hood, so it will have a lot of familiar methods, but it takes a lot more work to build complicated figures with `matplotlib`. The alias `plt` is used by convention for the `pyplot` module of `matplotlib`.

```
import matplotlib.pyplot as plt
```

Then we can scatter plot the true data and make a line of the predictions in blue.

```
plt.scatter(diabetes_X_test,diabetes_y_test, color='black')
plt.plot(diabetes_X_test,diabetes_y_pred, color='blue')
```

```
[<matplotlib.lines.Line2D at 0x7ff5625c5210>]
```



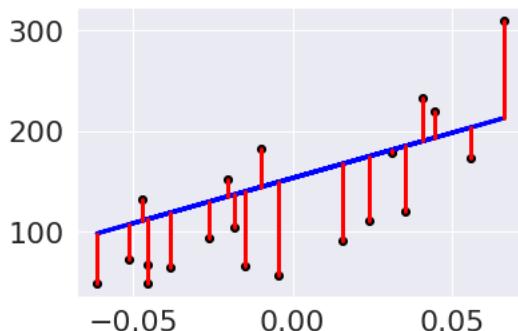
We can modify this plot to highlight the errors, too.

```
plt.scatter(diabetes_X_test, diabetes_y_test, color='black')
plt.plot(diabetes_X_test, diabetes_y_pred, color='blue', linewidth=3)

[plt.plot([x,x],[yp,yt], color='red', linewidth=3)
    for x, yp, yt in zip(diabetes_X_test,
    diabetes_y_pred,diabetes_y_test)];
```

💡 Tip

the semicolon here prevents it from printing out the axis information for each element of the list comprehension



Try it yourself

Fit another regression model to this data that uses all of the features instead of only one

```
diabetes_X_train2 = diabetes_X[:-20]
diabetes_X_test2 = diabetes_X[-20:]
diabetes_y_train2 = diabetes_y[:-20]
diabetes_y_test2 = diabetes_y[-20:]

regr2 = linear_model.LinearRegression()
regr2.fit(diabetes_X_train2,diabetes_y_train2)
diabetes_y_pred2 = regr2.predict(diabetes_X_test2)
r2_score(diabetes_y_test2,diabetes_y_pred2)
```

0.5850753022690575

This does better on r^2 , let's also check the mse.

```
mean_squared_error(diabetes_y_test2,diabetes_y_pred2)
```

2004.5676026898207

More practice

1. try out `regr2.score` what does scikite learn use for the score for `linear_model.LinearRegression` models?
2. repeat the above using `test_train_split`, can you make the test size 20 samples again?
3. try another single feature model using a different feature than we did above. Is your new feature a better or worse predictor? Make plots to compare and try to build intuition for the performance metrics geometrically.

Questions After class

What is considered a good r^2 value?

This will depend on context in some sense. You can read about it in the `r2_score` docs and the corresponding [wikipedia page](#)

Here's a quick visual of what different r^2 score values look like. I simulated data by randomly picking 10 points, then made the "predicted" y values by picking a slope of 3 and computing $3*x$. Then I simulated various levels of noise, by sampling noise and multiplying the same noise vector by different scales and adding all of those to a data frame with the column name the r score for if that column of target values was the truth.

Then I added some columns of y values that were with different slopes and different functions of x . These all have the small amount of noise.

I used the pandas `melt` method to restructure the DataFrame so that I could use `FacetGrid` with `col` and `col_wrap` to show all of the results.

```

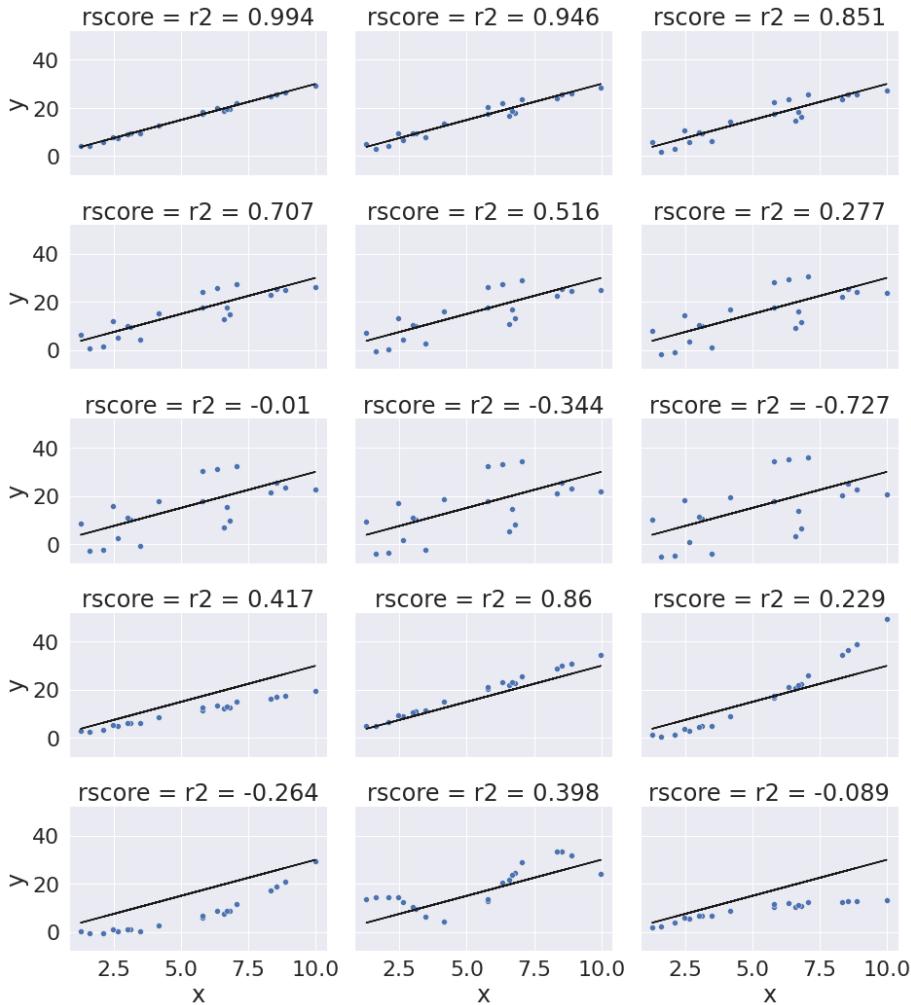
x = 10*np.random.random(20)
y_pred = 3*x
ex_df = pd.DataFrame(data = x,columns = ['x'])
ex_df['y_pred'] = y_pred
n_levels = range(1,18,2)
noise = (np.random.random(20)-.5)*2
for n in n_levels:
    y_true = y_pred + n* noise
    ex_df['r2 = '+ str(np.round(r2_score(y_pred,y_true),3))] = y_true

f_x_list = [2*x,.5*x**2, .03*x**3, 10*np.sin(x)+x*3,3*np.log(x**2)]
for fx in f_x_list:
    y_true = fx + noise
    ex_df['r2 = '+ str(np.round(r2_score(y_pred,y_true),3))] = y_true

xy_df = ex_df.melt(id_vars=['x','y_pred'],var_name='rscore',value_name='y')
# sns.lmplot(x='x',y='y', data = xy_df,col='rscore',col_wrap=3, )
g = sns.FacetGrid(data = xy_df,col='rscore',col_wrap=3,aspect=1.5,height=3)
g.map(plt.plot, 'x','y_pred',color='k')
g.map(sns.scatterplot, "x", "y",)

```

<seaborn.axisgrid.FacetGrid at 0x7ff56b27e750>



The `r2_score` alone doesn't tell the whole picture. We'll also often want to analyze the errors in greater detail. We'll look at how to do that to investigate how the model really fits more later this week.

Can some of the methods applied in regression can be used in the same format in classification?

the `sklearn` estimator objects all have the same methods, you can see details on the API on the [sklearn developer page](#).

Why couldn't we use `train_test_split` for this model?

We absolutely can. Splitting this way was to show *another* way to split the data, and point out under which conditions this way is ok.

I am still confused about machine learning

Try reading the notes from the [introduction class](#).

The scikit-learn [choose an estimator](#) graph can also be helpful.

What is the difference between test and train data?

We split data, sample-wise, into two pieces, in order to evaluate what we are doing.

There is nothing inherently different between test data and train data. In fact, by applying the model we learned with the training data to predict on the test data we are assuming that they are very similar.

We split our data so that we can see how well we expect our trained model to work on new data, since the goal is to send our learned model into the world and use it for something else. For example, a model fit on today's data, about diabetes, might be used to predict which patients are going to have the most severe disease in a year based on their stats today and then enroll them in a program to help them learn to better manage their diabetes.

The test set is meant to simulate future data that we've never seen before, but it's usually data we have seen, but our learning algorithm did not use to fit the model.

What does cross validation mean?

Cross validation will come up again later, but basically cross validation is a more elaborate evaluation. It repeatedly splits the data into test and train sets, fits the model, scores the model, and then returns all fo the results.

What does `mean_squared_error` do?

It is a popular performance metric for regression. To understand it, let's look back at what regression is doing. Regression is trying to predict a continuous valued quantity, so unlike in classification where we can just check if it matches or not, to compute performance, it makes sense consider the size of the error, because it's unlikely to get *exactly* the right value for any prediction.

So, we might be tempted to compute error like this:

```
diabetes_y_test2-diabetes_y_pred2

array([ 35.38153092, -64.43979328, -61.88665147,  40.46462721,
       -44.80054784, -64.06954875,  50.87762239,  -6.47935157,
      65.9398948 , -58.30503555, -45.36632793,  10.80168716,
     -83.25046751, -56.3332925 , -4.54458691, -16.03798088,
      1.42860298,   8.43395013,   8.9653683 ,  4.39664326])
```

this, however is a whole vector of quantities, so we would need to take the average. Taking the average here though, will be very small, because some are positive and some are negative and they will cancel one another out.

```
np.mean(diabetes_y_test2-diabetes_y_pred2)

-13.941182850978748
```

In this case, it turns out to even be negative.

Instead, we can square each error, to make them all positive. We *could* also take the absolute value, but in a lot of calculations, we want to take the derivative, so squaring instead of absolute value became more popular.

```
(diabetes_y_test2-diabetes_y_pred2)**2

array([1.25185273e+03, 4.15248696e+03, 3.82995763e+03, 1.63738606e+03,
       2.00708909e+03, 4.10490708e+03, 2.58853246e+03, 4.19819967e+01,
      4.34806973e+03, 3.39947717e+03, 2.05810371e+03, 1.16676446e+02,
     6.93064034e+03, 3.17343984e+03, 2.06532702e+01, 2.57216831e+02,
      2.04090647e+00, 7.11315149e+01, 8.03778287e+01, 1.93304719e+01])
```

now we can take the mean of these to get a total average error.

```
mse_manual = np.mean((diabetes_y_test2-diabetes_y_pred2)**2)
mse_manual
```

```
2004.5676026898207
```

we can confirm this gets the same answer as the sklearn function

```
mse_sk = mean_squared_error(diabetes_y_test2,diabetes_y_pred2)
mse_sk
```

```
2004.5676026898207
```

We can even assert that they're the same, they're always going to be exactly the same

```
assert mse_manual == mse_sk
```

Class 22: More Regression, More Evaluation and LASSO

1. log prismia chat
2. say hello in the zoom chat

```
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
import pandas as pd
from sklearn import datasets, linear_model
from sklearn.metrics import mean_squared_error, r2_score
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import train_test_split

# Load the diabetes dataset
diabetes_X, diabetes_y = datasets.load_diabetes(return_X_y=True)
```

Questions after class Monday

Some good questions were asked on the form Monday. Check the [notes](#) for insight on the `r2_score` and `mean_square_error`.

Review of Test Train splits

```
X_train,X_test, y_train,y_test = train_test_split(diabetes_X, diabetes_y ,
test_size=20,random_state=0)
```

What metric does the `score` method of `LinearRegression` use?

```
regr = linear_model.LinearRegression()
regr.fit(X_train, y_train)
```

```
LinearRegression()
```

```
regr.score(X_test,y_test)
```

```
0.5195208400616668
```

```
y_pred = regr.predict(X_test)
r2_score(y_test,y_pred)
```

```
0.5195208400616668
```

```
mean_squared_error(y_test,y_pred)
```

```
2850.3176917525775
```

Digging in deeper to the Linear Regression model

Linear regression fitting involves learning the coefficients

```
regr.coef_
```

```
array([-32.3074285, -257.44432972, 513.31945939, 338.46656647,
       -766.86983748, 455.85416891, 92.55795582, 184.75163454,
       734.92318647, 82.7231425])
```

and an intercept

```
regr.intercept_
```

```
152.39189054201842
```

The linear regression model is

$$y = wx + b$$

where it stores `w` in the `coef_` attribute and `b` as the `intercept_`

We can check how this works by multiplying one sample by the coefficients, to get a vector

```
X_test[0]*regr.coef_
```

```
array([-0.64334474, -13.0473092, 53.80033982, 23.71721361,
       27.58260658, -12.16168908, -2.31326921, -0.47892464,
       2.72784249, 3.33733048])
```

then taking the sum and adding the intercept

```
np.sum(X_test[0]*regr.coef_) + regr.intercept_
```

```
234.9126866563482
```

and then comparing to the predictions.

```
y_pred[0]
```

```
234.91268665634823
```

These are not exactly the same due to float rounding errors, but they're very close.

We can also check for the whole test set

```
errors = np.sum(X_test*regr.coef_,axis=1) + regr.intercept_ - y_pred
```

```
array([-2.84217094e-14, 0.00000000e+00, 0.00000000e+00, 0.00000000e+00,
       0.00000000e+00, 0.00000000e+00, 0.00000000e+00, 0.00000000e+00,
       0.00000000e+00, 0.00000000e+00, 0.00000000e+00, 0.00000000e+00,
       1.42108547e-14, 0.00000000e+00, 0.00000000e+00, -1.42108547e-14,
       0.00000000e+00, -1.42108547e-14, -2.84217094e-14, 0.00000000e+00])
```

and confirm these are very small

```
np.max(errors)
```

Tip

by default `np.sum` sums across both axes so it gets a single value. we want to only sum on 1 so that we get t

```
1.4210854715202004e-14
```

It's doing linear regression as we expected.

Changing the complexity in Linear regression

It can be important to also know what the model is really doing and see how different features are used or not. Maybe, for example all of the features are expensive to measure, but for testing we measured a lot of them. We might want to simultaneously learn which features we actually need *and* the linear model.

LASSO can do that for us it's objective is like linear regression, but it adds an extra term. This term forces some of the learned coefficients to be 0. Multiplying by 0 gives 0, so that's like throwing away that feature. The term is called the $1 - norm$, the details of the math are not important here, just the idea that it can reduce the number of features used as well.

$$\|y - wx\|_2^2 + \alpha \|w\|_1$$

```
lasso = linear_model.Lasso()  
lasso.fit(X_train, y_train)  
lasso.score(X_test,y_test)
```

```
0.4224899448032938
```

It uses many fewer

```
lasso.coef_
```

```
array([ 0.        , -0.        , 358.27498703,  9.7141813 ,  
       0.        , 0.        , -0.        ,  0.        ,  
      309.50796119,  0.        ])
```

It has a parameter alpha that must be > 0 that we can use to control how many features it uses. When `alpha = 0` it's the same as linear regression, but the regular linear regression estimator uses a more numerically stable algorithm for the `fit` method

```
lasso2 = linear_model.Lasso(alpha=.5)  
lasso2.fit(X_train, y_train)  
lasso2.score(X_test,y_test)
```

```
0.5272242529276977
```

We see that fewer are 0 now.

```
lasso2.coef_
```

```
array([ 0.        , -0.        , 466.09039819, 140.20195776,  
       -0.        , -0.        , -61.96474668,  0.        ,  
      405.95829094,  0.        ])
```

```
regr.score(X_train, y_train)
```

```
0.5170933599105026
```

```
lasso2.score(X_train, y_train)
```

```
0.4516267981295532
```

```
lasso_cv = cross_val_score(lasso2, diabetes_X, diabetes_y)  
lasso_cv
```

```
array([0.3712605 , 0.45675648, 0.45066569, 0.42134051, 0.47736463])
```

```
regr_cv = cross_val_score(regr,diabetes_X, diabetes_y )  
regr_cv
```

```
array([0.42955643, 0.52259828, 0.4826784 , 0.42650827, 0.55024923])
```

```
np.mean(lasso_cv), np.mean(regr_cv)
```

```
(0.43547756049731523, 0.48231812211149394)
```

```
cross_val_score(regr,diabetes_X, diabetes_y ,cv=10)
```

```
array([0.55614411, 0.23056092, 0.35357777, 0.62190498, 0.26587602,  
0.61819338, 0.41815916, 0.43515232, 0.43436983, 0.68568514])
```

We can do cross validation for all three of these models and look at both the mean and the standard deviation. The mean tells us how well on average each model does on different subsets of the data. The standard deviation is a measure of *spread* of the scores. For intuition, another measure of spread is `max(cv_scores) - min(cv_scores)`.

```
regr_cv = cross_val_score(regr,diabetes_X, diabetes_y ,cv=10)  
np.mean(regr_cv), np.std(regr_cv)
```

```
(0.461962361958337, 0.14698789185375885)
```

```
lasso_cv = cross_val_score(lasso,diabetes_X, diabetes_y ,cv=10)  
np.mean(lasso_cv), np.std(lasso_cv)
```

```
(0.3211351084864853, 0.09122225780232662)
```

```
lasso2_cv = cross_val_score(lasso2,diabetes_X, diabetes_y ,cv=10)  
np.mean(lasso2_cv), np.std(lasso2_cv)
```

```
(0.41808972445133297, 0.11927596370400496)
```

💡 Tip

To keep things concise, we can put two values on one line. This creates a `tuple` and prints out both values since it's on the last line of the cell.

Questions after class

What does the `cv` parameter in `cross_val_score` do?

First, let's look at the help.

```
help(cross_val_score)
```

```
Help on function cross_val_score in module sklearn.model_selection._validation:

cross_val_score(estimator, X, y=None, *, groups=None, scoring=None, cv=None,
n_jobs=None, verbose=0, fit_params=None, pre_dispatch='2*n_jobs', error_score=np.nan)
    Evaluate a score by cross-validation

    Read more in the :ref:`User Guide <cross_validation>`.

Parameters
-----
estimator : estimator object implementing 'fit'
    The object to use to fit the data.

X : array-like of shape (n_samples, n_features)
    The data to fit. Can be for example a list, or an array.

y : array-like of shape (n_samples,) or (n_samples, n_outputs),
default=None
    The target variable to try to predict in the case of
    supervised learning.

groups : array-like of shape (n_samples,), default=None
    Group labels for the samples used while splitting the dataset into
    train/test set. Only used in conjunction with a "Group" :term:`cv`
    instance (e.g., :class:`GroupKFold`).

scoring : str or callable, default=None
    A str (see model evaluation documentation) or
    a scorer callable object / function with signature
    ``scorer(estimator, X, y)`` which should return only
    a single value.

    Similar to :func:`cross_validate`
    but only a single metric is permitted.

    If None, the estimator's default scorer (if available) is used.

cv : int, cross-validation generator or an iterable, default=None
    Determines the cross-validation splitting strategy.
    Possible inputs for cv are:

    - None, to use the default 5-fold cross validation,
    - int, to specify the number of folds in a `(Stratified)KFold`,
    - :term:`CV splitter`,
    - An iterable yielding (train, test) splits as arrays of indices.

    For int/None inputs, if the estimator is a classifier and ``y`` is
    either binary or multiclass, :class:`StratifiedKFold` is used. In all
    other cases, :class:`KFold` is used.

    Refer :ref:`User Guide <cross_validation>` for the various
    cross-validation strategies that can be used here.

.. versionchanged:: 0.22
    ``cv`` default value if None changed from 3-fold to 5-fold.

n_jobs : int, default=None
    The number of CPUs to use to do the computation.
    ``None`` means 1 unless in a :obj:`joblib.parallel_backend` context.
    ``-1`` means using all processors. See :term:`Glossary <n_jobs>`
    for more details.

verbose : int, default=0
    The verbosity level.

fit_params : dict, default=None
    Parameters to pass to the fit method of the estimator.

pre_dispatch : int or str, default='2*n_jobs'
    Controls the number of jobs that get dispatched during parallel
    execution. Reducing this number can be useful to avoid an
    explosion of memory consumption when more jobs get dispatched
    than CPUs can process. This parameter can be:

    - None, in which case all the jobs are immediately
        created and spawned. Use this for lightweight and
        fast-running jobs, to avoid delays due to on-demand
        spawning of the jobs

    - An int, giving the exact number of total jobs that are
        spawned

    - A str, giving an expression as a function of n_jobs,
        as in '2*n_jobs'

error_score : 'raise' or numeric, default=np.nan
    Value to assign to the score if an error occurs in estimator fitting.
```

```

If set to 'raise', the error is raised.
If a numeric value is given, FitFailedWarning is raised. This parameter
does not affect the refit step, which will always raise the error.

.. versionadded:: 0.20

Returns
-----
scores : array of float, shape=(len(list(cv)),)
    Array of scores of the estimator for each run of the cross validation.

Examples
-----
>>> from sklearn import datasets, linear_model
>>> from sklearn.model_selection import cross_val_score
>>> diabetes = datasets.load_diabetes()
>>> X = diabetes.data[:150]
>>> y = diabetes.target[:150]
>>> lasso = linear_model.Lasso()
>>> print(cross_val_score(lasso, X, y, cv=3))
[0.33150734 0.08022311 0.03531764]

See Also
-----
:func:`sklearn.model_selection.cross_validate`:
    To run cross-validation on multiple metrics and also to return
    train scores, fit times and score times.

:func:`sklearn.model_selection.cross_val_predict`:
    Get predictions from each split of cross-validation for diagnostic
    purposes.

:func:`sklearn.metrics.make_scorer`:
    Make a scorer from a performance metric or loss function.

```

It does different things, depending on what type of value we pass it. We passed it an `int (10)`, so what it did was split the data into 10 groups and then trains on 9/10 of those parts and tests on the last one. Then it iterates over those folds.

For classification, it can take into consideration the target value (classes) and an additionally specified group in the data. A good visualization of what it does is shown in the [sklearn docs](#).

It uses StratifiedKFold for classification, but since we're using regression it will use `KFold.test_train_split` uses `ShuffleSplit` by default, let's load that too to see what it does.

⚠ Warning

The key in the following is to get the concepts not all of the details in how I evaluate and visualize. I could have made figures separately to explain the concept, but I like to show that Python is self contained.

```

from sklearn.model_selection import KFold, ShuffleSplit

kf = KFold(n_splits = 10)

```

When we use the `split` method it gives us a generator.

```

kf.split(diabetes_X, diabetes_y)

<generator object _BaseKFold.split at 0x7f332e706ad0>

```

We can use this in a loop to get the list of indices that will be used to get the test and train data for each fold. To visualize what this is doing, see below.

```

N_samples = len(diabetes_y)
kf_tt_df = pd.DataFrame(index=list(range(N_samples)))
i = 1
for train_idx, test_idx in kf.split(diabetes_X, diabetes_y):
    kf_tt_df['split ' + str(i)] = ['unused']*N_samples
    kf_tt_df['split ' + str(i)][train_idx] = 'Train'
    kf_tt_df['split ' + str(i)][test_idx] = 'Test'
    i +=1

```

We can count how many times 'Test' and 'Train' appear

How would you use those indices to get a out actual test and train data?

```
count_test = lambda part: len([v for v in part if v=='Test'])
count_train = lambda part: len([v for v in part if v=='Train'])
```

When we apply this along `axis=1` we to check that each sample is used in exactly 1 test set how many times each sample is used

```
sum(kf_tt_df.apply(count_test, axis = 1) == 1)
```

442

and exactly 9 training sets

```
sum(kf_tt_df.apply(count_test, axis = 1) == 9)
```

0

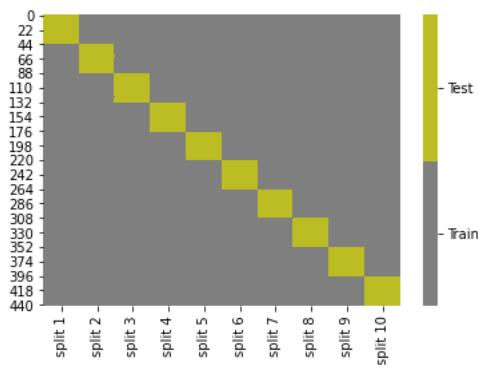
the describe helps ensure that all of the values are exa

We can also visualize:

```
cmap = sns.color_palette("tab10", 10)
g = sns.heatmap(kf_tt_df.replace({'Test':1, 'Train':0}), cmap=cmap[7:9], cbar_kws=
{'ticks':[.25,.75]}, linewidths=0,
linecolor='gray')
colorbar = g.collections[0].colorbar
colorbar.set_ticklabels(['Train', 'Test'])
```

Tip

`sns.heatmap` doesn't work on strings, so we can replace them for the plotting



Note that unlike `test_train_split` this does not always randomize and shuffle the data before splitting.

If we apply those `lambda` functions along `axis=0`, we can see the size of each test set

```
kf_tt_df.apply(count_test, axis = 0)
```

```
split 1    45
split 2    45
split 3    44
split 4    44
split 5    44
split 6    44
split 7    44
split 8    44
split 9    44
split 10   44
dtype: int64
```

and training set:

```
kf_tt_df.apply(count_train, axis = 0)
```

```

split 1    397
split 2    397
split 3    398
split 4    398
split 5    398
split 6    398
split 7    398
split 8    398
split 9    398
split 10   398
dtype: int64

```

We can verify that these splits are the same size as what `test_train_split` does using the right settings. 10-fold splits the data into 10 parts and tests on 1, so that makes a test size of $1/10 = .1$, so we can use the `train_test_split` and check the length.

```

X_train2,X_test2, y_train2,y_test2 = train_test_split(diabetes_X, diabetes_y ,
                                                    test_size=.1,random_state=0)

[len(split) for split in [X_train2,X_test2,]]

```

Under the hood `train_test_split` uses `ShuffleSplit`. We can do a similar experiment as above to see what `ShuffleSplit` does.

```

skf = ShuffleSplit(10)
N_samples = len(diabetes_y)
ss_tt_df = pd.DataFrame(index=list(range(N_samples)))
i = 1
for train_idx, test_idx in skf.split(diabetes_X, diabetes_y):
    ss_tt_df['split ' + str(i)] = ['unused']*N_samples
    ss_tt_df['split ' + str(i)][train_idx] = 'Train'
    ss_tt_df['split ' + str(i)][test_idx] = 'Test'
    i +=1

ss_tt_df

```

	split 1	split 2	split 3	split 4	split 5	split 6	split 7	split 8	split 9	split 10
0	Train	Test	Train							
1	Train									
2	Train									
3	Test	Train	Test	Train	Test	Train	Train	Train	Test	Train
4	Train	Test	Train							
...
437	Train	Train	Train	Test	Test	Train	Train	Train	Train	Train
438	Train									
439	Train									
440	Train									
441	Train	Train	Train	Train	Test	Train	Train	Train	Train	Train

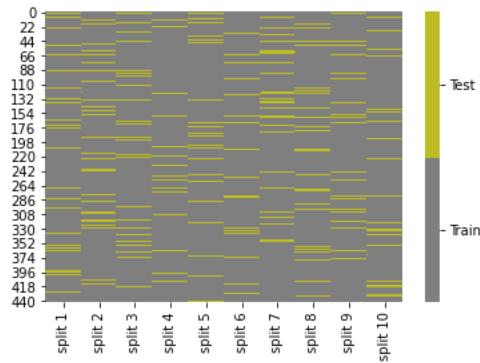
442 rows × 10 columns

And plot

```

cmap = sns.color_palette("tab10",10)
g = sns.heatmap(ss_tt_df.replace({'Test':1,'Train':0}),cmap=cmap[7:9],cbar_kws=
{'ticks':[.25,.75]},linewdiths=0,
                linecolor='gray')
colorbar = g.collections[0].colorbar
colorbar.set_ticklabels(['Train','Test'])

```



Now, we see the samples in each training set (gray along the columns) are a random subset of all of all of the samples.

And check the usage of each sample

```
sum(ss_tt_df.apply(count_test,axis = 1) ==1)
```

```
167
```

and exactly 9 training sets

```
sum(ss_tt_df.apply(count_test,axis = 1) ==9)
```

```
0
```

And the size of the splits

```
ss_tt_df.apply(count_test,axis = 0)
```

```
split 1    45
split 2    45
split 3    45
split 4    45
split 5    45
split 6    45
split 7    45
split 8    45
split 9    45
split 10   45
dtype: int64
```

and training set:

```
ss_tt_df.apply(count_train,axis = 0)
```

```
split 1    397
split 2    397
split 3    397
split 4    397
split 5    397
split 6    397
split 7    397
split 8    397
split 9    397
split 10   397
dtype: int64
```

Again the same sizes

Class 23: Interpreting Regression Evaluations

1. Snow in October. Share your thoughts in the zoom chat
2. Log onto prismia

```
# %load http://drsmb.co/310
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
import pandas as pd
from sklearn import datasets, linear_model
from sklearn.metrics import mean_squared_error, r2_score
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import train_test_split
```

```
X, y = datasets.load_boston(return_X_y= True)
```

```
X[:5]
```

```
array([[6.3200e-03, 1.8000e+01, 2.3100e+00, 0.0000e+00, 5.3800e-01,
       6.5750e+00, 6.5200e+01, 4.0900e+00, 1.0000e+00, 2.9600e+02,
       1.5300e+01, 3.9690e+02, 4.9800e+00],
      [2.7310e-02, 0.0000e+00, 7.0700e+00, 0.0000e+00, 4.6900e-01,
       6.4210e+00, 7.8900e+01, 4.9671e+00, 2.0000e+00, 2.4200e+02,
       1.7800e+01, 3.9690e+02, 9.1400e+00],
      [2.7290e-02, 0.0000e+00, 7.0700e+00, 0.0000e+00, 4.6900e-01,
       7.1850e+00, 6.1100e+01, 4.9671e+00, 2.0000e+00, 2.4200e+02,
       1.7800e+01, 3.9283e+02, 4.0300e+00],
      [3.2370e-02, 0.0000e+00, 2.1800e+00, 0.0000e+00, 4.5800e-01,
       6.9980e+00, 4.5800e+01, 6.0622e+00, 3.0000e+00, 2.2200e+02,
       1.8700e+01, 3.9463e+02, 2.9400e+00],
      [6.9050e-02, 0.0000e+00, 2.1800e+00, 0.0000e+00, 4.5800e-01,
       7.1470e+00, 5.4200e+01, 6.0622e+00, 3.0000e+00, 2.2200e+02,
       1.8700e+01, 3.9690e+02, 5.3300e+00]])
```

```
y[:5]
```

```
array([24. , 21.6, 34.7, 33.4, 36.2])
```

```
X_train, X_test, y_train, y_teest = train_test_split(X,y)
```

```
regr = linear_model.LinearRegression()
regr.fit(X_train, y_train)
```

```
LinearRegression()
```

```
regr.score(X_test,y_teest)
```

```
0.6645457841499955
```

```
y_pred = regr.predict(X_test)
```

```
# %load http://drsmb.co/310
col_name = ['ZN',
'INDUS',
'CHAS',
'NOX',
'RM',
'AGE',
'DIS',
'RAD',
'TAX',
'PTRATIO',
'B',
'LSTAT',
'MEDV']
```

```
test_df = pd.DataFrame(data=X_test, columns = col_name)
test_df.head()
```

	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	B	LST
0	9.82349	0.0	18.10	0.0	0.671	6.794	98.8	1.3580	24.0	666.0	20.2	396.
1	22.05110	0.0	18.10	0.0	0.740	5.818	92.4	1.8662	24.0	666.0	20.2	391.
2	0.29819	0.0	6.20	0.0	0.504	7.686	17.0	3.3751	8.0	307.0	17.4	377.
3	0.13914	0.0	4.05	0.0	0.510	5.572	88.5	2.5961	5.0	296.0	16.6	396.
4	0.02055	85.0	0.74	0.0	0.410	6.383	35.7	9.1876	2.0	313.0	17.3	396.

```
test_df['y_test'] = y_teest
test_df['y_pred'] = y_pred
```

```
error = lambda r: r['y_test'] - r['y_pred']
test_df['pred_error'] = test_df.apply(error, axis=1)
```

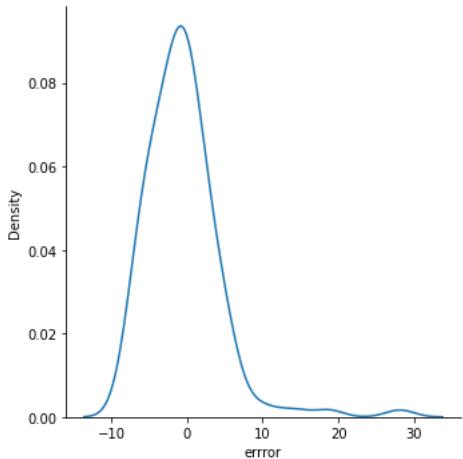
```
test_df['error'] = y_teest - y_pred
```

```
test_df.head()
```

	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	B	LST
0	9.82349	0.0	18.10	0.0	0.671	6.794	98.8	1.3580	24.0	666.0	20.2	396.
1	22.05110	0.0	18.10	0.0	0.740	5.818	92.4	1.8662	24.0	666.0	20.2	391.
2	0.29819	0.0	6.20	0.0	0.504	7.686	17.0	3.3751	8.0	307.0	17.4	377.
3	0.13914	0.0	4.05	0.0	0.510	5.572	88.5	2.5961	5.0	296.0	16.6	396.
4	0.02055	85.0	0.74	0.0	0.410	6.383	35.7	9.1876	2.0	313.0	17.3	396.

```
sns.displot(data=test_df, x='error', kind = 'kde')
```

```
<seaborn.axisgrid.FacetGrid at 0x7fa49d53eed0>
```



Class 24: Clustering

1. Log onto prismia
2. Say hello on Zoom

```
# %load http://drsmb.co/310
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns
from sklearn import datasets
from sklearn.cluster import KMeans
import string
import pandas as pd
```

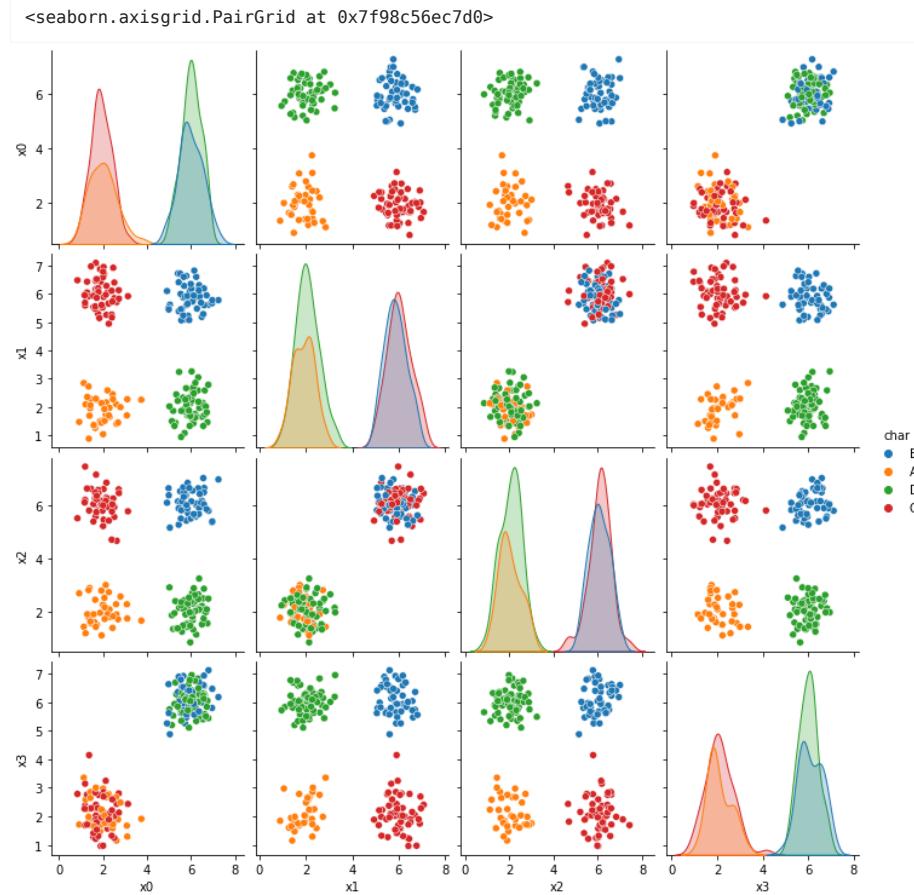
How Does Kmeans work?

We'll start with the data from dataset5 in assignment 6.

```
C = 4
N = 200
classes = list(string.ascii_uppercase[:C])
mu = {c: i for c, i in zip(classes,[[2,2, 2,2], [6,6,6,6], [2,6,6,2],[6,2,2,6]])}
sigma = {c: i*.5 for c, i in zip(classes,np.random.random(4))}

target5 = np.random.choice(classes,N)
data5 = [np.random.multivariate_normal(mu[c],.25*np.eye(C)) for c in target5]
df5 = pd.DataFrame(data = data5,columns = ['x' + str(i) for i in range(C)]).round(2)
rand_target = np.random.choice(classes,N)
obs_target = [np.random.choice([t,r],p=[.85,.15]) for t,r in zip(target5,rand_target)]
df5['char_noisy'] = obs_target
df5['char'] = target5

sns.pairplot(data =df5, hue='char')
```



We'll work with just two columns and the true labels

```
data_cols = ['x0','x1']
df = df5[data_cols]
df.head()
```

	x0	x1
0	6.24	6.59
1	3.08	1.71
2	1.12	2.85
3	6.21	5.87
4	5.83	1.96

We'll also setup some things for making our plots how we'd like in advance

```

def mu_to_df(mu,i):
    mu_df = pd.DataFrame(mu,columns=data_cols)
    mu_df['iteration'] = str(i)
    mu_df['class'] = ['M'+str(i) for i in range(K)]
    mu_df['type'] = 'mu'
    return mu_df

cmap_pt = sns.color_palette('tab20',8)[1::2]
cmap_mu = sns.color_palette('tab20',8)[0::2]

```

Now, we'll start our clustering. First we set how many clusters we want and then we'll try out randomly assigning the samples and see where that puts the means

```

K = 4

df['0'] = np.random.choice(K,size=N)
sfig = sns.scatterplot(data =df,x='x0',y='x1',hue='0',palette=cmap_pt,legend=False)

mu = df.groupby('0')[data_cols].mean().values
mu_df = mu_to_df(mu,0)

# sns.scatterplot(data
=mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig,legend=False)
sfig.get_figure().savefig('kmeans00.png')
# sfig = sns.scatterplot(data =df,x='x0',y='x1',hue='1'
sns.scatterplot(data =mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig)

```

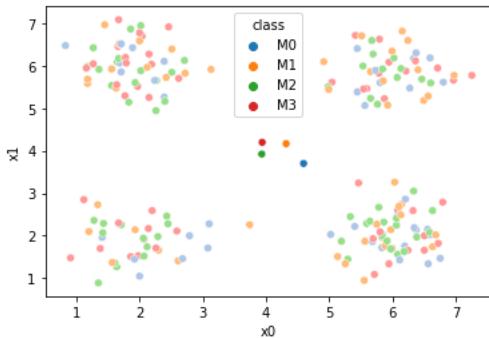
```

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/ipykernel_launcher.py:3: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-
docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
This is separate from the ipykernel package so we can avoid doing imports until

```

<AxesSubplot:xlabel='x0', ylabel='x1'>



This puts the means all really close to the center, so we'll try a different initialization for those: randomly sampling 4 points

```

mu = df[data_cols].sample(n=K).values
mu

```

```

array([[5.59, 2.29],
       [1.45, 6.96],
       [1.97, 5.51],
       [6.06, 2.25]])

```

Now we can compute the distance from each point to each of the four means and use the closest one as an assignment. We'll look at the assignments again.

```

i = 1
df[str(i)] = pd.concat([(df[data_cols]-mu_i)**2).sum(axis=1) for mu_i in
mu],axis=1).idxmin(axis=1)

sfig = sns.scatterplot(data =df,x='x0',y='x1',hue='1',palette=cmap_pt,legend=False)
# plt.plot(mu[:,0],mu[:,1],marker='s',linewidth=0)
mu_df = mu_to_df(mu,i)
sns.scatterplot(data
=mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig,legend=False)
sfig.get_figure().savefig('kmeans01.png')

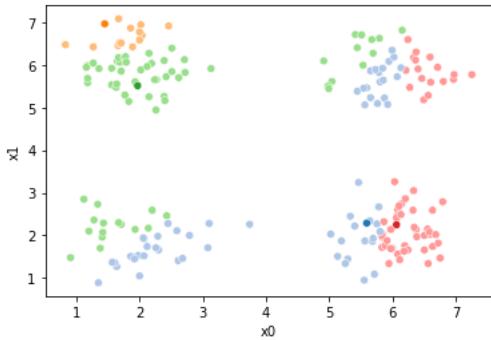
```

```

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-
packages/ipykernel_launcher.py:2: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-
docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy

```

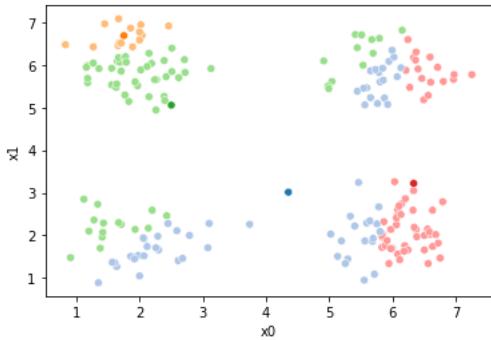


Now, we can use those assignments, to compute new means:

```

mu = df.groupby('1')[data_cols].mean().values
# sfig = sns.scatterplot(data =df,x='x0',y='x1',hue='1')
# plt.plot(mu[:,0],mu[:,1],marker='s',linewidth=0)
fig = plt.figure()
mu_df = mu_to_df(mu,i)
sfig = sns.scatterplot(data =df,x='x0',y='x1',hue='1',palette=cmap_pt,legend=False)
sns.scatterplot(data
=mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig,legend=False)
sfig.get_figure().savefig('kmeans02.png')

```



Now we can set up a while loop to continue the iterations

```

mu_list = [mu_to_df(mu,i)]
cur_old = str(i-1)
cur_new = str(i)
while sum(df[cur_old] != df[cur_new]) >0:
    cur_old = cur_new
    i +=1
    cur_new = str(i)
    #      update the assignments and plot with the associated means
    df[cur_new] = pd.concat([(df[data_cols]-mu_i)**2).sum(axis=1) for mu_i in
mu],axis=1).idxmin(axis=1)
    fig = plt.figure()
    sfig = sns.scatterplot(data
=df,x='x0',y='x1',hue=cur_new,palette=cmap_pt,legend=False)
    sns.scatterplot(data
=mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig,legend=False)
    file_num = str(i*2 -1).zfill(2)
    sfig.get_figure().savefig('kmeans' +file_num + '.png')

#      update the means and plot with current generating assignments
mu = df.groupby(cur_new)[data_cols].mean().values
mu_df = mu_to_df(mu,i)
mu_list.append(mu_df)

fig = plt.figure()
sfig = sns.scatterplot(data
=df,x='x0',y='x1',hue=cur_new,palette=cmap_pt,legend=False)
    sns.scatterplot(data
=mu_df,x='x0',y='x1',hue='class',palette=cmap_mu,ax=sfig,legend=False)
#      plt.plot(mu[:,0],mu[:,1],marker='s',linewidth=0)

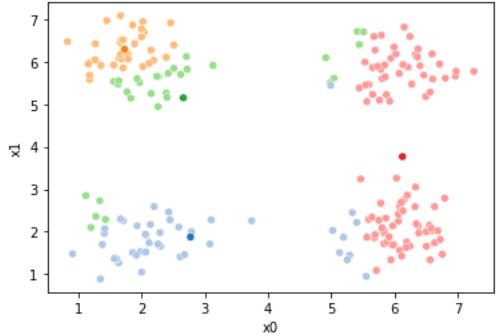
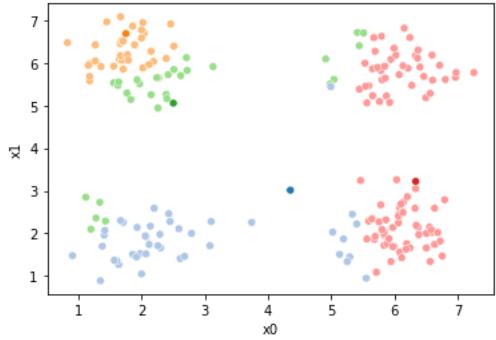
file_num = str(i*2).zfill(2)
sfig.get_figure().savefig('kmeans' +file_num + '.png')

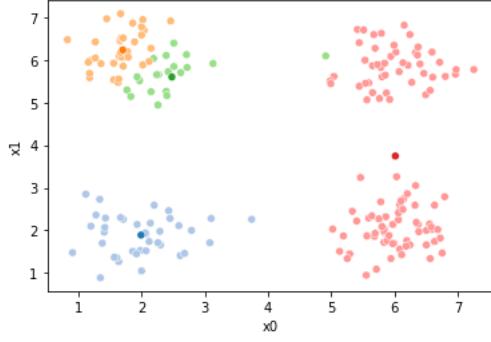
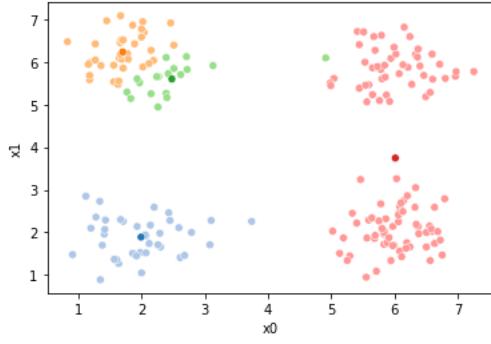
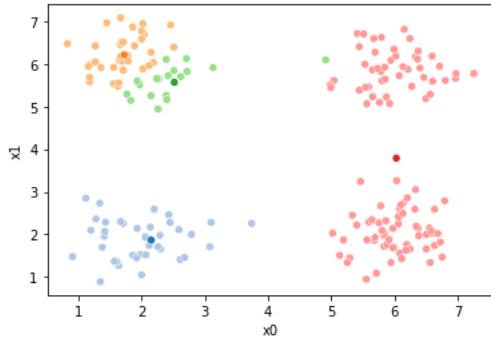
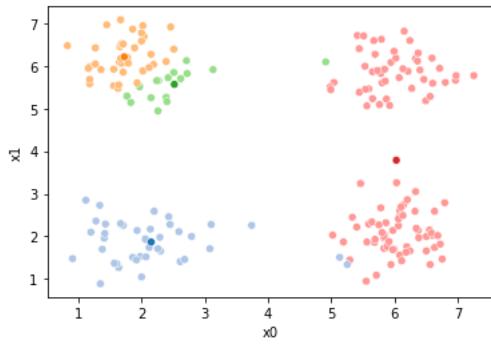
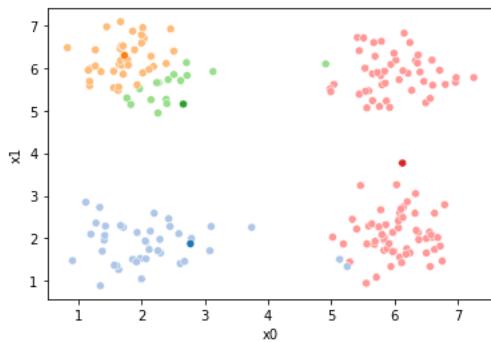
n_iter = i

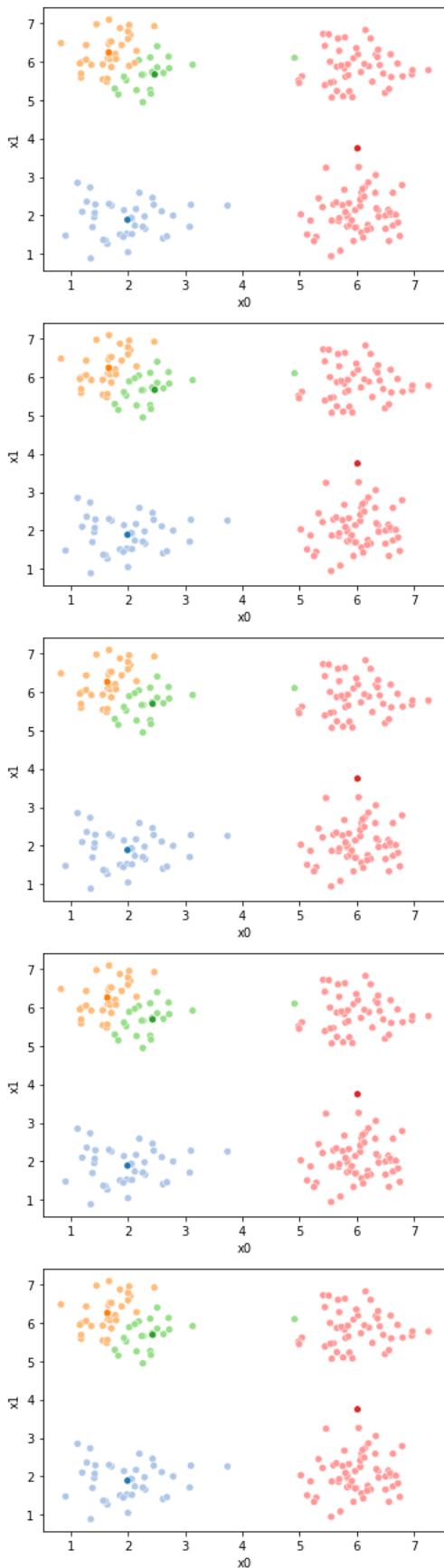
```

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-packages/ipykernel_launcher.py:9: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
if __name__ == '__main__':







These plots can be saved and merged into a gif with, for example [imagemagick](#) to create the gifs above. Since we've saved them in a single dataframe, we can also look at that

```
df.head()
```

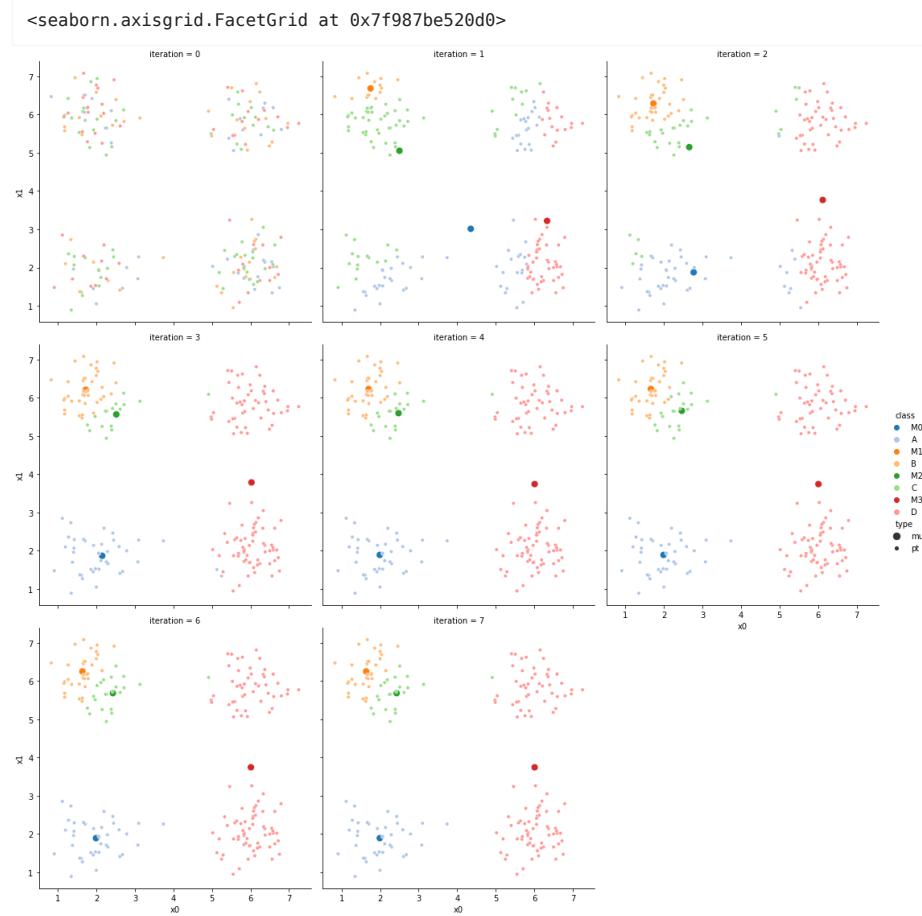
	x0	x1	0	1	2	3	4	5	6	7
0	6.24	6.59	1	3	3	3	3	3	3	3
1	3.08	1.71	0	0	0	0	0	0	0	0
2	1.12	2.85	3	2	2	0	0	0	0	0
3	6.21	5.87	3	3	3	3	3	3	3	3
4	5.83	1.96	1	3	3	3	3	3	3	3

We can also manipulate the dataframe to make plotting them all together easier.

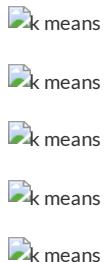
```
df_vis = df.melt(id_vars = ['x0','x1'], var_name ='iteration',value_name='class')
df_vis.replace({'class':{i:c for i,c in
enumerate(string.ascii_uppercase[:C])}},inplace=True)

df_vis['type'] = 'pt'
df_mu_vis = pd.concat([pd.concat(mu_list),df_vis])
cmap = sns.color_palette('tab20',8)
n_iter = 5

sns.relplot(data=df_mu_vis,x='x0',y='x1',hue='class',col='iteration',
            col_wrap=3,hue_order = ['M0','A','M1','B','M2','C','M3','D'],
            palette = cmap,size='type',col_order=[str(i) for i in range(n_iter+1)])
```



Here are a few different runs of the algorithm on the same dataset.



KMeans with Sklearn

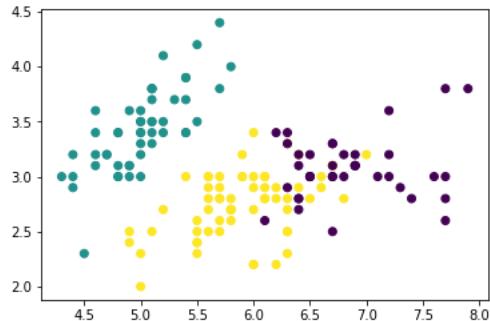
```
iris_X, iris_y = datasets.load_iris(return_X_y=True)
```

```
km3 = KMeans(n_clusters=3)
```

```
iris_cluster3 = km3.fit_predict(iris_X)
```

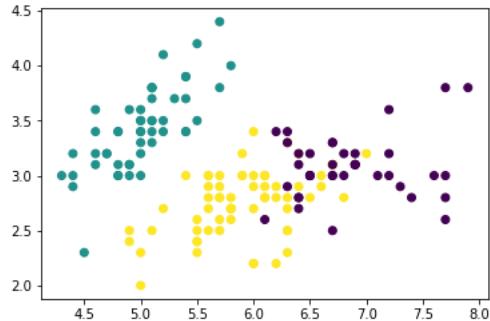
```
plt.scatter(iris_X[:,0],iris_X[:,1],c=iris_cluster3)
```

```
<matplotlib.collections.PathCollection at 0x7f987b7504d0>
```



```
iris_cluster3 = km3.fit_predict(iris_X)
plt.scatter(iris_X[:,0],iris_X[:,1],c=iris_cluster3)
```

```
<matplotlib.collections.PathCollection at 0x7f987b5cce90>
```



Class 25: Evaluating Clustering

1. Log onto Prismia

2. (forgot to use this on monday, honoring post-halloween) share your favorite candy in the zoom chat (or just say hi)

```
# %load http://drsmb.co/310
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns
from sklearn import datasets
from sklearn import cluster
from sklearn import metrics
```

```
iris_X, iris_y = datasets.load_iris(return_X_y=True)
```

```
km3 = cluster.KMeans(n_clusters=3)
```

```
km3.fit(iris_X)
```

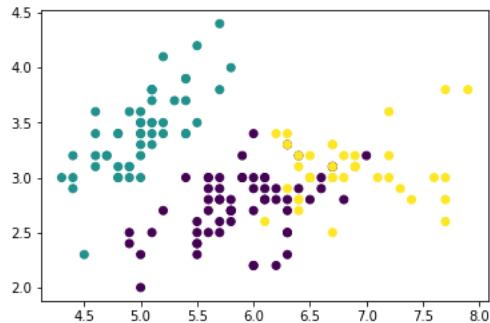
```
KMeans(n_clusters=3)
```

```
km3.cluster_centers_
```

```
array([[5.9016129 , 2.7483871 , 4.39354839, 1.43387097],  
      [5.006      , 3.428      , 1.462      , 0.246      ],  
      [6.85       , 3.07368421, 5.74210526, 2.07105263]])
```

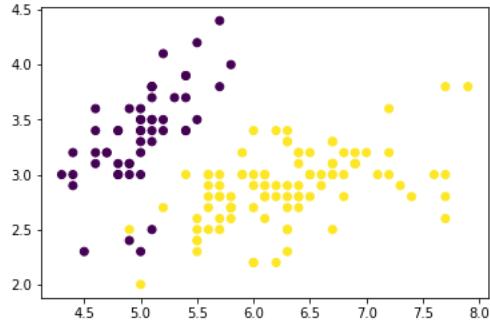
```
plt.scatter(iris_X[:,0],iris_X[:,1],c=km3.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f553cc6f210>
```



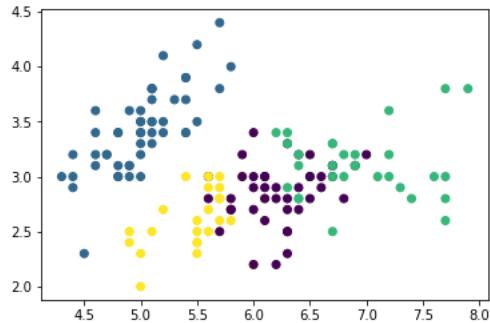
```
km2 = cluster.KMeans(n_clusters=2)  
km2.fit(iris_X)  
plt.scatter(iris_X[:,0],iris_X[:,1],c=km2.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f553cb6ca50>
```



```
km4 = cluster.KMeans(n_clusters=4)  
km4.fit(iris_X)  
plt.scatter(iris_X[:,0],iris_X[:,1],c=km4.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f553cae8d0>
```



$$s = \frac{b - a}{\max(a, b)}$$

a: The mean distance between a sample and all other points in the same class.

b: The mean distance between a sample and all other points in the next nearest cluster.

```
metrics.silhouette_score(iris_X, km3.labels_)
```

```
0.5528190123564091
```

```
metrics.silhouette_score(iris_X, km2.labels_)
```

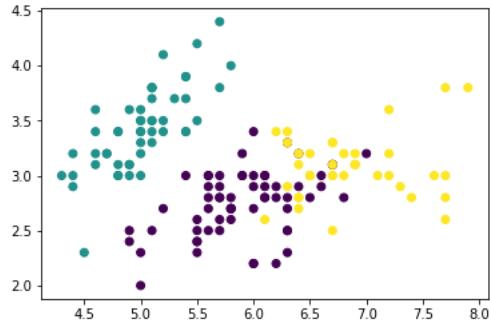
```
0.681046169211746
```

```
metrics.silhouette_score(iris_X, km4.labels_)
```

```
0.49745518901737446
```

```
km3a = cluster.KMeans(n_clusters=3)
km3a.fit(iris_X)
plt.scatter(iris_X[:,0],iris_X[:,1],c=km3.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f553ca5cf90>
```



```
metrics.adjusted_rand_score(iris_y,km3.labels_)
```

```
0.7302382722834697
```

```
metrics.adjusted_rand_score(iris_y,km2.labels_)
```

```
0.5399218294207123
```

```
metrics.adjusted_rand_score(iris_y,km4.labels_)
```

```
0.6460787233460947
```

```
metrics.adjusted_mutual_info_score(iris_y,km3.labels_)
```

```
0.7551191675800485
```

```
metrics.adjusted_mutual_info_score(iris_y,km2.labels_)
```

```
0.653838071376278
```

```
metrics.adjusted_mutual_info_score(iris_y,km4.labels_)
```

```
0.7151742230795869
```

Class 26: More Clustering Models

```
# %load http://drsmmb.co/310
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns
from sklearn import datasets
from sklearn import cluster
from sklearn import metrics
```

```
iris_X, iris_y = datasets.load_iris(return_X_y = True)
```

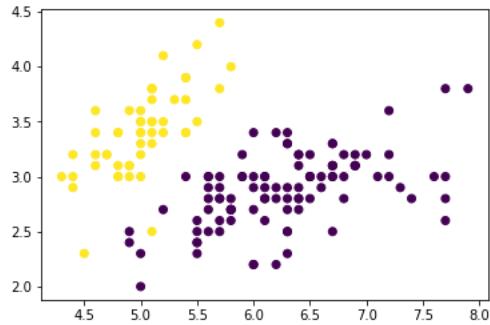
```
msc = cluster.MeanShift()
```

```
msc.fit(iris_X)
```

```
MeanShift()
```

```
plt.scatter(iris_X[:,0],iris_X[:,1], c= msc.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f4c5dfa2f50>
```



```
sns.pairplot(iris_X, hue= msc.labels_)
```

```
-----  
TypeError                                                 Traceback (most recent call last)  
<ipython-input-6-7e1512556191> in <module>  
----> 1 sns.pairplot(iris_X, hue= msc.labels_)

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-  
packages/seaborn/_decorators.py in inner_f(*args, **kwargs)  
    44         )  
    45     kwargs.update({k: arg for k, arg in zip(sig.parameters, args)})  
--> 46     return f(**kwargs)  
    47     return inner_f  
    48

/opt/hostedtoolcache/Python/3.7.9/x64/lib/python3.7/site-packages/seaborn/axisgrid.py  
in pairplot(data, hue, hue_order, palette, vars, x_vars, y_vars, kind, diag_kind,  
    markers, height, aspect, corner, dropna, plot_kws, diag_kws, grid_kws, size)  
    1915     raise TypeError(  
    1916         "'data' must be pandas DataFrame object, not:  
{typefound}'.format(  
-> 1917             typefound=type(data)))  
    1918  
    1919     plot_kws = {} if plot_kws is None else plot_kws.copy()

TypeError: 'data' must be pandas DataFrame object, not: <class 'numpy.ndarray'>
```

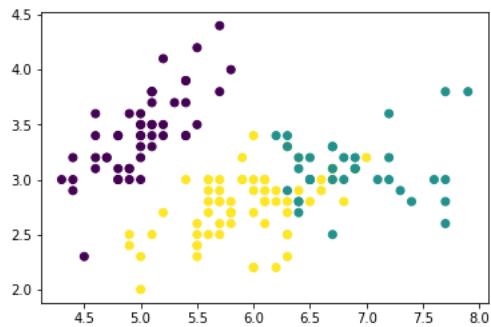
```
spc = cluster.SpectralClustering(n_clusters=3)
```

```
spc.fit(iris_X)
```

```
SpectralClustering(n_clusters=3)
```

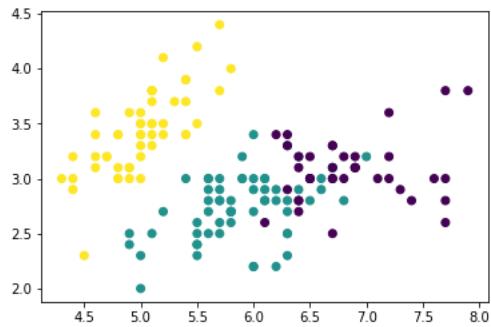
```
plt.scatter(iris_X[:,0],iris_X[:,1], c=spc.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f4c5dd9cd50>
```



```
km3 = cluster.KMeans(n_clusters=3)
km3.fit(iris_X)
plt.scatter(iris_X[:,0],iris_X[:,1], c=km3.labels_)
```

```
<matplotlib.collections.PathCollection at 0x7f4c5bd19b10>
```



```
from sklearn.neighbors import NearestNeighbors
```

```
nbrs = NearestNeighbors()
```

```
nbrs.fit(iris_X)
```

```
NearestNeighbors()
```

```
dists, indx = nbrs.kneighbors(iris_X)
```

```
dists
```

```

array([[0.          , 0.1         , 0.14142136, 0.14142136, 0.14142136],
[0.          , 0.14142136, 0.14142136, 0.14142136, 0.17320508],
[0.          , 0.14142136, 0.24494897, 0.26457513, 0.26457513],
[0.          , 0.14142136, 0.17320508, 0.2236068 , 0.24494897],
[0.          , 0.14142136, 0.14142136, 0.17320508, 0.17320508],
[0.          , 0.33166248, 0.34641016, 0.36055513, 0.37416574],
[0.          , 0.2236068 , 0.26457513, 0.3         , 0.31622777],
[0.          , 0.1         , 0.14142136, 0.17320508, 0.2         ],
[0.          , 0.14142136, 0.3         , 0.31622777, 0.34641016],
[0.          , 0.1         , 0.17320508, 0.17320508, 0.17320508],
[0.          , 0.1         , 0.28284271, 0.3         , 0.33166248],
[0.          , 0.2236068 , 0.2236068 , 0.28284271, 0.3         ],
[0.          , 0.14142136, 0.17320508, 0.2         , 0.2         ],
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[0.          , 0.36055513, 0.54772256, 0.6164414 , 0.6164414 ],
[0.          , 0.34641016, 0.36055513, 0.38729833, 0.38729833],
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[0.          , 0.28284271, 0.3         , 0.36055513, 0.36055513],
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[0.          , 0.45825757, 0.50990195, 0.50990195, 0.51961524],
[0.          , 0.2         , 0.26457513, 0.37416574, 0.38729833],
[0.          , 0.3         , 0.37416574, 0.41231056, 0.42426407],
[0.          , 0.17320508, 0.2         , 0.2236068 , 0.2236068 ],
[0.          , 0.2         , 0.2236068 , 0.2236068 , 0.24494897],
[0.          , 0.14142136, 0.14142136, 0.14142136, 0.17320508],
[0.          , 0.14142136, 0.14142136, 0.14142136, 0.17320508],
[0.          , 0.14142136, 0.17320508, 0.2236068 , 0.2236068 ],
[0.          , 0.14142136, 0.14142136, 0.17320508, 0.2236068 ],
[0.          , 0.28284271, 0.3         , 0.3         , 0.31622777],
[0.          , 0.34641016, 0.34641016, 0.37416574, 0.42426407],
[0.          , 0.34641016, 0.36055513, 0.38729833, 0.41231056],
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[0.          , 0.2236068 , 0.3         , 0.31622777, 0.33166248],
[0.          , 0.3         , 0.31622777, 0.33166248, 0.34641016],
[0.          , 0.14142136, 0.24494897, 0.26457513, 0.26457513],
[0.          , 0.14142136, 0.2         , 0.24494897, 0.3         ],
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[0.          , 0.6244998 , 0.71414284, 0.76811457, 0.78102497],
[0.          , 0.2         , 0.2236068 , 0.3         , 0.3         ],
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[0.          , 0.14142136, 0.14142136, 0.2236068 , 0.2236068 ],
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[0.          , 0.14142136, 0.17320508, 0.2236068 , 0.2236068 ],
[0.          , 0.26457513, 0.33166248, 0.43588989, 0.45825757],
[0.          , 0.26457513, 0.31622777, 0.34641016, 0.37416574],
[0.          , 0.26457513, 0.28284271, 0.31622777, 0.34641016],
[0.          , 0.2         , 0.3         , 0.31622777, 0.43588989],
[0.          , 0.24494897, 0.31622777, 0.37416574, 0.37416574],
[0.          , 0.3         , 0.31622777, 0.31622777, 0.33166248],
[0.          , 0.26457513, 0.37416574, 0.42426407, 0.45825757],
[0.          , 0.14142136, 0.38729833, 0.45825757, 0.72111026],
[0.          , 0.24494897, 0.24494897, 0.31622777, 0.31622777],
[0.          , 0.38729833, 0.50990195, 0.51961524, 0.52915026],
[0.          , 0.36055513, 0.45825757, 0.67082039, 0.71414284],
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[0.          , 0.42426407, 0.4472136 , 0.50990195, 0.51961524],
[0.          , 0.14142136, 0.31622777, 0.31622777, 0.34641016],
[0.          , 0.2         , 0.3         , 0.38729833, 0.41231056],
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[0.          , 0.33166248, 0.34641016, 0.37416574, 0.4         ],
[0.          , 0.36055513, 0.36055513, 0.41231056, 0.42426407],
[0.          , 0.2236068 , 0.3         , 0.38729833, 0.43588989],
[0.          , 0.2         , 0.26457513, 0.36055513, 0.38729833],
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```

```

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```

indx

```
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```

```
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[145, 141, 147, 139, 112],
[146, 123, 111, 126,  72],
[147, 110, 111, 116, 145],
[148, 136, 115, 110, 147],
[149, 127, 138, 101, 142]])
```

Class 27: Model Optimization- Choosing K

1. On the zoom chat, say hello
2. Log onto Prismia

```
# %load http://drsmb.co/310
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns
import pandas as pd
from sklearn import datasets
from sklearn import cluster
from sklearn import metrics

url ='https://raw.githubusercontent.com/rhodyprog4ds/06-naive-
bayes/main/data/dataset2.csv'
```

```
df = pd.read_csv(url)
```

```
X = df.values[:, :-1]
```

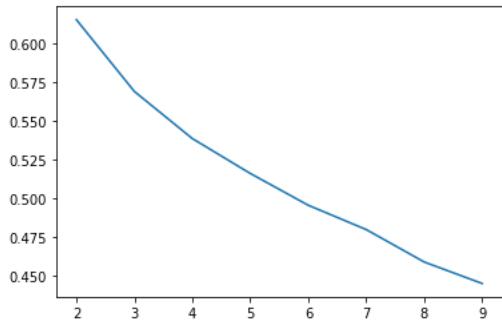
```
score = []
for k in range(2, 10):
    km = cluster.KMeans(n_clusters=k)
    km.fit(X)
    score.append(metrics.silhouette_score(X, km.labels_))
```

```
score
```

```
[0.6152428695158447,
 0.5689398715407455,
 0.5385291617893929,
 0.5160098177580555,
 0.4954892476648668,
 0.4797917245143165,
 0.4588725916710707,
 0.444987563684623]
```

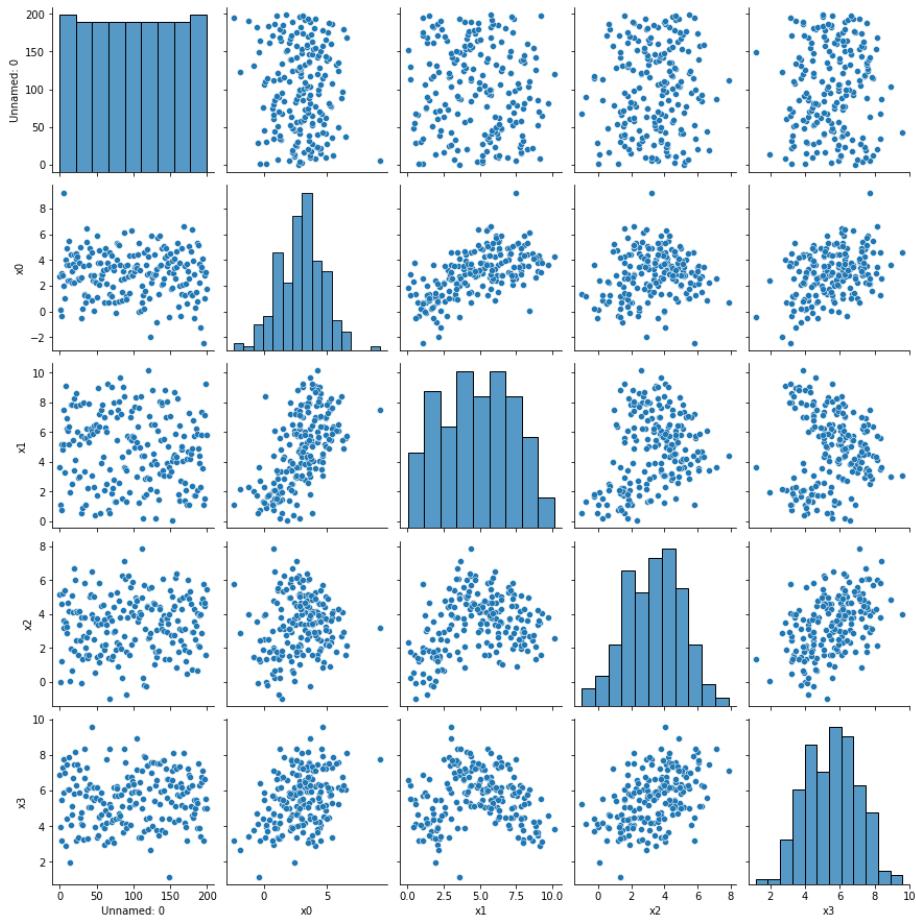
```
plt.plot(range(2, 10), score)
```

```
<matplotlib.lines.Line2D at 0x7f8f95a67090>
```



```
sns.pairplot(df)
```

```
<seaborn.axisgrid.PairGrid at 0x7f8f95a0cdd0>
```



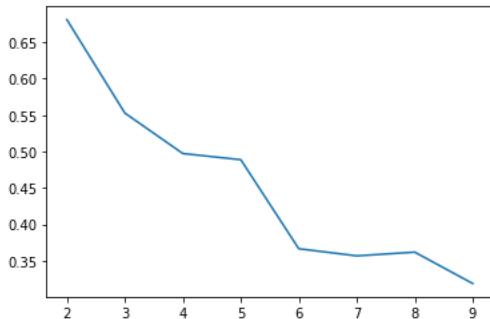
```
iris_X, _ = datasets.load_iris(return_X_y = True)
```

```
score_iris = []

for k in range(2,10):
    km = cluster.KMeans(n_clusters=k)
    km.fit(iris_X)
    score_iris.append(metrics.silhouette_score(iris_X,km.labels_))
```

```
plt.plot(range(2,10), score_iris)
```

[<matplotlib.lines.Line2D at 0x7f8f95850c10>]

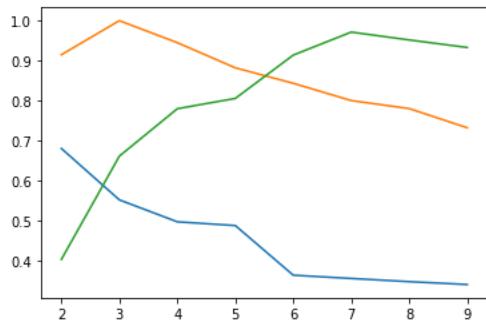


```
silohette_iris = []
ch_iris = []
db_iris = []

for k in range(2,10):
    km = cluster.KMeans(n_clusters=k)
    km.fit(iris_X)
    silohette_iris.append(metrics.silhouette_score(iris_X,km.labels_))
    ch_iris.append(metrics.calinski_harabasz_score(iris_X,km.labels_))
    db_iris.append(metrics.davies_bouldin_score(iris_X,km.labels_))
```

```
ch_iris = np.asarray(ch_iris)/np.max(ch_iris)
plt.plot(range(2,10), silhouette_iris)
plt.plot(range(2,10), ch_iris)
plt.plot(range(2,10), db_iris)
```

[<matplotlib.lines.Line2D at 0x7f8f94de7390>]



Assignments

All assignments are due on Sunday at 11:59pm, via github unless otherwise noted.

Assignment TOC:

- [Assignment 1](#) Due September 13
- [Assignment 2](#) Due September 20
- [Assignment 3](#) Due September 29
- [Assignment 4](#) Due October 4
- [Assignment 5](#) Due October 11
- [Assignment 6](#) Due October 20
- [Assignment 7](#) Due October 27
- [Assignment 8](#) Due November 3

Assignment 1: Portfolio Setup, Data Science, and Python

Due: 2020-09-13

Objective & Evaluation

This assignment is an opportunity to earn level 2 achievements for the `process` and `python` and confirm that you have all of your tools setup, including your portfolio.

To Do

Your task is to:

1. Install required software
 2. Setup your portfolio, by [accepting the assignment](#) and following the instructions in the README file on your repository.
 3. Add your own definition of data science to the introduction of your portfolio, in `about/index.md`
 4. Add a Jupyter notebook called `grading.ipynb` to the `about` folder and write a function that computes a grade for this course, with the following docstring. Include:
 - a Markdown cell with a heading
 - your function called `compute_grade`
 - three calls to your function that verify it returns the correct value for different number of badges that produce at three different letter grades.
1. Uncomment the line `# - file: about/grading` in your `_toc.yml` file.

Note

If you get stuck on any of this after accepting the assignment and creating a repository, you can create an issue on your repository, describing what you're stuck on and tag us with `@rhodyprog4ds/fall20instructors`.

To do this click Issues at the top, the green "New Issue" button and then type away.

```

...
Computes a grade for CSC/DSP310 from numbers of achievements at each level

Parameters:
-----
num_level1 : int
    number of level 1 achievements earned
num_level2 : int
    number of level 2 achievements earned
num_level3 : int
    number of level 3 achievements earned

Returns:
-----
letter_grade : string
    letter grade with possible modifier (+/-)
...

```

Here are some sample tests you could run to confirm that your function works correctly:

```

assert compute_grade(15,15,15) == 'A'
assert compute_grade(15,15,13) == 'A- '
assert compute_grade(15,14,14) == 'B- '
assert compute_grade(14,14,14) == 'C- '
assert compute_grade(4,3,1) == 'D'
assert compute_grade(15,15,6) == 'B+'

```

Warning

your function can have a different name than `compute_grade`, but make sure it's your function name, with those parameter values in your tests.

Note

when the value of the expression after `assert` is `True`, it will look like nothing happened. `assert` is used for testing

Submission Instructions

Create a Jupyter Notebook with your function and Add the notebook to your portfolio by uploading it to your repository, or adding to the folder off line and committing and pushing the changes.

View the `gh-pages` branch to see your compiled submission, as `portfolio.pdf` or by viewing your website.

There will be a pull request on your repository that is made by GitHub classroom, [request a review](#) from the team `rhydyp4ds/Fall20instructors`.

Solutions

One solution is added to the [Detailed Mechanics](#) part of the Grading section of the syllabus.

Assignment 2: Practicing Python and Accessing Data

Due: 2020-09-20

Objective & Evaluation

This assignment is an opportunity to earn level 1 or 2 achievements in `python`, `process` and `access` and begin working toward level 1 in `summarize`.

Accept the assignment on [GitHub Classroom](#). It contains a notebook with some template structure (and will set you up for grading). The template will also convert notebooks that are added to markdown, which makes reading on GitHub for easier grading. If you want to incorporate feedback you receive back into a notebook file, [Jupytext](#) can do that.

To work with this notebook you can either:

- download the repository as .zip from the green code button, unzip, and re-upload, OR
- clone the repository with git and the push your changes. See Git/GitHub help on cloning, committing, and pushing, for example this [tutorial on git](#) to learn more about git.

Accessing Data with Python and pandas

(for `python` and `access`)

Find 3 datasets of interest to you that are provided in different file formats. Choose datasets that are not too big, so that they do not take more than a few seconds to load. At least one dataset must have non-numerical (e.g. string or boolean) data in at least 1 column. Complete a dictionary for each with the url, a name, and what function should be used to load the data into a `pandas.DataFrame`.

Use a list of those dictionaries to iterate over the datasets and build a table that describes them, with the following columns [`'name'`, `'source'`, `'num_rows'`, `'num_columns'`, `'source_file_name'`]. The `source` column should be the URL where you loaded the data from or the source if you downloaded it from a website first. The `source_file_name` should be the part of the URL after the last `/`, you should extract this programmatically. Display that summary table as a DataFrame and save it as a CSV, named `dataset_summary.csv`.

💡 Tip

URLs are strings. The `string` class in Python has a lot of helpful methods for manipulating strings, like `split`.

ℹ️ Note

If you download the datasets (or find them as .zip and need to) you can use the local path instead of the URL, but include a Markdown cell with links to where you got your data from.

💡 Tip

You can create a `pandas.DataFrame` using the `constructor` and you can build lists (or lists of lists) using the `append` method

For one dataset (must include non-numerical data):

- display the heading with the last seven rows
- make and display a new DataFrame with only the non-numerical columns
- was the format that the data was provided in a good format? Why or why not?

💡 Tip

For a second dataset:

- display the heading and the first three rows
- display the datatype for each column
- Are there any variables where pandas may have read in the data as a datatype that's not what you expect (e.g. a numerical column mistaken for strings)?

For the third dataset:

- display the first 5 even rows of the data for three columns of your choice

For any dataset:

- try reading it in with the wrong `read_` function. If you had done this by accident, how could you tell?

Data Science Process

(for the `process` skill)

Make a list of a data science pipeline and denote which types of programming might be helpful at each stage. Include this in a Markdown cell in the same notebook with your analysis.

💡 Tip

Remember that this will be graded based on the rubric and that it should reflect your understanding, not be simply copied from a source. Also, always cite your sources, informal linking is OK.

To make a link in Markdown

```
[text to display](http://url.com/of/the/site/your/are/linking/)
```

Assignment 3: Exploratory Data Analysis

Due: 2020-09-27

Objective & Evaluation

This assignment is an opportunity to earn level 1 or 2 achievements in `summarize`, `visualize`, or `access`. You can earn level 2 in `python`.

Accept the assignment on [GitHub Classroom](#). The template will convert notebooks that are added to markdown, which makes reading on GitHub for easier grading. It will sync between .ipynb and .md style notebooks stored in your repository.

This week I encourage you to try working with git, but if you're not comfortable with that you can work via upload again.

Exploratory Data Analysis

This week your goal is to do a small exploratory data analysis for two datasets of your choice. One dataset must include at least two continuous valued variables and at least one categorical variable(d1). One dataset must include at least two categorical variables and at least one continuous valued variable(d2).

Use a separate notebook for each dataset, name them `dataset_01.ipynb` and `dataset_02.ipynb`.

For each dataset:

1. Include a markdown header with a title for your analysis
2. Load the data to a notebook as a `DataFrame` from url.
3. Explore the dataset in a notebook enough to describe its structure
 - shape
 - columns
 - variable types
4. Write a short description of what the data contains and what it could be used for
5. Complete an exploratory analysis with statistics and plots. Your analysis should include markdown cells describing the results you see, not only what you did. Your analysis should be structured to follow the steps below, for the corresponding dataset.

For d1:

1. Display all of the summary statistics for a subset of 5 of your choice or all variables if there are fewer than 5 numerical values
2. Display all of summary statistics grouped by a categorical variable
3. For two continuous variables make a scatter plot and color the points by a categorical variable
4. Pose one question for this dataset that can be answered with summary statistics, compute a statistic and plot that help answer that exploratory question.

For d2:

1. Display two individual summary statistics for one variable
2. Group the data by two categorical variables and display a table of one summary statistic
3. Use a seaborn plotting function with the `col` parameter or a `FacetGrid` to make a plot that shows something informative about this data, using both categorical variables and at least one numerical value. Describe what this tells you about the data.
4. Produce one additional plot of a different plot type that shows something about this data.

In total, in each of your notebooks you will have:

- a loaded dataset
- a basic description
- at least two summary statistic calculations
- at least two plots

💡 Tip

A continuous valued variable is one that can take on infinitely many values. Examples include temperature, length, age, etc. A categorical value is one that can only take on one of a subset of specific values. These require a bit more context to define often. For example color could be recorded in a basically continuous way as RGB values in HEX, but color of the stoplight will only be red, green, or yellow. Town of birth in a dataset of celebrities would not be a good categorical variable, there might not be many duplicates to do analysis with, but metro area of current residence for celebrities would be because we'd see mostly major cities and could do analyses.

💡 Tip

If you use a local copy of the dataset you can load from a `relative_path` instead of a url. Please include the data in a data folder that's in the folder where your notebook is so that the path is something like: `data/best-dataset-ever.csv` and upload the data to your assignment repository as well.

💡 Tip

Recall how we used a single function to see many statistics in class, use that for dataset 1. For dataset 2, you can make your own combination of summary statistics with `pandas.DataFrame.agg`

Assignment 4: Preparing Data for Analysis

Due: 2020-10-04

Objective & Evaluation

You can earn prepare level 2 by cleaning two datasets as outlined in the `cs_degrees.ipynb` and `travel_time.ipynb` notebooks.

To earn access level 2 you must clean a third dataset, following the `airlines.ipynb` or `safi_full.ipynb` notebooks. Note that this is the SAFI dataset we've been working with in class, but a messier version of it. You know what some of the fixes are because they're in the notes and some of what it will look like when it's done.

To earn python level 2, you must complete [safi_full.ipynb](#). Specifically, you must successfully use list or dictionary comprehensions and conditional statements, beyond those provided as hints.

To earn level 2 for summarize and visualize, include additional analyses after cleaning the datasets. You'll need at least two types of plots and two different ways of using summary statistics and to interpret the results.

Instructions

For this assignment, your assignment is to clean datasets in notebooks and include narrative description of how you're making decisions about the data cleaning. At the end of each cleaning, save the cleaned dataset to the data folder with an informative file name.

When you [accept the assignment](#), there will be template notebooks in the repository. These have significant hints to guide your efforts.

There is an issue type with a todo list, try that out to plan and track questions you have.

You can work with this assignment in 2 ways:

Git Workflow

1. Click the green code button and copy the url.
2. clone the repository as below where the url is the one you copied from GitHub. In your terminal on Linux or Mac or on the GitBash on Windows ([install instructions on tools section of syllabus](#))

```
cd path/where/you/want/a/new/folder/for/this/assignment  
git clone http://github.com/rhodyprog4ds/04-prepare-username
```

1. then launch your notebook from the newly created folder.

on Linux or Mac, in the same terminal (remember you can use tab complete)

```
cd 04-prepare-username  
jupyter notebook
```

or on windows, in your anaconda prompt

```
cd path/where/you/want/a/new/folder/for/this/assignment/04-prepare-username  
jupyter notebook
```

1. work on your assignment, by opening the notebooks there and editing them.
2. commit your changes when you want to save a point in your progress. Either you have part working and want to save that, or you want feedback, or you're done. On whichever terminal you used for the `git clone` command above

```
git add .  
git commit -m 'description of current status of project'
```

1. push your changes when you want to share them on GitHub, (eg need help, want feedback or complete)

```
git push
```

1. if you will keep working after pushing, first pull down the .md conversion that was added by GitHub actions. If you don't you'll have to merge, it should be fine, but ask if you're not sure.

```
git pull
```

Download and upload workflow

1. Click the green code button
2. choose the download via zip option
3. unzip
4. launch a notebook in the folder where you unzipped
5. Upload only the files you changed into the repository to replace their previous versions with the add file via upload button on GitHub. Put the notebooks at the top level and the data files in the data folder.

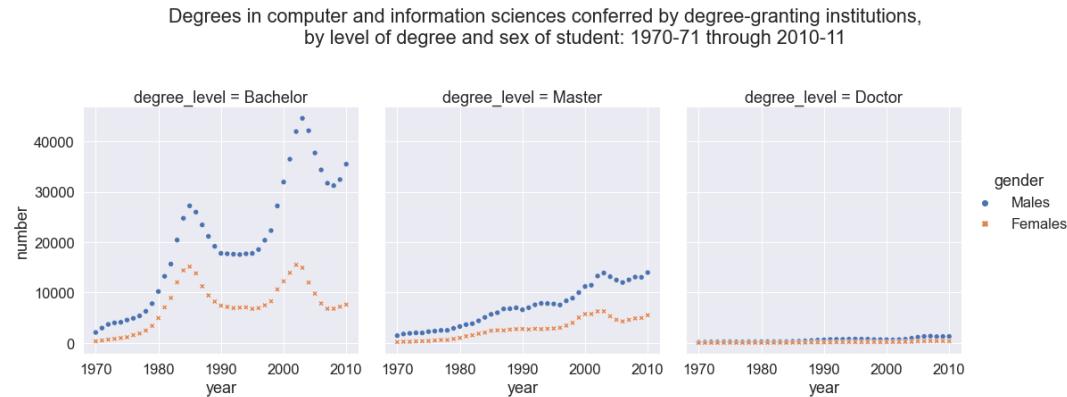
Tip

You can download any of the files and open them with another program if you need to by pasting the url in your browser. You'll need to display relevant parts in a notebook for grading, but if looking another way helps, you can try that.

Tips and hints

- Pandas can [convert string to dates](#) to be able to work with them more flexibly
- With python [date](#) objects you can get the [day of the week](#) from a date
- Pandas [read_excel](#) has:
 - a `header` parameter can take a list as its value
 - two parameters that allow you to skip rows when you read in the data
- `list(range(7))` check out what this line does
- Pandas [drop](#) can work a lot of different ways

The goal of the plot for the CS degrees dataset is:



Assignment 5: Constructing Datasets and Using Databases

Due: 2020-10-11

there will be no tolerance of late submissions on this assignment because it will be graded Monday, so that you can use that feedback to finish up your portfolio.

Accept the [assignment](#)

Instructions

this will earn level 2 for construct

Your goal is to build and prepare two ready to analyze datasets. You will submit a notebook that describes how you built and prepared each dataset.

- Each finished dataset must be produced from two or more DataFrames.
- At least one must come from an sqlite database, either by merging results from multiple queries or multiple tables.
- You should use at least two different merges and one concatenate

Your completed datasets should have:

- column names that are well formatted (only lowercase letters, numbers and `_`)
- an added column that is derived from one or more other columns (string operation or calculation)

For each dataset, pose one question that could not be directly answered from the input data files as provided and demonstrate how to answer it with the dataset you built. This could be something that can be answered with using only `shape` of the merged data, but if you need summarize and visualize level 2 achievements, you should use more statistics and plots.

Your notebooks must be in the top level of the repository, not in a subfolder.

Additional Achievements:

if you already earned prior achievements you can ignore the following

To earn level 2 for prepare, one of your analyses must use datasets with missing values and one must be provided as excel files with merged columns (for example from [NCES](#)). You may use one dataset with both merged columns and missing data or one of each. You must also use datasets that have column names that need repair.

For the merged column data, either before or after merging, you must additionally:

- create separate tables for original and aggregate values (eg percentages or sums that can be recovered from the other columns)
- unstack all levels of the data to create a single level index over the columns

To earn level 2 for summarize and/or visualize, include additional analyses after building the datasets. You'll need at least two types of plots for visualize and/or two different ways of using summary statistics for summarize and to interpret the results for either.

Assignment 6: Naive Bayes

Due: 2020-10-20

Remember, even if you aren't sure how to do every part of the assignment, submitting pseudocode, a list of your questions indicating where you got stuck, or a partial solution will get you personalized feedback. You might even earn level 1 achievements for the partial effort.

For Classification Level 2:

[Accept the Assignment](#)

Create one notebook where you load examine each of the provided datasets for for suitability to use with Gaussian Naive Bayes. Label the sections **Dataset #**. Use exploratory data analysis (visualisation and statistics) in pandas and Gaussian Naive Bayes from scikit learn to answer the following for each dataset:

1. Do you expect Gaussian Naive Bayes to work well on this dataset, why or why not? (think about the assumptions of naive bayes and classification in general) *explanation is essential here, because you can actually use the classifier to check*
2. How well does a Gaussian Naive Bayes classifier work on this dataset? Do you think a different classifier might work better or do you think this data cannot be predicted any better than this?
3. How does the actual performance compare to your prediction? If it performs much better or much worse than you expected, what might you use to figure out why? (*you do not have to figure out why your predictions were not correct, just list tools you've learned in class that might help you figure that out*)

This assignment will be easiest if you take advantage of the template via git:

Git Workflow

1. Click the green code button and copy the url.
2. clone the repository as below where the url is the one you copied from GitHub. In your terminal on Linux or Mac or on the GitBash on Windows ([install instructions on tools section of syllabus](#))

```
cd path/where/you/want/a/new/folder/for/this/assignment  
git clone http://github.com/rhodyprog4ds/06-classification-username
```

1. then launch your notebook from the newly created folder.

on Linux or Mac, in the same terminal (remember you can use tab complete)

```
cd 06-classification  
jupyter notebook
```

or on windows, in your Anaconda prompt

```
cd path/where/you/want/a/new/folder/for/this/assignment/06-classification-username  
jupyter notebook
```

1. work on your assignment, by opening the notebooks there and editing them.
2. commit your changes when you want to save a point in your progress. Either you have part working and want to save that, or you want feedback, or you're done. On whichever terminal you used for the **git clone** command above

```
git add .
git commit -m 'description of your changes since last commit'
```

1. push your changes when you want to share them on GitHub, (eg need help, want feedback or complete)

```
git push
```

1. if you will keep working after pushing, first pull down the .md conversion that was added by GitHub actions. If you don't you'll have to merge, it should be fine, but ask if you're not sure.

```
git pull
```

For construct level 2

Build your own dataset for classification by merging data from separate sources that you're interested in.

Summarize and Viz level 2

Show some extra summary stats and plots

Assignment 7: Decision Trees

Due: 2020-10-27

Decision Trees

[Accept this repo for submission](#)

Choose a datasets that is well suited for classification and that has all numerical features. (eg the Wisconsin Breast Cancer data from UCI)

Part 1: DT Basics

1. Include a basic description of the data(what the features are)
2. Write your own description of what the classification task is and why a decision tree is a reasonable model to try for this data.
3. Include one summary visualization of the data.
4. Fit a decision tree with the default parameters on 50% of the data
5. Test it on 50% held out data and generate a classification report
6. Inspect the model by visualizing and interpreting the results
 - Does this model make sense?
 - Are there any leaves that are very small?
 - Is this an interpretable number of levels?
1. Repeat with the entropy criterion. Does using the entropy criterion make a big difference or small difference in the overall classifier?

💡 Tip

See the documentation for the [sklearn.DecisionTreeClassifier](#). One of the parameters is called `criterion`

Part 2: DT parameters

Do an experiment to see how `max_depth`, `min_values_split`, or `min_values_leaf` impacts the model.

1. Choose one of these and say explain why and how you hypothesize it will impact the performance
2. Use the model you fit above and EDA to choose minimum and maximum values for your parameter. Choose a total of 3 values for the parameter.
3. Retrain the model for each value of the parameter
4. Test and use at least 3 metrics to describe the performance, compiling your results into a DataFrame
5. Plot and interpret your results

you should use a loop for this part

Part 3: Test and Train Sizes

Do an experiment to compare test set size vs performance:

1. Train a decision tree on 20%, 30%, ..., 80% of the data, using one of the training parameter combinations you tried above and explain why you chose the one you chose.
2. Save the results of both test and train accuracy for each size training data in a DataFrame with columns ['train_pct','n_train_samples','n_test_samples','train_acc','test_acc']
3. Plot the accuracies vs training percentage.
4. Explain these results. What is the best test/train split. Why?

you should use a loop for this part

Evaluation

This assignment's focus is Classification and Evaluation:

- part 1 will show that you understand classification and can apply it for level 2
- parts 2 & 3 are opportunities to earn level 2 for evaluation.
- If you don't successfully complete the whole assignment, pseudocode or partial answers to the questions could earn you level 1 for either skill.

Tip

Start thinking about level 3 for evaluate by trying this experiment with cross validation.

Additionally:

- Using functions, loops, and conditionals appropriately while completing those tasks could earn level 2 for python if needed.
- You can earn level 2 for viz, summarize, prepare, or construct by choosing a messy dataset and/or including extra exploratory data analysis.

FAQ

How do I find a good dataset?

Look for a dataset with numerical features and a categorical target variable.

If you look at the [UCI website](#) you can search for datasets for Classification and numerical and look through what those search filters give you some options. You might want to choose a dataset with less than about 10 attributes to make your decision tree readable. To make the training fast, try to find a dataset with 1000 samples or less.

Assignment 8: Linear Regression

Due: 2020-11-03

Linear Regression

Find a dataset suitable for regression. We recommend a dataset from the UCI repository (see below)

Fit a linear regression model, measure the fit with two metrics, and make a plot that helps visualize the result.

Examine the [coefficients](#) or the residuals to try to interpret the results and explain what this regression model is doing.

Try fitting the model only on one feature. Justify your choice of feature based on the results above. Plot this result.

[Accept the assignment](#) to create your submission repository

Part 2: Test Train Splits

if you successfully completed this experiment in assignment 7, you don't need to repeat it, but it might be interesting to try

Do an experiment to compare test set size vs performance:

1. Train a the linear regression model on 20%, 30%, ..., 80% of the data.

2. Save the results of both test and train accuracy for each size training data in a DataFrame with columns

```
['train_pct','n_train_samples','n_test_samples','train_acc','test_acc']
```

3. Plot the accuracies vs training percentage.

4. Explain these results. What is the best test/train split. Why?

Part 3: Other models

Try fitting LASSO and explaining what it does. Does LASSO make better predictions on your data?

Do you think a model more complex than linear would be better? How could you tell?

Grading

Include description of what you're doing, why you're doing it, and what the results mean at each step of your analysis so that we can tell that you understand.

For regression level 2, complete part 1.

For evaluate level 2, complete part 2.

If you're curious, try a more complex regression model as in part 3. This will get you an early start on level 3 for regression.

If you don't successfully complete the whole assignment, pseudocode or partial answers to the questions could earn you level 1 for either skill.

FAQ

How do I find a good dataset?

Look for a dataset with numerical features and a categorical target variable.

If you look at the [UCI website](#) you can search for datasets for Classification and numerical and look through what those search filters give you some options. You might want to choose a dataset with less than about 20 attributes. To make the training fast, try to find a dataset with 1000 samples or less, or use read functions to use only a small chunk of the data. Remember to read about the dataset and note what you're predicting.

Assignment 9

Due: 2020-11-010

[assiginemnt submission](#)

1. Apply Kmeans to the dataset you used for assignment 7.

2. Evaluate how well clustering worked on the data:

- using clustering metrics and
- using visualization.
- the ground truth labels

1. Include a discussion of your results that addresses the following:

- describes what the clustering means
- what the metrics show
- Does this clustering work better or worse than expected based on the classification performance (if you didn't complete assignment 7, also apply a classifier)

Evaluation and Grading

The goal of this is for clustering level 2.

You may also earn evaluation level 2. To do this, use at least two metrics, explain them thoroughly, and use train-test splits.

You can earn visualize and summarize by adding and interpreting visualizations and summary statistics.

As always, psuedo code or an outline can earn level 1

Portfolio

This section of the site has a set of portfolio prompts and this page has instructions for portfolio submissions.

Starting in week 3 it is recommended that you spend some time each week working on items for your portfolio, that way when it's time to submit you only have a little bit to add before submission.

The portfolio is your only chance to earn Level 3 achievements, however, if you have not earned a level 2 for any of the skills in a given check, you could earn level 2 then instead. The prompts provide a starting point, but remember that to earn achievements, you'll be evaluated by the rubric. You can see the full rubric for all portfolios in the [syllabus](#). Your portfolio is also an opportunity to be creative, explore things, and answer your own questions that we haven't answered in class to dig deeper on the topics we're covering. Use the feedback you get on assignments to inspire your portfolio.

Each submission should include an introduction and a number of 'chapters'. The grade will be based on both that you demonstrate skills through your chapters that are inspired by the prompts and that your summary demonstrates that you know you learned the skills. See the [formatting tips](#) for advice on how to structure files.

The second submission will be graded on the following criteria and due on November 13:

Level 3

keyword	
python	reliable, efficient, pythonic code that consistently adheres to pep8
process	Compare different ways that data science can facilitate decision making
access	access data from both common and uncommon formats and identify best practices for formats in different contexts
construct	merge data that is not automatically aligned
summarize	Compute and interpret various summary statistics of subsets of data
visualize	generate complex plots with pandas and plotting libraries and customize with matplotlib or additional parameters
prepare	apply data reshaping, cleaning, and filtering manipulations reliably and correctly by assessing data as received
classification	fit and apply classification models and select appropriate classification models for different contexts
regression	can fit and explain nonlinear regression
clustering	apply multiple clustering techniques, and interpret results
evaluate	Evaluate a model with multiple metrics and cross validation

On each chapter(for a file) of your portfolio, you should identify which skills by their keyword, you are applying.

You can view a (fake) example [in this repository](#) as a [pdf](#) or as a [rendered website](#)

For this portfolio check I encourage you to dig in on either data from one domain and think about how clustering, regression, and classification apply in this domain OR to inspect these more carefully in a performance prospective, taking a more CS-basics approach.

Formatting Tips

Your portfolio is a [jupyter book](#). This means a few things:

- it uses [myst markdown](#)
- it will run and compile Jupyter notebooks

This page will cover a few basic tips.

Organization

The summary of for the **part** or whole submission, should match the skills to the chapters. Which prompt you're addressing is not important, the prompts are a *starting point* not the end goal of your portfolio.

Data Files

Also note that for your portfolio to build, you will have to:

- include the data files in the repository and use a relative path OR
- load via url

using a full local path **will not work** and will render your portfolio unreadable.

Structure of plain markdown

Use a heading like this:

```
# Heading of page
```

in the file and it will appear in the sidebar.

File Naming

It is best practice to name files without spaces. Each **chapter** or file should have a descriptive file name (**with_no_spaces**) and descriptive title for it.

Syncing markdown and ipynb files

To sync feedback received to your runnable notebook files, change the related GitHub Actions file: [.github/workflows/](#) In the step named convert that looks like:

```
- name: convert
  run: |
    jupytext */*.ipynb --to myst
```

change it to:

```
- name: convert
  run: |
    jupytext --set-formats ipynb,md */*.ipynb # Turn .ipynb into a paired ipynb/py notebook
    jupytext --sync */*.ipynb # Update whichever of .ipynb/notebook.md is
    outdated
```

This means if you accept suggestion commits from the the **.md** file, the action will update your **.ipynb** file. If you update your **.ipynb** file the action will update the **.md** file.

Adding annotations with formatting or margin notes

You can either install [jupytext](#) and convert locally or upload /push a notebook to your repository and let GitHub convert. Then edit the **.md** file with a [text editor](#) of your choice. You can run by uploading if you don't have jupytext installed, or locally if you have installed jupytext or jupyterbook.

In your **.md** file use backticks to mark [special content blocks](#)

```
```{note}
Here is a note!
````
```

```
```{warning}
Here is a warning!
````
```

```
```{tip}
Here is a tip!
````
```

```
```{margin}
Here is a margin note!
```
```

For a complete list of options, see [the sphinx-book-theme documentation](#).

Links

Markdown syntax for links

```
[text to show](pathor/url)
```

Configurations

Things like the menus and links at the top are controlled as `settings`, in `_config.yml`. The following are some things that you might change in your configuration file.

Show errors and continue

To show errors and continue running the rest, add the following to your configuration file:

```
# Execution settings
execute:
    allow_errors : true
```

Using additional packages

You'll have to add any additional packages you use (beyond pandas and seaborn) to the `requirements.txt` file in your portfolio.

Reflective Prompts

These prompts are more reflective to help demonstrate your understanding of skills. These are more writing than new coding.

Correcting a Prior Assignment

Python Process Access Summarize Visualize Prepare Construct Classification Clustering Regression Evaluation

Yes No Yes Yes Yes Yes Yes Yes Yes Yes

Table 1 Eligible Skills

Choose an assignment that you did not achieve the target level for. Write a blog style notebook analysis that corrects what you could have done better, what you learned, and addresses the misconception if applicable.

Data Science Pipeline

Like the day 1 activity, find two different sources that describe the data science pipeline or lifecycle. Write a blog style post that discusses their differences and hypothesizes about why they may be different? Are they for different audiences? Is one domain specific. How do they emphasize different modeling tasks?

Podcast

Python Process Access Summarize Visualize Prepare Construct Classification Clustering Regression Evaluation

No Yes, L2 No No No No No No

Table 2 Eligible Skills

Watch an episode of a high quality[\[1\]](#) podcast and write a blog style summary and review of the episode. Highlight what you learned and how it relates to topics covered in class.

Approved Podcasts:

- [Pod of Asclepius, Fall Series: The Philosophy of Data Science](#)

What Can Data Science do?

Write a blog post describing different ways

[\[1\]](#) approved by Dr. Brown by creating a pull request to add it to the list on this page that is successfully merged. To create a PR, use the suggest an edit button at the top of this page.

Analysis Prompts

Loading Data

| Python | Process | Access | Summarize | Visualize | Prepare | Construct |
|--------|---------|--------|-----------|-----------|---------|-----------|
| No | No | Yes | No | No | No | No |

Table 3 Eligible Skills

Look at all of the different ways pandas can load data. Consider some of the questions below and add a notebook that's styled like a report that answers a few of them with text and code.

- Which seem like good idea? Are any dangerous?
- Which seem more or less common?
- Can you compare them on speed? Is it ever worth transforming a dataset before loading?
- How much can you repair a dataset using the parameters of the load functions?

CheatSheet

| Python | Process | Access | Summarize | Visualize | Prepare | Construct | Classification | Clustering | Regression | Evaluation |
|--------|---------|--------|-----------|-----------|---------|-----------|----------------|------------|------------|------------|
| No | No | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes |

Table 4 Eligible Skills

Make a cheatsheet with examples of the several different parameter settings for common operations for one topic.

This [cheatsheet](#) is an example, it's too broad, but it's the same idea. Yours should

Deeper Analysis

| Python | Process | Access | Summarize | Visualize | Prepare | Construct | Classification | Clustering | Regression | Evaluation |
|--------|---------|--------|-----------|-----------|---------|-----------|----------------|------------|------------|------------|
| Yes | No | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes | Yes |

Table 5 Eligible Skills

For one of the assignments, if there was something you were curious about. Try it out and investigate how to answer it. Vary parameters and document your investigation.

New Analysis

Python Process Access Summarize Visualize Prepare Construct Classification Clustering Regression Evaluation

Yes No Yes Yes Yes Yes Yes Yes Yes Yes

Table 6 Eligible Skills

For one a topic of interest, clean, explore and model the data. Work with messy data or data provided in multiple files to earn prepare and construct achievements or use clean data to earn only EDA and modeling achievements.

Modeling Experiments

Python Process Access Summarize Visualize Prepare Construct Classification Clustering Regression Evaluation

No No Yes Yes Yes Yes Yes Yes Yes Yes

Table 7 Eligible Skills

FAQ

This section will grow as questions are asked and new content is introduced to the site. You can submit questions:

- via e-mail to Dr. Brown (brownsarahm) or Beibhinn (beibhinn)
- via Prismia.chat during class
- by creating an [issue](#)

Syllabus FAQ

How much does assignment x, class participation, or a portfolio check weigh in my grade?



Can I submit this assignment late if ...?



GitHub FAQ

My portfolio won't compile



Help! I accidentally merged the Feedback Pull Request before my assignment was graded



Common Debugging Issues

Key Error



Resources

This section will compile resources for the course over time into various sections:

- [Data](#) sources of data to use for assignments
- [Programming](#) info on the tools and libraries we're using in class, background info, etc
- Reference example tips
- [Tips](#) general, semi-related information
- [Cheatsheet](#) patterns and checkpoints

General Tips and Resources

This section is for materials that are not specific to this course, but are likely useful. They are not generally required readings or installs, but are options or advice I provide frequently.

on email

- [how to e-mail professors](#)

References on Python

- [Course Text](#)

Data Sources

- [Kaggle](#)
- [Google Dataset Search](#)
- [UCI Data Repository](#)
- [Json Datasets](#)
- [Databases](#)

If you have others please share by creating a pull request or issue on this repo (from the GitHub logo at the top right).

By Professor Sarah M Brown
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