Text Analytics and Natural Language Processing in Finance and Fintech

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Introduction

1.1 Why Is This Course Special?

This course is special because to the best of my knowledge, it was developed as the **first course worldwide** that covers text analytics and natural language processing (NLP) in finance. There are courses in NLP and courses in finance. But none of them cover how both of them work together and interact. This is increasingly becoming important, not only with the rise of fintech, but also in many other fields such as investment management, where often vast amounts of so-called alternative data is used, e.g. text from social media, company filings, or the financial press.

Another reason why this course is special is because it is **project-focused**. Students throughout the course work on projects of their own choosing where they write code and analyze text in a financial setting.

Finally, there are many resources available that explain isolated issues in NLP, but it is very difficult to find a coherent approach. Beginners are mostly overwhelmed by the material they can find in various places, and don't know how to tie everything together in a way that makes sense. This course fills this gap by showing you how to develop a **consistent and well thought-out workflow** for text analytics in finance, see for example Chapter 8 starting on page 43.

1.2 Why Are Text Analytics and NLP Useful in Finance?

Why are text analytics and natural language processing (NLP) useful in finance? I briefly would like to provide two examples. The first is that you can use NLP in investment management. For example, you could analyze financial media, social media (e.g. Twitter), or company reports to make better investment decisions and ultimately get higher investment returns. Second, you could for example analyze the minutes of the Federal Reserve and calculate the probability of a raise interest rates, which is a trillion-dollar question.

For clarification, in this book we focus on the analysis of existing text, as this is a common use case in financial applications. In particular, we do not spend a lot of time on

natural language generation or machine translation.

1.3 Important Announcements via HKU Email

Each student in this course is **expected to check his/her HKU email on a daily basis**. The reason is that this is the way the instructor might contact students. Furthermore, important announcements will be made on Moodle, and these announcements will not be repeated in class. These announcements will be forwarded to your HKU email address only. If a student misses important information or announcements sent to the HKU email address, the student will have to bear the consequences, which may include but are not limited to receiving a lower grade in this course, failing this course, or delaying/endangering the graduation.

1.4 Course Logistics

- As Lev Konstantinovskiy has succinctly pointed out, "NLP is 80% preprocessing." So please keep in mind that NLP is mostly grunt work. This is a course about getting your "hands dirty" with NLP, not about a fancy marketing brochure about how fashionable NLP is.
- Furthermore, it is very important to set up the right workflow and use the right tooling. While we talk a lot about text analytics and NLP, it is in my opinion at least **equally important to get up to speed with the right tools** to use, so we will spend a fair amount of time on that. We discuss more of this in Chapter 4 starting on page 23.
- Text analytics and NLP is a very applied topic that is learned best by actually doing it. In analogy, you cannot learn to play the piano just by reading books. You need to actually do it and play the piano. That is why this course will put emphasis on students working on their own projects and presenting them in class.
- The instructor can point you into the right direction, help you to understand the key concepts, tell you which packages to use (these choices are very important as they can make or break your project), and provide examples. All of these are things you often cannot easily find online. Furthermore, the lecture notes and website contain many code examples.
- But for specific programming problems during your projects you should find out yourself how to write your code. Most (if not all) the packages we use are very well documented (e.g. on ReadTheDocs.org or on their own website) and surrounded by an active community (e.g. on StackOverflow.com, blog posts, etc.), so please make good use of it.

- We are mostly going to focus on analyzing **English** text. However, many of the
 methods we discuss also work with other languages, e.g. **Chinese**, sometimes with
 modifications.
- If anything in this document is written in italics (*like this*), it means that either I want to emphasize something, or it is a new concept mentioned/defined for the first time. It will be clear from the context which one it is, emphasize or define. Sometimes I will also write something in bold (**like this**) instead of using italics.

1.5 Links

- The electronic PDF version of these lecture notes contains clickable links in blue color. If you happen to have a printed version, feel free to try the electronic version of the lecture notes as they might be more convenient.
- Furthermore, reference to different chapters/sections and their respective pages are also clickable in the PDF version and will take you directly to the relevant page.
- Why do many links in these lecture notes point to GitHub? Because that is where
 the original source code lives and the GitHub repos usually also contain brief introductions and pointers to other webpages. Usually you can find a link to the main
 website at the top of the GitHub page or in the README file that is displayed after
 the code listing.

1.6 Important Deadlines and Dates

- The group leader should hand in the group information to the instructor at the latest on the day after the add/drop period. For details see Section 2.4 starting on page 17.
- The **midterm** is tentatively scheduled in-class for **Saturday, March 5, 2022**. The format and date of the midterm is subject to change due to the ongoing Covid-19 pandemic.
- Blog post(s): See Sections 2.2 and 2.4 starting on pages 13 and 17, respectively.
- Presentation dates: See Section 2.3 starting on page 15.
- Group project: See Section 2.4 starting on page 17.

Group Project

Throughout this course, you will work on projects related to text analytics and NLP in finance. This chapter contains detailed instructions and suggestions about the group projects.

2.1 Version Control and Collaborative Coding

- For more information on version control see Chapter 5 starting on page 31.
- Using version control for your project is **optional and not evaluated**. However, it might make your life easier in the long run if you put your code under version control and host it on GitHub.
- If you would like to use version control and if you do not already have a GitHub account, please sign up now. Please use sensible names for your GitHub user name and your GitHub repositories (or "repos" in short).

2.2 Blogging

2.2.1 Instructions

- From Wikipedia: "A blog (a truncation of the expression "weblog") is a discussion or informational website published on the World Wide Web consisting of discrete, often informal diary-style text entries (posts)."
- Each group should write one or several blog post(s) about the project the group is working on. The blog post(s) can be written throughout the course (recommended) or at the end of the course (at the latest before the group project is handed in).
- The topic of the blog may be about the work you do for your group project. However, the blog's content should be significantly different from the content of your presentation slides and group report, in other words, it should show a different perspective.

- The blog should be **reflective** in nature. For example, you can describe some technical or conceptual problems you encounter, and describe your journey and how you solved (or intend to solve) these problems.
- One reason why blogging is a good idea is that, similar to posting your code on GitHub, it will increase your visibility and your chances on the job market. Blogging really is community engagement. Its usefulness cannot be overstated in establishing yourself as a thought-leader.
- It is not required, but feel free to mention your names on the blog if you like (e.g. for promoting yourself to potential future employers). Also feel free to mention the course and the instructor's name on the blog post if you like (optional). But my recommendation is to NOT include your university student numbers on your blog for privacy reasons.
- The **evaluation criteria** are that the blog post
 - should be of high quality in terms of content, writing style, and execution,
 - should contain a clear description of what's going on and why it is important,
 - should include well-documented code snippets with explanations that illustrates key things about your project (any code posted should be self-contained and be fully reproducible in the sense that anyone reading the blog post can run the code and verify the results), and
 - should have a word count of at least 700 (source code is not included in the word count).
 - If you prefer to split up the description of your work across several smaller blog posts (instead of one big blog post) it is also fine. The evaluation of the blog is based on the blog's overall content. It is not based on the number of posts.
- While your blog should look professional, it is important that the **content** is of high quality in every conceivable way. It's not necessary to have a very fancy styling in terms of layout. A minimalistic layout is often a good idea if you want to focus on the written bog content and your code.
- Once you have written a new blog post, please **notify the instructor by email** and announce your blog post within three days of its publication on the **online forum of the course webpage** so that your classmates can take a look as well.

2.2.2 Blogging Software

• There is a blog website for this course, where students can publish their blog articles. It is available at buehlmaier.github.io/MFIN7036-student-blog-2022-02

- We use Pelican, which is a static website generator written in Python. You can find usage instructions for Pelican at docs.getpelican.com.
- The workflow for creating a new blog post is as follows (no knowledge of Git is required):
 - 1. Download the source code of the blog at buehlmaier/MFIN7036-student-blog-2022-02 (click on "Code" and then click on "Download ZIP")
 - 2. Write a first draft of your blog post in a Markdown file (e.g. myblogpost.md) and add it to the content directory; the easiest way to generate this file is to make a copy of the demo-blog-post.md file, rename it to myblogpost.md, and start modifying its content
 - 3. Open a terminal in the parent directory of the content directory
 - 4. Type the following command in the terminal:

```
pelican content -t themes/elegant-5.4.0 --autoreload --listen
```

- 5. Point your web browser to http://127.0.0.1:8000
- 6. You can make changes to your markdown blog file (e.g. myblogpost.md as mentioned previously) and then reload your browser by pressing F5. Repeat this step as often as you like
- 7. Once you are satisfied with your blog post, you can email the file(s) to the course instructor/TA for upload to the blog. The file(s) should include the markdown file of the blog post and potential additional files that are included in your blog post, e.g. picture(s)
- Please keep in mind that if a blog post is broken in any way (i.e. does not work with the existing source code of the blog) or is of insufficient quality, the course instructor/TA will refuse to put the blog post on the blog website and this blog post will not be counted towards the evaluation, i.e. the student(s) will receive zero points for this blog post.

2.3 Presentations

2.3.1 General Points

• When conducting your presentations, please always leave enough time for discussion, questions, and the instructor's feedback. Based on previous experiences, around 20%-25% of the presentation time should be set aside for this purpose. For example, if your presentation is scheduled to last for 20 minutes, you should prepare for around 15 minutes presentation and 5 minutes of discussion, questions, and feedback. Students lose 5% of the overall presentation grade for each minute overtime (on the group level). No points will be deducted for presentations that finish a bit early.

- The **durations of the group presentations** will be announced on the course website in due time. The reason why we cannot announce them now is that before the add/drop period, we do not know how many students will take the course and therefore how many groups we have in this course.
- All **presentations will be evaluated** based on the following criteria:
 - Confidence of the speaker. Is he comfortable, can easily connect with the audience, make eye contact, and put the audience at ease?
 - Quality of the information presented. Enough details to support the main point, but not too many unnecessary details that confuse or distract.
 - Level of clarity. Should easily be able to convey the main point, vocabulary should be easy to understand and all words should be spoken in a clear and fluent manner.
 - Level of organization. A good presentation should have structure and organization, including a proper introduction and conclusion.
- The group presentations should consist of going through slides and sometimes through short software demonstrations (showing the group's progress or results) on a student's laptop connected to the projector.
- Presentations will be evaluated on an individual basis according to the syllabus. Although it is encouraged that students present frequently, it is not required that each group member presents during each presentation. But each student should present at least one time during the whole course. For each student, the overall presentation grade will be the average of all presentations done by this student. The decision about which students present during which presentation should be made within the group.
- During a presentation, please do not switch back and forth between students; if a student is done with his/her part, he/she should not come back later during the same presentation. (Of course, a student can come back to present in a subsequent presentation on another date.)

2.3.2 Presentation Dates

- We will have group presentations in **classes 7 and 12 (the last class)**. On each of these dates, all groups should present.
- In the presentation in **class 7**, you start by laying out **your plan** (i.e. the roadmap and schedule) of the group project for the rest of the course. See also Section 8.1 starting on page 43.
 - What are the goals you would like to accomplish?
 - Why are they important?

- What are the sub-tasks you need to do to accomplish your goals and what is the timeline for each sub-task? (You can show a Gantt chart for example.)
- Who is responsible for which task?
- Optional: What have you done so far to accomplish the first sub-tasks, are there any partial results you can show?
- Keep in mind that the plan is not set in stone. Think of your project as an iterative process. Most likely it will be necessary to go through several rounds of updates and changes. But it is still important to have a plan, even if it needs to be modified during the course of your project.
- The presentation in **class 12** is the final presentation where you tie everything together and summarize what you have accomplished throughout the course.
- Important **potential exception to the timing**: In case of a large number of students enrolled in this class and/or a large number of groups, we might spread the second group presentations over the last two classes of the course (instead of having the second group presentation in the last class only). An announcement will be made in due course.

2.4 Group Project General Points

- A central element of the project should be the analysis of textual data. Often you may want to include structured data as well, e.g. company fundamentals, macroeconomic data, etc., but the main focus of the project should be on textual data.
- Probably you have many questions where to get the data from. In Chapter 6 starting on page 33 there is a list with many potential **data sources**. Furthermore, there are a lot of **application examples of text analytics and NLP in finance** in Chapter 7 starting on page 37.
- For the data sources and project ideas, keep in mind that these are only suggestions. If you have a great idea or a new dataset/source you would like to analyze, feel free to pursue it further. However, I strongly recommend consulting with me first to find out whether this project is feasible and suitable, as there are often "hidden" challenges that are difficult to identify in advance if you have not worked on text analytics and NLP before.
- Students should use a database that is freely available or a database subscribed to by the University. **Students should not have to pay for any database used in the group project.** For a list of potential data sources please see Chapter 6 starting on page 33.
- For the group project, form groups as soon as possible, nominate a group leader, and email a spreadsheet to the instructor (at the latest on the day after the add/drop period) containing the following information:

- Group name (please choose a professional and sensible name)
- Student ID numbers
- Last names
- First names
- English first names (if any)
- GitHub user names (if any)
- Preferred presentation date of second group presentation: Either the second-to-last class or the last class of this course. (This is only valid if there is a large number of students and/or groups, see Section 2.3.2. Usually the second presentation would take place in the last class. Note: We will try to accommodate the presentation date preferences as far as possible, first-come, first-served, but we cannot guarantee to fulfill the preferences of everyone.)

Next to the group leader's name put a label so that it's easy to identify him/her from the spreadsheet. In case several groups choose the same group name, it is first-come, first-served (in other words, the group who handed in later has to change their group name).

- The accepted group size is 3-5 students per group. You may form your groups already at the beginning of the course, but in general you might want to wait a bit with the group formation until students have committed to taking this course (at the latest until the end of the add/drop period). In terms of group size, you should try to strike a balance depending on the strengths and weaknesses of your group members. If you have too few students, each student will have more work to do, and no "bonus" in terms of grading will be given if you choose this arrangement. And if you have too many students, it might be difficult to coordinate the group work. Furthermore, you might not have enough time in your group presentations for everyone to present, which might have a negative impact on the evaluation of your presentation in case you have to rush and/or don't have enough time to get your point across or answer questions.
- To manage the workload efficiently, I suggest to split up your team by **assigning specific responsibilities from the start**. Whether and how you split up the workload is up to you, but you might want to orient yourself at the list from Section 8.1 starting on page 43.
- Please choose a project based on **textual analysis of the English language only**. The reason is simply that most software libraries are originally designed with English in mind. Furthermore, to ensure fairness to all students, it might sometimes be difficult to evaluate projects based on text other than English.
- When choosing a project, please keep in mind that it is often not easy to predict market prices in the future. So while it is tempting to work on a project that potentially could make a lot of money, you should have a **backup plan** if you fail to

find predictive power for prices. For example, you could look at predicting volatility. Or you could focus on other variables that are of financial interest, e.g. predicting accounting fraud or others.

- In general, it is a good idea to **start simple**. The "simple" ideas in NLP are often already difficult enough to implement. Once you have finished the "simple" analysis, you still have ample opportunities to make it more complex.
- If some of your attempts do not work out (e.g. you cannot find predictive power for the question you're trying to analyze such as predicting whether the FED is going to raise rates), you should still describe what you did, what problems you encountered, and your attempts to solve these problem, even if those attempts were at the end not successful. What matters mostly is whether you followed proper process, not so much whether your hypothesis finds support in the data in the end.
- All files created with an office suite such as LibreOffice (Linux), iWork (Apple), or Office (Windows), with the possible exception of spreadsheets, should be converted and **submitted in PDF format** to ensure consistent behavior across different platforms. Every document that should be submitted in PDF format but is submitted in another format results in a **deduction of 1% of the overall course grade for all students in the group**.
- Source code files, Jupyter notebooks, and data files should stay in their original format and do not have to be converted to PDF.
- If you use GitHub, the programming code and data should be in repo(s) of the group leader. The group leader should give write access to the repo(s) to all group members who contribute source code or collect data. Be careful not to mess things up when several people contribute to the same codebase. See also Section 2.1 starting on page 13.
- You should also hand in a project report summarizing what you have done in your group project. The cover page should contain all the names and UIDs of your group members as well as the name of your group.
- The project report should not only describe technical aspects of your work, but also provide information about the economic or financial question you are addressing and explain why your topic is important.
- Furthermore, there should be a separate "who did what" document that describes in half a page the way you split the work. Please **indicate the percentage of work contributed by each team member at the end of the project**. For example, if a group member was not a free-rider, then his/her percentage would be 100%, meaning that the group member contributed 100% of what was agreed before when the work allocation was determined by the group. If no percentage is given, all group members are assumed to have contributed equally (independent of the written description of the work done by each team member). **If a free-rider is identified**

according to the percentage listed on the "who did what" document, the free-rider who contribute sufficiently less than their group mates will receive proportionately fewer points in the evaluation of the group project.

- Finally, the slides of all group presentations held should also be handed in together with the project report.
- In summary, the group project handed in should consist of the following files (combined in .zip compression format), handed in at the latest at 11:59pm on the seventh day after the last class of this course in one email to the instructor and TA(s) with the email subject "Group Project Submission:"
 - 1. Project report (PDF format, max. five pages excluding cover page; the page limit includes the appendix, if any; a references section, if any, can be on an additional sixth page)
 - 2. "Who did what" document (PDF format, half a page)
 - 3. Slides of all presentations held (PDF format)
 - 4. A list (PDF format) containing links to all blog posts; links should be clickable
 - 5. Printouts (PDF format) of all blog posts
 - 6. Code files
 - 7. Data files necessary to run the code; if the size of the data is larger than one megabyte (MB), you may provide a link to a file sharing service where the data is stored online, e.g. Git Large File Storage, Dropbox, Google Drive, etc.; if a file sharing service is used, it should be used for data files only, and all other files should be submitted (combined in .zip compression format) by email as as mentioned above
 - 8. Supplementary files, e.g. Jupyter notebooks (this is optional)
- The group projects are evaluated on the group level (unless some free-riders are identified, in which case adjustments will be made accordingly in the evaluation of the group report as mentioned above). So unless there is a free-rider detected, all students receive the same points for the group project (with the exception of the in-class group project presentations, which are evaluated on the individual level according to the syllabus).
- The **group project is evaluated** according to the following criteria:
 - 1. Quality of economic/financial idea
 - 2. Execution of economic/financial idea
 - 3. Quality of programming code

Please keep in mind that the blog post(s) are evaluated separately as described in Section 2.2 starting on page 13. Moreover, while the presentation slides are part of the group project, the presentation skills of each student are also evaluated separately as described in Section 2.3 starting on page 15.

Common Terms/Abbreviations and their Explanations

This chapter is meant to be used as a reference. When reading the lecture notes you come across a term you don't fully understand, come back here to look it up.

- NLP: Natural language processing.
- AI: Artificial intelligence.
- Semantics tries to capture the intrinsic meaning of natural language.
- Valence: Often it just refers to whether a word is positive or negative.
- "Parsing" refers to the process of analyzing a string of symbols conforming to the rules of a formal grammar.
- "Tokenization" turns a string or document into tokens, i.e. smaller parts. These could be, for example, words, word combinations (e.g. n-grams), sentences, paragraphs, etc.
- Corpus (plural corpora) or text corpus: A large and structured set of texts, typically a set of text documents. You can think of a text document as a book and the corpus as a library containing many books.
- Term: It basically means "word," although it sometimes might refer to a more abstract word representation (e.g. a linear combination of words or an n-gram), see for example Section 14.6.
- BoW: Bag-of-words, see Section 14.5 and Chapter 15.
- Stop word: A word that occurs frequently in text but does not have significant meaning such as "the," "is," "at," "which," or "on." Often these words are removed when preprocessing text documents. See also Section 14.4 starting on page 98.
- Render: It basically means to "display" something, e.g. a browser renders a HTML document.

- API: Short for "application programming interface." It means that you can write a programm to access the application, e.g. to download some data. An example is the Twitter API that lets you download tweets programmatically instead of scraping the Twitter website (see Section 11.3 starting on page 73).
- Command line: This is a user interface navigated by typing commands at the prompt instead of using a mouse. The command line plays an important role in Unix/Linux because it can be very powerful and it lets you get a lot of things done that would otherwise be be difficult to do or would take a longer time (e.g. renaming files in batches, using version control, searching for files, etc.). See also the explanation in Chapter 4 starting on page 23.

Python Installation

Whenever you work on software projects, it is really crucial to use the right tooling. By "tooling" I mean all the software you use to write your own software. Using the right tooling can mean a great increase in productivity because you can get things done faster and with fewer errors. Using the right tools also means that you (hopefully) have a larger community around your software tools so you can get help and/or feedback much faster and more effectively.

4.1 Quick Summary

- This section (Section 4.1) should be all you need to get started with Python. However, if you need to dive in deeper, you can also take a look at the other sections in this chapter.
- Operating system: It is recommended to use Linux or macOS, although using Windows is also possible.
- To install the programming language Python, use Anaconda (click on the link and follow the instructions there).
- Once you have installed Anaconda, you can type commands into the Anaconda prompt (if you're on Windows) or in the command line (if you're on Linux or macOS). The following bullet points explain how you can type commands to install an IDE and various Python packages.
- To write Python code, use the Spyder integrated development environment (IDE). You can install Spyder with

```
conda install -c anaconda spyder
```

• You can install Python packages using conda, e.g. typing the following commands will install some of the fundamental Python packages required for this book:

```
conda install -c anaconda numpy
conda install -c anaconda scipy
conda install -c anaconda pandas
conda install -c anaconda scikit-learn
conda install -c anaconda tensorflow
conda install -c anaconda keras
conda install -c anaconda nltk
conda install -c anaconda spacy
conda install -c anaconda gensim
conda install -c anaconda requests
conda install -c anaconda bs4
conda install -c anaconda selenium
conda install -c anaconda seaborn
conda install -c anaconda bokeh
conda install -c anaconda dash
conda install -c conda-forge pelican
```

4.2 Operating System

- It is up to you which operating system you use in this class, but in general I recommend **Linux or macOS**. It is usually easier to set up a good programming environment in those operating systems. **Windows** is also allowed in this course, although its use is not encouraged.
- The popularity of Linux distributions can vary quite a bit as time goes by. Distrowatch has a popular ranking of Linux distributions based on page hits. Some of the most popular Linux distributions these days are MX Linux, Manjaro, and Mint. If you would like to use Linux and are not sure which Linux distribution to use, you can give one of them a try.
- As usual, **make a backup** of all your data before installing Linux.
- You can install Linux inside a **virtual machine** or as **dual boot**. The advantage of a virtual machine over dual boot is that it might be easier to set up and that you don't mess up your existing system if something fails in the Linux installation. But the disadvantage is that you'll have less memory available to run your code (this might or might not be an issue depending on what you want to do with Linux).
- If you install Linux, use a 64-bit Linux (not 32-bit, assuming that your laptop is not very old).
- You can also use macOS as an alternative to Linux in this course. macOS is a Unix-based operating system tracing its roots back to the original BSD (Berkeley Unix), so it has very similar philosophical roots as Linux.

- What is "root?" It is the name of the superuser. This is a special user account used for system administration. It is very powerful as there are almost no restrictions to what root can do on a computer. So use it with care.
- So **what is "sudo?"** It is a program on Unix-like systems (such as macOS or Linux) that allows you to run another program with the security privileges of another user, typically the superuser.
- What is a "command line" or "terminal" or "shell?" The command line is an ancient but very powerful way of interacting with your computer. It originated in the 1960s when people were working on computer terminals. The basic idea is that instead of using a mouse and clicking your way through the operating system, you type commands on a keyboard and look at the text output generated by your computer. It can be very powerful because you can combine various commands in ways that are difficult to replicate using a point-and-click interface. The command line can also be a powerful tool for renaming files in batches, using version control, searching for files, etc. You can even program most shells.

4.3 Working With Python

- You should use the <u>IPython</u> command shell. It is a powerful interactive shell with support for interactive data visualization and parallel computing. It also has a kernel for <u>Jupyter</u>, which is a notebook for Python allowing you to run code in.
- Unless mentioned explicitly otherwise, we use **Python 3** instead of Python 2.

4.4 Python Language Installation

You should use Python 3. Python version 3.0 was released in 2008. It made some backwards incompatible changes relative to Python 2 to solve some long-standing issues and design flaws in the language. This means you cannot run some code written for Python 2 in Python 3 and vice versa. For example:

```
print 'hello'  # Python 2
print('hello')  # Python 3
```

The following list shows how to install the Python programming language on all major platforms:

- Windows: Probably the easiest way to install Python and manage packages is with Anaconda.
- 2. **macOS**: Python 2 should be pre-installed on macOS, but you want Python 3 for this course. One alternative is Anaconda, which should work out of the box. Alternatively you can use Homebrew and type the following command in the terminal. You do not need to add sudo as brew itself is owned by root.

```
brew install python3
```

3. **Linux**: Anaconda should work, but in Linux, people often want to install software themselves from the command line using a Linux package manager such as apt or dnf. The Linux package manager depends on the Linux distribution you are using. For example, in a Debian-based system such as MX Linux, Mint, Ubuntu, or Debian you can type the following commands, which should install a set of Linux packages related to Python.

```
sudo apt-get update
sudo apt-get -y upgrade # This is optional. It will upgrade your system.
sudo apt-get install -y python3 python3-pip python3-venv
sudo apt-get install -y build-essential libssl-dev libffi-dev python-dev
```

4.5 Python Package Installation

• In general I recommend using the Anaconda Python distribution. It was created specifically with the needs of a data scientist in mind. It includes Python itself and hundreds of packages. It is relatively easy to set up, and uses Conda to manage the packages. There is also Miniconda, which is a bare-bones version of Anaconda containing only Python and Conda, so you can install whatever package you need from scratch. In any case, you typically install Anaconda as user, not as root. Anaconda is freemium open-source, which means that the core is open source, but money is charged for add-on services. For our purposes, we don't need these extra services, so Anaconda is free.

Please keep in mind that **if you end up using Anaconda**, **you should use Anaconda** (**not pip**) **to install Python packages**. The only reason you should use pip to install packages is if the package is not available through Anaconda. The usual way to install a package is by typing in the Anaconda prompt (if you're on Windows) or in the command line (if you're on Linux or macOS; no need to use sudo) the following command:

```
conda install package-name
```

You should replace package-name with the name of the package you would like to install, e.g. scipy.

• If you already have an existing Python installation (e.g. your Linux distribution usually should already have a system-wide Python installation in place), you can install packages directly with pip or Conda without using Anaconda. Unless you need to manage non-Python dependencies, it is usually **best to go with pip+virtualenv** (see Section 4.6 starting on page 27).

- pip (or pip3 for Python 3) is the historically older installer. Often people use pip together with virtualenv. The basic idea is that you use virtualenv to manage packages for different projects, e.g. you could be using different package versions for different projects by putting them in different environments. Virtualenv does not need root permissions, so you don't have to touch the global, system-wide Python installation. Alternatively, you could also run pip as root (in which case the package would be installed system-wide), but this practice is generally discouraged as it messes with your system-level Python installation. If there are any problems resulting from the package installation, it might be difficult to undo the changes you have made.
- Conda is the younger relative of pip. It is also open-source. In contrast to pip, it can also handle non-Python library dependencies. It is the package manager for the Anaconda Python distribution, but can also be used in isolation, independently of Anaconda. It is possible to install (some) Conda packages within a virtualeny, but it is better to use Conda's own environment manager. Conda's environment manager works seamlessly with both Conda and pip. Just like pip, you usually run Conda as user, not as root.

4.6 Installing Packages With pip And virtualenv

Here we go into more details of using pip and virtualenv in Linux and macOS, which is the recommended way to install packages (unless you are using Anaconda as explained above). We'll go through the process step by step.

But before we start, I would like to explain why we use virtualenv in the first place. There are two reasons:

- 1. You can use it to manage Python packages for different projects. For example, you might have two projects, in the first you need for backwards compatibility an old version of the gensim package, while for your other project you use a new version of gensim. With virtualenv, you just keep the two version of gensim separate and each project can use the version of gensim it needs.
- 2. With virtualenv you install packages as a normal user (no sudo or root required). This means you avoid installing Python packages globally on your whole system, which could potentially break system tools or other projects.

First of all you want to make sure that your pip and virtualenv are up to date by typing the following commands into the command line:

```
sudo python3 -m pip install -U pip
sudo python3 -m pip install -U virtualenv
```

The above commands perform a system-wide installation/upgrade of pip and virtualenv, as the installation is run as root user. Alternatively, if you prefer an installation/upgrade of pip and virtualenv as a normal user (and install into site.USER_SITE), you can add the --user option and run the commands *without* sudo like so:

```
python3 -m pip install --user -U pip
python3 -m pip install --user -U virtualenv
```

Now that pip and virtualenv are up to date, we can next explain how to install packages into a virtual environment.

When you create a new project, you usually put it into a new directory. For example, let's assume you that to put your project into the myproject directory. The following code lets you create that directory, change into that directory, and then create a virtual environment there.

This code will set up a directory called env inside your project directory that contains the virtual environment. If you're using Git, you should exclude env from your version control system using .gitignore.

Your virtual environment is now set up. Note that you when you have activated the environment using the steps below, you do not need to use pip3 or python3. You can simply use pip and python because inside this environment they will automatically use Python 3, as the virtual environment has been created with Python 3 above.

In any case, whenever you want to work on your project, you follow these steps:

1. Activate the virtualenv. You change into your project directory (e.g. cd myproject) and then activate the virtualenv.

```
source env/bin/activate
```

2. You write code for your project or install packages you need for your project. For example, if you want to install or upgrade the requests package, you can now type at the command line:

```
pip install -U requests
```

Note that this command simply a shortcut for alternatively running the more verbose

```
python -m pip install -U requests
```

3. When you're done (e.g. want to switch to another project or leave your virtualenv for some reason), type

```
deactivate
```

Throughout, you can always check the location of your Python interpreter by typing which python.

4.7 Python IDEs

An IDE is an integrated development environment. It enables a programmer to bring together the different aspects of computer programming into a single application. Examples include editing source code, running your programs, and debugging, which can all be done within a single IDE.

There are many IDEs for Python. If you use Anaconda, I recommend using Spyder, as Spyder is already available in Anaconda. On the other hand, if you are not using Anaconda, I recommend using Atom because it is the most popular IDE.

- Atom is a very popular code editor developed by GitHub. As you can imagine, it has
 very tight integration with Git and GitHub. Furthermore, Atom supports many
 other programming languages besides Python. Useful add-on packages for data
 science:
 - You can combine Atom with Hydrogen, which lets you run your code directly in Atom using Jupyter.
 - Data Atom lets you write and execute SQL queries.
 - Markdown Preview Plus (MPP) has support for editing and visualizing Markdown files.
- Spyder is also built with a focus on data science and has an interface similar to RStudio, just like Rodeo. If you are using Anaconda, I recommend using Spyder as it is available in Anaconda.
- Rodeo was built specifically for data analysis. It is also relatively similar to RStudio, so switching between both is easy.
- PyCharm is made by JetBrains, which is well-known for a famous Java IDE called IntelliJ IDEA. If you have used other JetBrains products, PyCharm might be more familiar to you. However, PyCharm is not all open-source.
- AWS Cloud9 is an interesting alternative from Amazon. It is cheap, but not free. It lets you code directly in your browser and you don't need to configure your local machine. It also enables collaboration between different people in real-time.

4.8 Text Editors

- For beginners I generally recommend using one of the Python IDEs mentioned in Section 4.7 starting on page 29.
- Vim and Emacs are two alternative code editors for the brave.
- Instead of going with a full-fledged IDE, you can alternatively use Vim or Emacs, which are relatively ancient: vi and Emacs were both initially released in 1976.
 Nonetheless, they are very powerful text editors that still very widely used today.

- In fact, Vim started as a clone of the original vi editor and has added more features. Nowadays Vim is more popular than the original vi. On most Linux distributions, if you type vi on the command line, you will be redirected to Vim.
- There is a long history of mostly friendly rivalry between Vim and Emacs, just google "Editor Wars." In terms of popularity, Vim is by now the clear winner over Emacs, although that doesn't necessarily make Vim better than Emacs. I personally tend to favor Emacs slightly due to its internal use of a Lisp dialect to provide a deep extension capability. (Lisp is a very powerful family of programming languages with a long and distinctive history originating from MIT.) On the other hand, Vim has its own extension language as well, Vimscript.
- Whether you decide on using Vim or Emacs is really up to you and based on your personal preferences. You will make a good choice in any case.
- The basic idea about Vim and Emacs is that you can be more productive with them because you avoid taking your hands off the keyboard all the time to do something with your mouse. It is possible to control Vim and Emacs from your keyboard alone, without having to click around with your mouse. Ideally you get into a **flow-like state of mind**, where your ideas flow naturally from your thoughts through the keyboard onto the screen.
- An advantage of Vim and Emacs is that you can use both with all major programming and markup languages. This means the if you work e.g. with both Python and R, you don't have to switch between different IDEs all the time. For example, these lecture notes are written in Emacs (using the document preparation system LATEX) and the Python programming code for this course is written in Emacs as well.
- Both Vim and Emacs are extremely extensible and you can customize them ad infinitum. **You can even play Tetris on Emacs** if you wish! There is the joke by Vim advocates that Emacs is a great operating system, lacking only a decent editor.
- No matter whether you use Vim/Emacs for Python or R, you should consider utilizing specific add-on packages. For example, if you use R with Emacs, you should use the ESS add-on.
- There are several so-called "**Emacs starter kits**" out there, the most popular being Spacemacs. You can combine it nicely with Magit, which is an interface to the Git version control system, see Chapter 5 starting on page 31.

Version Control

A version control system lets you track changes in computer files, i.e. mostly source code files, but also other files such as documentation. Moreover, it allows you to coordinate the work performed on those files among several people. And even if you just code by yourself, it can be immensely helpful to put your code under version control. For example, it allows you to track past changes you made in your code. And of course if you are working in teams on a software project, then version control is a must. It allows you to keep your codebase in a consistent state, even if several people work at it at the same time.

5.1 Git

There are many version control systems out there, the most popular one for open source being Git. You can put your code in a so-called **repository**, which contains your code as well as the metadata that keeps track of the changes you have made. Different **revisions** act like snapshots of your codebase. Git is often operated from the command line, although there are also graphical clients available. There are a number of useful online resources available:

- Setting up Git
- Using Git
- Git cheat sheets
- Git and GitHub learning resources

5.2 GitHub

• GitHub is the leading hosting service for repositories. As the name indicates, it is mainly targeted at Git, but by now it also supports other version control systems. In any case, Git plus GitHub nowadays is the gold standard and you are encouraged to use this combination.

- Why use GitHub to host your repositories? For three reasons:
 - 1. It increases your visibility and helps you with your career and job market. You can reference your work displayed on GitHub directly on your CV and in your application package to impress potential employers.
 - 2. It is best industry practice to use version control.
 - 3. GitHub fosters collaboration and an open culture.
- For instructions see the official documentation, or google for something like
 "GitHub tutorial Linux" or "GitHub command line tutorial" (without the quotes).

 I recommend using GitHub from the command line and setting up SSH keys. It just makes your life much simpler compared to HTTPS, where you always have to re-enter your password.
- Your **data** can go on GitHub as well. If it is not too big you can just include it in your normal code repo. However, a better practice might be to use Git Large File Storage or Dropbox. In any case, the **data should be publicly accessible** so that the instructor can run your code as well.
- If you need to share some **code snippets** (i.e. small regions of re-usable source code) or notes, put them on GitHub Gist. This can come in handy for blog posts, see also Section 2.2 starting on page 13.
- Keep in mind that GitHub can render GitHub Flavored Markdown (GFM), and Jupyter Notebooks, so you can make a lot of documentation etc. available online directly in your GitHub repo (or reference it in your blog, see Section 2.2).
- Keep in mind that **everything you put on GitHub** is available for the whole world to see (unless you use a private repo, which you should not do for this course). This is part of the reason why open-source software often is of high quality. People cannot hide crappy code in their drawer any more, so they have incentives to polish it. Another reason is that coding is much more collaborative, so your learning curve increases because you can learn from other coders much more easily; furthermore, it is easier to identify star-coders whom you can follow and you can read and learn from their code.

Data Sources

This chapter contains a few data sources for obtaining textual and financial data. The list is not all-inclusive. Its main purpose is to inspire you about what is possible. Keep in mind that I have not checked for each and every data source whether they allow bulk downloading and/or web scraping. You should always stay within legal limits regarding download policies.

6.1 Example Texts

If you need some example texts to run your code on (e.g. if you have not downloaded your data yet but would like to start writing code to analyze text), you can look at the janeaustenr and gutenbergr packages in R that contain example texts. If you need the data in Python, you can either save it to a file from within R, or you can call the R packages directly from within Python using rpy2. (Please keep in mind to install R before you install rpy2; if you use Anaconda you can install both R and rpy2 from Anaconda directly.) For example, the following code extracts Jane Austen's novel *Sense and Sensibility* into Python from the janeaustenr R package:

```
from rpy2.robjects.packages import importr
from rpy2.robjects import r
importr("janeaustenr")  # Load 'janeaustenr' R package.
x = list(r['sensesensibility']) # Convert R dataset to Python.
x[0:10]  # Take a look at beginning of novel.
```

6.2 Text Data

Chapter 9 starting on page 47 provides an overview of Python packages that can be used to download and web scrape from the various data sources discussed in this section.

• Factiva by Dow Jones & Company has newspaper articles and newswire articles (e.g. press releases of companies) as well as earnings calls transcripts

- Capital IQ by S&P Global has various business-related transcripts such as earnings calls
- Seeking Alpha hosts a wealth of information, for example earnings calls and articles about the stock market.
- Project Gutenberg
- Internet archive
- · Glassdoor company reviews
- LinkedIn
- SEC EDGAR: 10-K, 10-Q, Form ADV, etc.
- FOMC statements/minutes of the Federal Reserve
- Wikipedia: Download the dump of all Wikipedia articles from here (you want the file enwiki-latest-pages-articles.xml.bz2, or enwiki-YYYYMMDD-pages-articles.xml.bz2 for date-specific dumps). This file is about 10 GB in size and contains (a compressed version of) all articles from the English Wikipedia. Using gensim on the command line, you can convert the articles to plain text and store the results as sparse tf-idf vectors:

```
$ python -m gensim.scripts.make_wiki
```

It might take up to ten hours. You might want to compress the output with bzip2 as Gensim can work with compressed files directly to save disk space.

- Twitter and/or StockTwits, see also Section 11.3 starting on page 73. There are also some websites that provide data for a subset of Twitter, e.g. the Trump Twitter Archive.
- Sentiment140 is a dataset with 1.6 million tweets
- Facebook and Google+
- Medium.com
- Instagram
- Tumblr
- Reddit
- Blogs and blog aggregators/platforms like Blogspot and Blogger
- Online forums and discussion groups

- Company websites, including (but not limited to) press releases and investor relations
- PR Newswire or other newswires
- News, e.g. mine websites of major news providers or take a look at Google/Yahoo News, Google/Yahoo Finance, etc. or directly at news sources like NYT, AP, CNN, as well as international sources. See also the Python packages we have mentioned under web scraping in Section 9 starting on page 47.
- You can also obtain news from News API, which is probably the simplest way if you don't mind their download restrictions (e.g. currently one month's worth of data).
- There is also a list of other news media APIs on Wikipedia.
- You can use the WayBack Machine to download historical news, e.g. from the archives of Yahoo News, Google News, or obtain data from Reddit, which is a social news aggregation, web content rating, and discussion website. However, keep in mind that the WayBack Machine does not always take snapshots on a regular basis, so the coverage might have gaps.
- JSTOR
- Rotten Tomatoes
- YouTube, e.g. you can analyze the comments
- Bloomberg has company filings, analyst reports as well as legal and business documents. Some of them might be downloadable via the Bloomberg Python API.
- Product reviews (the exact location depends on what kind of product you're looking at, e.g. Amazon, Walmart, Ticketmaster, and TripAdvisor)
- In principle you could also look at VK or Sina Weibo, but we want to focus on the English language in this course to keep things simple
- There are also services that provide firehose access to social media, e.g. Six Apart, Spinn3r, Datasift, and GNIP (now part of Twitter).

6.3 Financial Data

This section provides a brief overview of some of the most popular financial databases. These data providers can be useful if you want to figure out how your textual data relates to various financial statistics and events, e.g. asset prices, corporate events, or fundamental data.

• **Bloomberg** has almost all financial data you could imagine, but it is also one of the most expensive data providers out there.

- If Bloomberg is too expensive, you can consider Thomson Reuters **Eikon** as an alternative.
- Capital IQ and FactSet are popular for fundamental data about companies (among other things). Capital IQ and FactSet are popular in investment banking, e.g. Mergers & Acquisitions, so if you need to analyze a company in detail, it is very useful to look at one of these databases.
- Morningstar, Compustat (via WRDS), Reuters Fundamentals, and Worldscope also have fundamental data.
- While both Capital IQ and Compustat have fundamental data and both are provided by S&P, they differ in the way they standardize the accounting variables and also in the time period covered (Capital IQ goes back to 1989 while Compustat goes back to 1950).
- For industry-specific metrics (e.g. ROAA or Tier-1 capital ratios for financial institutions) or news, you can use S&P's **Market Intelligence Platform**, formerly known as SNL Financial.
- I/B/E/S has earnings estimates.
- The **Wind** database is a leading provider for fixed income data for China.
- **AlphaVantage** is a free data provider for stock prices and digital/crypto-currencies (see also Section B.6 starting on page 214).
- Coinbase can be used for digital/crypto-currencies.
- Yahoo Finance and Google Finance provide stock prices, among other data.

Chapter 7

Applications of Text Analytics and NLP in Finance

This chapter contains examples of text analytics and NLP applications in finance and beyond. Its purpose is not to go into great detail for every application, but instead to provide an overview of what is possible. Furthermore, after going through this chapter, one should have a more intuitive understanding of the potential capabilities of text text analytics and NLP.

7.1 General NLP Applications

Most people are unaware that they have already used text analytics and NLP in one form or the other before. The following list provides an overview from general applications, while we discuss finance-specific examples further below.

- Spell checking in Microsoft Word
- Email spam filters
- Call centers transcribe conversations into text and analyze it to find out more about common complaints and problems
- Email messages with complaints to a municipal authority are automatically routed to the appropriate department (or returned if they contain inappropriate or obscene messages)
- Search engines such as Google and Bing
- Google Translate
- Amazon's Alexa, Google's Now, Apple's Siri, or Microsoft's Cortana respond to vocal prompts (which are transcribed to text internally) and do everything from finding a coffee shop to getting directions or turning on the lights at home
- Question answering, e.g. IBM Watson for healthcare, weather, and insurance.

7.2 Applications in Finance

The list below includes many applications to make it clear there are relevant topics in the real-world that can help you make money and/or make the world a better place in general:

- Fed watching, i.e. analyzing reports and minutes coming out of the Federal Reserve. For example, you can calculate the text-based probability that they are going to raise or lower rates, which is a trillion dollar question.
- Understanding and responding to consumer and investor sentiment in financial newspapers and social media. This is not only of interest to corporations for the public/investor relations departments, but can also provide important feedback loops for top managers, e.g. the C-suite (CEO, CFO, COO, and CIO).
- Improve risk-adjusted performance in asset/investment management by analyzing financial newspapers, social media, company filings, etc. The idea is to tap the "wisdom of the crowd" to extract additional information that is otherwise not readily available. This is important because it benefits investors (they get richer) and ultimately can give you as the portfolio manager a higher bonus and the hedge/mutual fund can generate higher income from fees.
- Auditing can be automated to some degree, as is currently being done by KPMG using data analytics. In particular, textual auditing content can also be analyzed using NLP and text analytics.
- Using the same data sources, one can estimate the profitability or the default risk of companies.
- Using online product reviews to gauge customer satisfaction as well as product quality and whether people are receiving a good service. Happy customers might translate into higher sales and generally a better company, which gets reflected in higher stock prices.
- Banks can use chatbots to automate customer service functionality or provide roboadvisory services (e.g. you're investing according to a quantitative model provided by your robo-advisor, and if you have questions you can simply go have an online chat with your virtual advisor).
- Monitoring of company events such as earnings announcements, share buybacks, or mergers and acquisitions. Potential data sources are company reports, analyst reports, newspaper and newswire articles, and social media. For example, if you are running a hedge fund for merger arbitrage, you can try to find out the ex-ante probability of a merger being announced or post-announcement you can find out the probability that the merger completes. Having sharper insights into these events can directly translate into higher profits.

- Due diligence of business/company and legal documents. If textual analysis produces a red flag, a lot of money can be saved by canceling a merger deal or an IPO, for example.
- Detecting insider trading. Again this may be done by analyzing company filings, business/legal documents, newspapers, social media, etc. Of course this kind of information can only be based on circumstantial evidence and is no substitute for a deep investigation using e.g. forensic accounting, but it is much easier and faster to do and allows you to monitor a much larger set of companies, and importantly, you can do this using public information only (no private information required).
- Monitor the news for key hire alerts, e.g. when a company (maybe your competitor)
 hires talented finance, development, marketing, or sales executives, or even people
 from the C-suite.
- Fraud detection, e.g. transcribing earnings calls and analyzing the choice of words used by C-level executives (CEO, CFO, COO). In fact, one could even go further and analyze their voices directly, but this would go beyond text analytics. Similarly, one could analyze regulatory filings for language that implies fraud, e.g. increased usage of negative-emotion words or reduced usage of first-person pronouns.
- Analyzing warranty or insurance claims.
- Analyze competitors by crawling their website and learn about what topics, terms, and features are most important.

7.3 Ideas for Further Work

The examples below go into greater detail and discuss potential applications in finance including the relevant databases. Please respect all legal limits regarding the downloading and the usage of the databases mentioned below.

- Realvision trade ideas. 1) Low likes, high like/dislike ratio, overwhelmingly negative comments: Go with it in size. 2) High like/dislike ratio, massive enthusiasm in comments: Sell as much as you can.
- How do discussions on social media differ during expansions and recessions? Can we use these differences to predict recessions?
- Do firms that have mostly good Glassdoor reviews perform better than firms with bad reviews?
- Can we learn something from the tweets of "famous" finance Twitter (or Stock-Twits) users about the future behavior of stock prices or stock market indices? For example, can we say something about future volatility or future returns?
- Does news coverage of bitcoin lead or lag the bitcoin price?

- Which articles on Medium are about finance-related topics? If they are, in what groups can they be clustered?
- Are stocks with more news coverage less risky?
- Which SEC EDGAR filings are most suitable for predicting fraud or scandals, e.g. in accounting (you could look at several famous scandals such as Enron to begin with and use them as a training sample)?
- Scan PR Newswire for merger announcements and extract the names of the target and the acquirer. How fast does the stock market react to this news?
- Do firms with more disclosure have less surprises in their earnings announcements?
- Calculate news sentiment based on streaming analysis of online news. Are news particularly euphoric/upbeat before stock market drawdowns?
- When analyzing the textual content of company filings from SEC EDGAR (e.g. of financial institutions), can we detect signs of financial market instability?
- Do companies that are frequently mentioned together with fintech terms such as "blockchain" do better on the stock market?
- Can we learn anything from FOMC statements/minutes about the probability of a future change in interest rates?
- Can textual content from the insider trading Forms 3, 4, 5 on SEC EDGAR predict strong selloffs in stock prices? See also here.
- Do firms that mention terms related to hedging and/or derivatives usage have riskier cash flows and/or riskier stock prices?
- Scan newspaper articles (e.g. from Google/Yahoo News or other sources) and/or newswire articles (e.g. from PR Newswire or other sources) and find out which companies have the most mentions of "hot" topics such as "blockchain," "fintech," or others. Do companies that are mentioned more frequently with these hot topics have higher stock returns than comparable companies? You should adjust for company size and/or the number of articles released for each company to ensure that your results are not driven by size or by media coverage.
- Extract product descriptions from 10-Ks and find out how similar companies are to each other in this space. It is possible to derive industry groups based on this similarity, see e.g. Hoberg and Phillips (2018 JPE). There are many questions that can be addressed with this data, e.g. do stocks of companies in similar text-based industries co-move together?
- Construct sentiment from conference call transcripts. Does positive sentiment predict higher stock prices?

- If there are more words related to "risk" in analyst reports (or other financial text documents), are stock prices or operating performance riskier/more volatile?
- Look at news articles and check what clusters they are falling into, e.g. using *k*-means clustering. Then find out what the market does on the following day depending on which cluster an article falls into. If you find a pattern, you can build a trading strategy based on that. For example, you find there are three clusters, and whenever an article falls into cluster two, the market goes up the next day on average. From now on, every time an article falls into that cluster, you go long the market.
- Does more disagreement on stock message boards (or Twitter or FinTwit) imply the stock is more volatile? Is it possible to trade profitably on this higher volatility, e.g. using option trading strategies such as straddles or strangles?

Chapter 8

Data Workflow

The following sections illustrate typical data workflows in finance. Of course, depending on the actual application there might be some differences. Furthermore, it is important to keep in mind that these steps are pretty general and might or might not include a textual analytics component.

8.1 General Data Workflow

An important point is that this workflow should not be thought of as going linearly through it. Instead, it requires a lot of iteration. For example, when you validate your data, you might discover some flaws and need to go back to data cleaning. Or when you analyze your data, you find out that you need a different variable, in which case you go back to data transformation. In any case you can tie much of the workflow together programmatically with tools such as Drake.

- 1. Data collection: This is obviously the first step, but unless you just download raw files, you should already at this stage start thinking about the next step. In any case, it is usually a good idea to separate the raw data from any data that is derived from the raw data. Put the raw data into a separate directory and write-protect it to avoid accidental alterations.
- 2. Data structure: This step is not so much about programming, but more about thinking and planning ahead on a conceptual level.
 - It is really important to have thought about how you want to structure your data. If you are in doubt, use **tidy data** (see also Hadley Wickham's article in the JSS).
 - Another big question is on database design, which does not always have to be fully laid out in all formality (depending on the size of your project), but you should have thought about this topic in any case.
 - Finally, you should also think about naming conventions for your data files and variable names. Often you will have a lot of files and variables at the end of

the project, and it makes a lot of sense to have consistent naming conventions for file names (actually for both data and code files) and variable names.

- 3. Data storage: Often it is simpler to just store your data in files without using a full-blown database. Examples of these files are binary files (e.g. .rds files in R or .pkl files in Python, see also Section A.13.3) as well as text files (plain text, XML, JSON, etc.). In some cases a database might be helpful or necessary, especially for large projects. Depending on your needs, I recommend looking at SQLite, PostgreSQL, MongoDB, or Elasticsearch, as these are some of the most popular databases in the main database categories out there.
- 4. Data cleaning, reshaping, preprocessing: This step is basically the programmatic implementation of the previous step (which was more conceptual). This step cannot be overestimated in terms of its importance because you know what happens to the output of your analysis when there is "garbage in."
- 5. Data validation: Does the data indeed have the properties you think it has? You will be surprised how often in the real world the data does NOT look the way you think it looks! Data validation often is not a single step where you write a separate program that performs a set of tests on your data. Instead, data validation should take place throughout your code. Whenever you make an assumption about your data, you should state this assumption explicitly at the appropriate point in the code. If the data does not satisfy your assumption, your program should throw an error. In this respect the assert statement in Python or the stopifnot function in R are valuable and practical tools. For example, if you assume that for each stock ticker symbol there is only one observation per date, you should explicitly state this assumption using assert or stopifnot.
- 6. Merging: This step consists of merging (or "joining") different datasets into a new dataset. For example, if you have data on stock prices and another dataset on accounting variables, you might want to bring both of them together and merge them into a single new dataset. Another example is if you have calculated some scores based on textual analysis and want to merge these scores to your stock market data. We discuss merging in detail in Appendix B starting on page 203.
- 7. Data transformation: This step takes the cleaned data and transforms it such that you can use it straight away in the following step about data analysis. For example, you might want to center some variables or apply other transformations. Depending on the task at hand, data transformation might take place before and/or after merging.

8. Data analysis

• Summary statistics, statistical methods, machine learning, etc.

- Data visualization (more details in Chapter 20 starting on page 147): This encompasses plotting and online apps. This step is actually a very important part and you should think early on how you want to present and visualize your results, be it a Jupyter Notebook or a dashboard made with Dash or Pyxley in Python, or ggplot2, Shiny and related packages in R.
- Summarize the information we can learn and the actions we should undertake based on our data analysis.

9. Community engagement and public relations:

- Documentation: Your work should be extremely well-documented, both in terms of commenting your source code (explain on a high level what it does and, more importantly, why it does so) as well as writing external documentation. This is not only important for other people who want to use your software or contribute to it, but also for future self when you get back to working on your project after taking a break.
- Set up a blog and write about your work. See also Section 2.2 starting on page 13 for more details.
- Give presentations about your work at user groups, fora, and conferences.

10. Quality control:

- Code review should take place on regular intervals where people review other people's commits or the source code.
- Pair programming means that two people sit in front of the computer, one is the "driver" who writes code, while the other is the "observer" who reviews the code as it is being typed in by the driver. The two people switch their driver/observer roles frequently. Pair programming can be extremely useful because it solves a couple of problems. First, it avoids that people get too tired. After all, our attention span is typically pretty short. Second, there is a lot of learning involved as it is a collaborative exercise. Third, it yields higher code quality.
- Code refactoring: This refers to rewriting the code in a better way without changing its functionality. This should be taking place basically all the time to constantly improve the code quality.
- TDD: Test-driven development (TDD) means that you add tests to your code and make sure they pass. Although you can never be 100% sure there are no bugs, it really increases your confidence if you have a lot of tests that pass. TDD makes mainly sense for larger projects, but even for smaller projects you can often add one-liners where some basic assumptions about the behavior of your code or your data are checked.

8.2 Investment Management Data Workflow

This workflow is more specifically tailored towards investment management. You can view it as a different flavor of the workflow from Section 8.1, looking at things from a slightly different and more specialized angle.

- 1. Identify and acquire the data
 - Formats: CSV, JSON, API, streaming, Text, HTML, etc.
 - People: Data managing team, legal/compliance
- 2. Store, structure, and preprocess the data
 - Database/computing infrastructure: SQL, NoSQL (MongoDB, Redis, Cassandra, etc.), HDFS, the "cloud," Spark, etc.
 - People: Software, system, data engineers
- 3. Analyze data via machine learning, design signals, backtest strategies
 - Analytics software: Excel, R, Python, Dask, Spark, Tableau, etc.
 - People: Data scientist, quantitative researchers, portfolio managers
- 4. Trade ideas, trading signals, risk analyses
 - Output: Report, alert, signal, dashboard, etc.
 - People: Traders, portfolio managers, execution system

Chapter 9

Core Python Packages

Python is a language that really shines when it comes to its ecosystem, i.e. the packages available. In fact, one reason why Python is so popular is that for many tasks you do not have to write a program from scratch, but instead can use an existing package written by someone else that provides the required functionality.

This chapter therefore provides a list of selected Python packages that are useful in the context of text analytics and NLP. The list is not set in stone, but it will give you a good overview of the most important Python packages for data science and NLP. I have ordered them roughly according to their popularity, in particular by the amount of stars on GitHub.

9.1 NLP

We introduce the following packages in detail throughout this book:

- SpaCy
- Gensim
- NLTK

9.2 Web Scraping

Many of the most interesting textual datasets are available from the internet. To obtain this data, one has to use a technique called "web scraping," which means to "scrape" the data from the web. This is a huge topic and the list below is an opinionated summary of the most important Python packages. See also Chapter 11 starting on page 65 for more details on how to use these packages.

9.2.1 Package Recommendations

1. If you need to scrape data from a specific web service such as Twitter, Facebook, Atom/RSS, etc., check some of the **specialized packages** listed below (or if the

service is not listed below, google around).

- 2. If you need to deal with JavaScript/Ajax-heavy sites and there is no specialized Python package available, use **Selenium**. If you like Requests, you can also combine Selenium with Requestium or Selenium-Requests (use Selenium to load the site, log in, fill out forms, etc., then Selenium-Requests moves all the cookies from Selenium into a Requests session where you can continue to use Requests as usual). Alternatively you could use requests_htlm, which can also deal with JavaScript.
- 3. If the website does not use a lot of problematic JavaScript/Ajax and if there is no specialized Python package available, check the following sequence whether it works for you:
 - If you just have to download and parse some webpages, use **Requests and Beautiful Soup**, see Chapter 11 starting on page 65 (or alternatively Requests and lxml if you prefer). Keep in mind that the requests. Session method (from the Requests package) allows you to persist parameters and cookies across requests made, which might come in handy if you need that sort of thing (although RoboBrowser might be easier in that case, see next bullet point).
 - If you need to "do" something with the webpage before getting your data (e.g. fill out forms, click buttons, logging in, etc.), use **RoboBrowser**.
 - If you need to crawl through the web, use **Scrapy**, maybe together with Scrapyz which tries to make Scrapy easier for simple spiders (or alternatively use PySpider or Pattern if you prefer).
- 4. Morph.io is great if you need to host your scraper in the cloud for free.
- 5. Important: No matter what you do, **do not overload the website you are trying to scrape**. If you hit it too often with your requests for data, it might thing you are an attacker and will shut down your IP address. In this case, you probably will have to wait for some time to access the website again.

9.2.2 Overview of Various Web Scraping Packages

- Obtaining data:
 - Requests
 - urllib2 used to be popular, but nowadays most people use Requests.
 - Selenium can be used if there is a dynamic website that uses JavaScript/AJAX,
 or if you need to click through some forms to get to the data you need.
 - RoboBrowser is similar to Selenium but is built on Requests and Beautiful Soup. For some of the older folks out there, RoboBrowser is similar to Mechanize (which has been around since 2004). It has browser history and cookies, is able to fill in forms and click links. However, if you need to deal with JavaScript/Ajax, Selenium is more suitable.

- requests_html is similar to the requests+BeautifulSoup combination, but it can deal with JavaScript. Effectively it is an alternative to Selenium.
- Extracting/parseing data from previously downloaded HTML or XML documents:
 - Beautiful Soup (bs4)
 - lxml: Some people are of the opinion that lxml is better than Beautiful Soup. For those people using lxml instead of Beautiful Soup, most use lxml.html, with lxml.html.html5parser if needed. Some people also use lxml.etree with HTMLParser class if needed. For CSS selectors, lxml.cssselect is popular.
 - requests_html, see above.
 - python-readability can extract the main body text from a HTML document and clean it up.
- Scrapy is a complete web framework, but for simple web scraping tasks it is overkill. It can be useful if you want to have a spider that crawls through entire websites in a systematic way. You can swap individual modules with other Python web scraping libraries, e.g. if you need to scrape dynamic websites (i.e. websites making heavy use of JavaScript/Ajax) you can use Selenium as a backend for Scrapy.
- PySpider is another popular web crawler similar to Scrapy (but less popular).
- Pattern is a Swiss Army knife, it can get data from web services such as Google, Twitter, and Wikipedia and can in general also crawl the web. Furthermore, it can do NLP, machine learning, and network analysis.
- Specific web services:
 - Get newspaper data: Newspaper, Goose, or news-please. If you look at the source code of news-corpus-builder you can see some example code using Goose (and Feedparser).
 - See also Pattern above for Google, Twitter, Wikipedia
 - Newspaper3k and Python-Goose for extracting the text of an article as well as meta information such as the title or author.
 - There are many Python packages for accessing Twitter. Broadly speaking, packages fall into two categories depending on whether they use the official Twitter API or reverse-engineer Twitter's JavaScript front-end (which is an undocumented, unofficial API). The official Twitter API is convenient to access, but it has several limitations, e.g. the amount of data you can download. In contrast, the unofficial JavaScript API often imposes fewer restrictions and typically does not require authentication (but one would need to check the legal terms for downloading). The following list is a partial record of Python packages for scraping Twitter, sorted according to popularity as measured by GitHub stars:

- 1. Twint does not require authentication, does not use Twitter's official API, and does not have rate limitations.
- 2. Tweepy, see also Section 11.3 starting on page 73. Tweepy conceptually is the "opposite" to Twint in the sense that Tweepy accesses the official Twitter API.
- 3. python-twitter for accessing the official Twitter API.
- 4. twitter-scraper, installed and imported as twitter_scraper, for accessing the unofficial API.
- 5. twitter for accessing the official Twitter API.
- 6. twython for accessing the official Twitter API.
- 7. twitterscraper for accessing the unofficial API.
- 8. snscrape for accessing the unofficial API. Besides Twitter, snscrape can scrape from various social networks such as Facebook, Instagram, Reddit, Telegram, VKontakte, and Weibo.
- 9. GetOldTweets3 if you need to retrieve older tweets. For accessing the unofficial API.
- 10. Scweet if everything else fails. It only uses Selenium, so it "pretends" to be a normal user, and it doesn't use the official or unofficial API.
- Praw lets you access the Reddit API
- facebook-page-post-scraper for scraping Facebook pages.
- Feedparser for parsing Atom and RSS feeds

9.3 Extracting Text from PDF

- Textract is a very popular package to extract text from any document. It is a wrapper for Poppler/Xpdf's pdftotext.
- Tika is an alternative that might be easier to install if you're running Windows.

```
from tika import parser
raw = parser.from_file('sample.pdf')
print(raw['content'])
```

- PyPDF2 is sometimes mentioned, but it seems to have some problems as it does not always recognize the text correctly.
- If nothing works, you can try to call the pdftotext binary (from Xpdf) from Python directly. The code could look similar to the following one, where you might have to adapt the path to pdftotext depending on where it is installed in your system:

```
import os, subprocess
SCRIPT_DIR = os.path.dirname(os.path.abspath(__file__))
```

9.4 Core Data Libraries

These are the packages most essential for data wrangling in Python. If your data has different data types (e.g. one column contains dates while another contains numbers), you should use Pandas. Pandas is explained in detail in Appendix B starting on page 203. On the other hand, for vectors, matrices, or arrays (i.e. data containing only one data type) you can use NumPy. SciPy provides sparse matrices and it has basic numerical tools such as integration, differentiation, and optimization.

- Pandas
- NumPy
- SciPy

9.5 Databases

As your data become larger, it often makes sense to store it in a database. Broadly speaking, there are two kinds of databases: SQL and NoSQL.

SQL databases are used when you can fit your data nicely in tables. Even if your data originally is not in a tabular format, it is often possible to convert it to tabular format and save it in an SQL database. If you need an SQL database, my recommendation is to use PostgreSQL or SQLite.

On the other hand, broadly speaking, NoSQL is any database that is not SQL. As you can imagine, there is a great variety of NoSQL databases, e.g. document-based, key-value pairs, graph databases, or wide-column stores. The most popular NoSQL database is MongoDB.

In Python, you can use the following popular packages to connect to a database:

- SQLAlchemy for SQL
- PyMongo for MongoDB

9.6 Big Data

• Dask is very similar to Pandas (in fact it builds on top of Pandas) and it can deal

with datasets that are larger than your computer's memory. It can also parallelize some operations and thus speed up your calculations.

• Spark if your dataset is really huge. Often Spark is not necessary as Dask is sufficient.

9.7 Finance

- Zipline for backtesting; other backtesting solutions include backtrader, pyalgotrade, qstrader, bt, and pysystemtrade (in decreasing order of popularity as measured by GitHub stars).
- PyFolio for portfolio and risk analysis
- TA-lib for technical analysis

9.8 Visualization and Plotting

See also Section 20.2 starting on page 148 as well as Section B.5 starting on page 213.

- Bokeh is immensely popular and can visualize your data interactively in the browser. You could also combine Bokeh with Chartify, which is built on Bokeh.
 - If you require interactivity, you should give Bokeh a try. On the other hand, for static plots it might be easier to get started with Matplotlib or Seaborn, for example.
- Matplotlib is another very popular plotting library in Python. One of the oldest plotting libraries in Python, it has been around since 2003 and therefore has received a lot of active development over the years. Matplotlib is good for basic plotting such as bars, pies, lines, and scatter plots.
- Seaborn is for statistical data visualizations. It is based on Matplotlib. If you would like to create statistical graphics, Seaborn is a very good starting point.
- Plotly can create interactive graphs in the browser, similar in spirit to Bokeh. Plotly is based on the JavaScript library Plotly.js.
- Altair is a declarative statistical visualization library.
- ggplot is a Python port of the wildly popular ggplot2 package from R.
- Holoviews for data visualization.

9.9 Interactive Web Apps

Sometimes you would like to create a web app that showcases the resuls of your analysis.

- Superset or Redash are very popular, but might be less suitable for smaller apps.
- For simple web apps, you can also use Bokeh or Plotly, as mentioned in Section 9.8 starting on page 52. Bokeh and Plotly are very good for interactive data visualizations.
- **Dash** is based on Plotly and can be used to build interactive web-based dashboards. If you're not sure which package to use, you won't make a mistake using Dash.
- Pyxley tries to be for Python what Shiny is for R
- Bokeh can also create interactive visualizations, although its main focus is more on plotting, not on creating web apps
- Jupyter notebook, maybe even with iPyWidgets for interactive widgets can be useful showing and visualizing what you're doing. It is more like a report that shows what you have done and it's less interactive, but if that's what you need, it's a great choice. Note that GitHub renders Jupyter notebooks, which is great if you put your code on GitHub anyway. You can also render a notebook on your computer (of course) but also at nbviewer.jupyter.org.

9.10 Miscellaneous

- Statsmodels for statistics
- Delorean for date and time

9.11 AI and Machine Learning

AI and machine learning are discussed in detail in Chapter 12 starting on page 79.

- Lime: If you need to explain to someone what your machine learning classifier is doing.
- SciKit-Learn: Great for ML, but for deep learning use another library.
- LightGBM and XGBoost: For distributed gradient boosting.
- Vowpal Wabbit (VW) is very popular, written mainly in C++ (with Python bindings), is pretty fast, and can deal with datasets that are larger-than memory.
- Annoy for approximate nearest neighbors.

- H2O.ai: Great for ML including deep learning. Written mostly in Java, has bindings for Python, R, Scala.
- MLlib on Spark if your dataset is truly huge

9.12 Deep Learning

- See also Chapter 12 starting on page 79 for further discussions on deep learning.
- Note that deep learning can be though of as being a subset of machine learning, although in the Python ecosystem, deep learning is often spread out to separate packages.
- My recommendation for most projects in Python is to use Keras if you're just getting started (maybe with the MXNet backend). The reason is that Keras (as well as MXNet) are usable from a higher level than TensorFlow or PyTorch. This means you don't have to worry about so many details as the most common use cases are relatively easy to implement. On the other hand, if you have a very specific application that requires a customized approach, you might be better off using TensorFlow or PyTorch.
- TensorFlow
- Keras
- PyTorch
- Theano (not recommended for new projects, as Theano is in maintenance mode since end of 2017; the main developer MILA stopped developing Theano.)
- H2O.ai as mentioned above.

Chapter 10

Text Processing

This chapter assumes basic Python knowledge. If you need to brush up your Python skills, check out the Python tutorial in Appendix A starting on page 151.

10.1 Common String Operations

- For more information see the official Python documentation.
- Strings exist withing single quotes 'or double quotes ". Most of the time it doesn't really matter whether you write 'hello world' or "hello world". One important exception is if you want to indicate possession, "This is Barry's shirt." in which case you want to use double quotes. On the other hand, if you have quotes, you need to do it the other way around, i.e. enclose the string with single quotes: '"Where are you going?" she asked.'
- If you have both single quotes and double quotes inside your string, you need to protect them with a backslash (\) like so (both ways give the same result):

```
print('"Where is Sam\'s jacket?" she asked.') # Enclose string in single quotes.
print("\"Where is Sam's jacket?\" she asked.") # Enclose string in double quotes.
```

• **Printing** a string:

```
print 'Hello world!' # Python 2.
print('Hello world!') # Python 3.
```

If you're running an interactive session in the Python shell, you can omit print and just type in the string and it will print it out.

• **Multiple lines** can be created by manually adding a newline (\n) or by using triple single quotes ('''') or triple double quotes ("""):

```
'This\nis a\nTest.'
# Or (same result):
'''This
is a
Test.'''
```

Of course it looks nicer as in the following examples, but in that case you would have additional newlines at the beginning and at the end of the string (which is fine if that is what you want):

```
This is a Test.
```

You can have the best of both worlds if you escape the newline using a backslash (\) like so, but keep in mind that there should be no space after the backslash:

```
'''\
This
is a
Test.\
'''
```

• If you don't want a multiline string but just want a long single line string without newlines, just enclose the strings in brackets *without* commas between the strings (otherwise you would be creating a tuple):

```
('Natural language processing (NLP) is a field of computer '
'science concerned with the '
'interactions between computers and human (natural) '
'languages, and, in particular, concerned with programming '
'computers to fruitfully process large natural language data.')
```

Alternatively you could also concatenate the strings together (see also the following bullet point). You can use + and then tell the interpreter that there is more coming in the following line by adding a backslash ("\") at the end of the line (make sure there is no space after the backslash).

```
'Natural language processing (NLP) is a field of computer ' + \
'science concerned with the ' + \
'interactions between computers and human (natural) ' + \
'languages, and, in particular, concerned with programming ' + \
'computers to fruitfully process large natural language data.'
```

• **String concatenation** can be done in several ways. In general, **f-strings are preferred**, but it's good to know the other ways as well.

Sometimes you need to be careful if you have different data types.

```
w3 = 8  # int (not str).
w1 + w3  # Does NOT work.
'{} {}'.format(w1, w3)  # Works.
'%s %s' % (w1, w3)  # Works.
f'{w1} {w3}'  # Works.
```

• If you want to dynamically generate labels, you can use %-formatting as well:

```
x = 8
mylabel = 'label_%s' % x
```

• String replication:

```
>>> 'hello ' * 8
'hello hello hello hello hello hello '
```

• **Raw strings** are useful if you want to shut down the interpretation of escape characters such as backslash (\). You do this by just adding r in front of the string.

```
>>> print(r'Hello \n The milk\'s color is white.')
Hello \n The milk\'s color is white.
```

• Length of a string (will include letters, numbers, whitespace, etc.):

```
len('hello world') # 11
```

• **Indexing** (or "**slicing**") can be done in the usual way (see also Section A.10 starting on page 179):

```
>>> w = 'I love NLP.'
>>> w[0]
                       # First character.
,I,
>>> w[len(w) - 1] # Last character.
·. ·
>>> w[-1]
                      # Last character (counting backwards).
, ,
>>> w[0:6]
                      # From index 0 (inclusive) to index 6 (exclusive).
'I love'
>>> w[:6]
                       # From beginning to index 6 (exclusive).
'I love'
                     # From index 6 (inclusive) to the end.
>>> w[6:]
, NLP.,
>>> w[-4:-2] # From index -4 (inclusive) to index -2 (exclusive).
'NL'
```

• Stride:

```
>>> w = 'I love NLP.'
>>> w[2:9]
              #
'love NL'
>>> w[2:9:1] # Stride of one (same as before).
'love NL'
>>> w[2:9:2] # Stride of two (skip every second).
'lv L'
>>> w[2:9:3] # Stride of three (skip every third).
'leL'
>>> w[::-1]
              # Negative stride, reverse string.
'.PLN evol I'
>>> w[::-2]
              # Skip every other letter in reversed string.
'.L vlI'
```

• **str.count** counts how often a character or a character sequence occurs. This can be used for BoW, see Section 15.1 starting on page 113.

```
>>> w = 'Hello World!'
>>> w.count('o')
2
>>> w.count('h')  # None because it's case sensitive.
0
>>> w = 'Dan likes to swim and he likes to smile.'
>>> w.count('like')
2
>>> w.count('like ')
0
```

• **str.find** finds the first occurrence of a character or character sequence:

```
>>> w = 'Danny likes to swim and he likes to smile.'
>>> w.find('m')
18
>>> w.find('likes')
6
>>> w.find('likes', 9)  # Start looking after index number 9.
27
>>> w.find('likes', 9, -3) # ... but stop looking at index -3.
27
>>> x = 'hello world'
>>> if x.find('llo') != -1: # Check whether can find 'llo' anywhere.
... print('contains string')
...
contains string
```

• Converting to **upper or lower case**:

```
w = 'Hello World!'
w.upper()  # 'HELLO WORLD!'
w.lower()  # 'hello world!'
```

• **Boolean methods**, i.e. checking whether some properties of a string are true:

Method	True if
str.isalnum	Only alphanumeric characters
str.isalpha	Only alphabetic characters
str.islower	All lower case
str.isnumeric	Only numeric characters
str.isspace	Only whitespace characters
str.istitle	String is in title case
str.isupper	All upper case

• For example:

```
'hello'.isalnum()  # True
'hello#'.isalnum()  # False
'Hello'.isupper()  # Fase
'HELLO'.isupper()  # True
```

• str.join:

```
w = 'I love NLP.'
''.join(w)  # Same string as before (no changes made).
''.join(w)  # 'I love NLP.'
''.join(reversed(w)) # '.PLN evol I'
','.join(['hello', 'world']) # Combine list of strings into single string.
```

Here's another example, where we join strings together with a given separator, in this example "_":

```
>>> '_'.join(['hello', 'world', 'this', 'is', 'great'])
'hello_world_this_is_great'
```

• **str.split** can be used as a rudimentary tokenizer, but it's usually better to use a more sophisticated solution, e.g. from Section 10.4 starting on page 63:

```
>>> w = 'I have a balloon.'
>>> w.split()  # Split on whitespace.
['I', 'have', 'a', 'balloon.']
>>> w.split('a')  # Remove letter 'a'.
['I h', 've ', 'b', 'lloon.']
```

• **str.splitlines** splits a string using the newline delimiter, for example:

```
>>> 'foo\nbar\n'.splitlines() # Equivalent to str.splitlines('foo\nbar\n')
['foo', 'bar']
```

In fact, this is (in this case) preferable to str.split because the final newline results in an empty string:

```
>>> 'foo\nbar\n'.split('\n')
['foo', 'bar', '']
```

• str.replace:

```
>>> w = 'I have a balloon.'
>>> w.replace('have', 'had')
'I had a balloon.'
```

• str.startswith

```
>>> x = 'hello world'
>>> x.startswith('hello') # Check whether it starts with 'hello'.
True
```

• Check whether a character is in a string:

```
>>> x = 'hello world'
>>> if 'e' in x:  # Check whether 'e' in string.
... print('yay')
...
yay
```

10.2 Regular Expressions

- Sometimes it is difficult to accomplish your text preprocessing tasks using the common string operations discussed in Section 10.1 starting on page 55. In this case regular expressions might help.
- Regular expressions, or **regex** for short, can take your text processing skills to the next level.
- Regex are strings with a special syntax which allow you to match patterns in other strings.
- You can use regex to find things in text documents (e.g. web links), parse email addresses, and remove/replace unwanted characters.
- In general, regex can be very useful if you have to **preprocess text**.

10.3 Python's re Module

The re module in Python provides regular expression matching operations. It allows you to operationalize some of your text preprocessing needs using regular expressions.

- You can load it with import re.
- The functions in the re module work in the same way. Always pass **pattern first** and **string second**. The functions may return a string, match object, or an iterator.
- The **pattern** tells the function what part of the string to search or replace.
- The **string** is contains the text the function operates on.
- It is often good practice to **prefix your regex patterns** with r (e.g. r"\n") to avoid misunderstandings. Otherwise \n for example would be interpreted as a newline, not as the raw string "\n" (i.e. the character "\" followed by the character "n").

We next introduce some of the most important functions and patterns, and then illustrate how to use them together. We begin by presenting the **most important functions**:

- sub replaces a pattern in a string with another pattern.
- match matches an entire string or substring based on a pattern.

- search searches for a pattern. Similar to match, but doesn't require to match from the beginning of the string.
- split splits a string on a regex. Can be used for tokenization (e.g. with the \s+ pattern).
- findall finds all patterns in a string.

Next we provide an overview of **important regex patterns**. These patterns control the behavior of the methods and functions we have just introduced. The patterns tell Python what kind of text you are looking for, or which part of the text you would like to replace with something else.

Pattern	Matches	Example
\w	Word	'Magic'
\d	Digit	9
\s	Space	, ,
\S	Anything that is NOT a space (capital letters negates)	'no_spaces'
+ or *	Greedy match (grabs repeats of single letters or whole patterns)	'aaaaa'
.*	Wildcard (match any letter or symbol)	'username74'
[]	Define explicit character ranges	
[a-z]+	Lowercase group (creates group of characters by putting them inside square brackets)	'abcdefb'
[A-Za-z]+	Upper and lowercase English alphabet	'ABCdef'
[0-9]	Numbers from 0 to 9	8
[A-Za-z\-\.]+	Upper and lowercase English alphabet, -, and . (using the escape character $\$ ("backslash") to protect the - and .)	'My-Website.com'
1	Logical OR	
()	Define a group, use to find explicit set of characters	
(a-z)	a, -, and z	'a-z'
(\s+ ,)	Spaces or comma	,,,

Now we tie everything together, the functions and patterns, to find out how they work together. The following code shows examples on how to use regex (see the file code-regex.py on the course website). Run the code to see what happens.

```
# This file contains a few examples of how to use the re module and
# how to deal with regular expressions.
import re
                                     Import 're' module.
re.sub(r'K', r'L', 'King Arthur') # Replace pattern in string.
re.match(r'abc', 'abcdef')
                                  # Match a substring.
re.search(r'cde', 'abcdef') # Fill also match in the middle of string.
re.match(r'\w+', 'Hello_world!') # Match a word.
re.match(r'[a-z0-9]+', 'lowercase\_and\_nums\_like\_8, \_but\_no\_commas.')
re.split(r'\s+', 'This_is_a_test.') # Returns list split on spaces, e.g. tokenization
re.findall(r'\w+', "Let's_write_regex!") # Find all words.
# Split into sentences.
re.split(r'[.?!]', "Hello_world!_Let's_write_regex._Isn't_this_great?")
# Find all capitalized words.
re.findall(r'[A-Z]\w*', 'Hello_world,_I_love_Hong_Kong.')
re. findall (r'\d+', 'The\_novel\_1984\_was\_published\_in\_1949.')
# Match digits and words (but not anything else, e.g. punctuation).
re.findall('\d+|\w+', 'He_has_12_cats.')
m = re.search(r'coconuts', 'I_love_coconuts.')
print(m.start(), m.end()) # Print start and end indices.
# Find square bracket containing a word (but no space or anything else).
re.search(r'\[\w+\]', 'Hello_[wind_bla]_this_is_[nice].')
```

10.4 Tokenization With NLTK

Often you need to split up a text into smaller parts ("tokens") such as words, sentences, or paragraphs. This process is called tokenization. The NLTK library provides a few tokenizers that are specialized for certain applications, such as word or sentence tokenization. Furthermore, there are special tokenizers such as for tweets. For an example of tokenization using gensim's tokenizer gensim.utils.tokenize see Section 19.5.2 starting on page 136.

- word_tokenize to tokenize text into words.
- sent_tokenize to tokenize a document into sentences.
- regexp_tokenize to tokenize a string or document based on a regex pattern.
- TweetTokenizer for tweet tokenization, to separate hashtags, mentions, and repeated punctuation marks!!!

Here are some examples on how to use tokenization with the NLTK package (see the file code-nltk-tokenization.py on the course website). Please run the code and see what happens.

```
# This file shows some tokenization examples using the NLTK package.

from nltk.tokenize import word_tokenize, sent_tokenize, regexp_tokenize, TweetToken
```

```
word_tokenize("Hi_there!")
sent_tokenize('Hello_world._I_love_HK!')
# Make set of unique tokens.
set(word_tokenize('I_love_HK._I_love_NYC'))
# Tokenize based on regular expression.
regexp_tokenize('SOLDIER_#1:_Found_them?', r'(\w+|#\d|\?|!)')
# Find hastags in tweets.
regexp_tokenize('This_is_a_great_#NLP_exercise.', r'#\w+')
# Find mentions and hashtags in tweets.
regexp_tokenize('great_#NLP_exercise_from_@user123.', r'[#@]\w+')
tknzr = TweetTokenizer()  # Create instance of TweetTokenizer.
[tknzr.tokenize(t) for t in ['thanks_@user123', '#NLP_is_fun!']]
```

Chapter 11

Web Scraping

This chapter covers the basics of web scraping. Web scraping deals with obtaining data from various pages on the internet. We cover this part relatively early on to enable you to start collecting the data you need at an early stage. If you would like to have an overview about which web scraping library to use, please see our previous discussion in Section 9.2 starting on page 47.

11.1 Primer on Web Technologies

Before we go into web scraping, we first need to understand the basics of the hyper text markup language (HTML). HTML is a markup language that tells your browser the structure of the webpage, e.g. it says that there should be a link to a given website or that some text should be displayed as a title or that a given JPEG file should be displayed as a picture. There are also cascading style sheets (CSS), which determine the visual layout of the webpage. Here we are mostly going to focus on HTML because for our purposes (i.e. NLP), this is where we find the information (i.e. text!) we're looking for. However, CSS (and maybe even XPath) might be important if you need to extract data from very specific parts of the webpage. We discuss this further below. For now we focus on HTML.

11.1.1 HTML

Here is an example of a very simple HTML webpage. The code below can be found in the file code-HTML-example.html on the course website. You can even open this file right in your browser and **take a look how this webpage is rendered (i.e. displayed) in your browser**. It won't look fancy, but you get the point of how HTML works. The following HTML code can be found in the file code-HTML-example.html on the course website.

```
<!doctype html>
<html>
    <head>
        <meta charset="UTF-8">
        <title>Text Mining</title>
```

You can see that all HTML **tags** such as "title," "h1," or "p" are opened using "<...>", then something happens in between, and then the tags are closed again using "</...>". For example, opens a "paragraph," then there is some text inside that paragraph, and then the tag gets closed again with (note the forward slash "/"). Another thing that is important is that some HTML tags have **attributes**. For example, the "a" tag (which is used for creating links) has the "href" attribute. Some websites use another attribute, the "id" attribute, for some tags to give them a unique ID. If that is the case, you can use this ID to extract specific elements from the webpage (otherwise you need to use CSS selectors or XPath selectors, see Section 11.1.4 on page 68).

11.1.2 XML

Some of you might have heard about XML and that HTML has something to do with it. There is a long history about XML and HTML (and we're not going to go into it). The current state of affairs is that nowadays HTML5 (which is the current version of HTML) and XML are very similar. You can write a HTML document either in "HTML syntax" or in "XML syntax." The example above is written in HTML syntax, but could relatively easily be transformed into XML syntax with only minor modifications (mainly at the top of the file, the rest of the file does not require any changes). The difference between the two syntaxes is that the former is roughly 99% valid XML while the latter is 100% valid XML.

Although their files look very similar, XML and HTML have very different conceptual origins. XML is used to save data in a file that is both human-readable and machine-readable. You can think of an XML document as a single-file mini-database. On the other hand HTML is used to create webpages. Specifically, HTML gives the browser the data and instructions needed to render a webpage. It just so happens that HTML gives this data in a format that is very similar to XML. That's why HTML and XML are so similar, even though they have very different purposes. Now why are they so similar? Because both of them originate historically from the same root, which is another markup language called SGML.

Has it always been the case that HTML and XML are so similar? Yes and no. They have always been somewhat similar, but in the past HTML had diverged from XML quite

a bit. The reason is that web browsers (e.g. Microsoft Edge, Mozilla Firefox, or Google Chrome) are relatively lenient when encountering malformed HTML. This has led to poor industry practices where some web designers for example wrote webpages where some HTML tags were not properly closed. In fact, you still might find websites in the wild with HTML that is strictly speaking invalid but still "works" because web browsers "understand" what the web developer is trying to do and still render the webpage, even if it contains invalid markup. This is, by the way, one reason why you need packages such as Beautiful Soup or lxml (discussed in Section 11.2 starting on page 70) is because they fix invalid HTML. In any case, there is an active push towards cleaning up this mess, especially starting from HTML5, which requires the markup language to be very close (or identical) to valid XML.

11.1.3 Document Object Model (DOM)

The Document Object Model, or **DOM** for short, is a way to represent HTML (or XML) in a tree structure. If you look at our HTML example from Section 11.1.1 starting on page 65, you can see that for example the parent of the "p" tag is the "body" tag. And the "body" tag is the child of the "html" tag. So you could actually build a **tree structure** containing parents and children that represents the whole HTML document. In other words, **the DOM** has the same information content as the HTML document, but this information is represented in a different way, i.e. in a tree structure.

Now why do we need the DOM tree? There are at least two reasons for that:

- 1. First, it is useful to identify specific elements of the webpage. We will discuss this further in Section 11.1.4 starting on page 68. Basically it can help you to **grab specific information you need from the webpage**. What happens is that the browser parses the HTML and internally represents the webpage as a DOM tree. So **the browser does actually not work with the HTML after it is parsed, but it actually works with the DOM internally**. Even though you might not directly refer to the DOM tree when grabbing data from the webpage (e.g. you are referring to an "id" attribute of a HTML tag or use a CSS selector or XPath selector), you are effectively using the DOM because that's how the webpage is represented internally.
- 2. Second, it is possible to change and update the DOM programmatically. Section 11.1.6 starting on page 68 discusses this in more detail, but basically you can write a JavaScript program that gets executed inside the browser and manipulates the DOM. For example, using JavaScript, you can remove a specific HTML element such as a link. If that update of the DOM occurs, the browser will change how the webpage is displayed accordingly (e.g. the link will disappear from the displayed webpage). As you can imagine, this is a very powerful way of updating a webpage since you can change any part of the webpage according to your liking. So the key point is that if you want to change the webpage, instead of loading a completely new HTML document (and thus reloading the whole webpage from the web server), you run a program inside the

browser that changes a specific element of the webpage (in the DOM) and leaves the rest of the webpage unchanged. This can be faster to do and requires less new data to download.

11.1.4 CSS and XPath Selectors

So given all that, how do we extract the information we need from a webpage? The simplest way is to simple get all the text from a website after stripping the HTML tags. On the other hand, if you are looking to extract very specific parts of a website, there are several approaches, but in my opinion the following usually works best:

- 1. First, you focus on **IDs** in the HTML markup.
- 2. If that doesn't work, you try **CSS selectors**. They are often more readable, brief, and concise than XPath.
- 3. Only if that doesn't work you use **XPath**. For example, walking *up* the DOM tree (i.e. going from a child to a parent node, see Section 11.1.3 starting on page 67) is not possible with CSS. For XPath you need to use lxml because Beautiful Soup does not support it.

How do you know specifically which elements to choose? You can either look at the raw HTML, which might work for relatively small and simple webpages. If the webpage is more complicated, you can open it in your browser and inspect it there. You can press **F12 in Google Chrome** or you can use the **Firebug extension in Firefox**.

11.1.5 HTTP

HTTP is the protocol through which your browser exchanges data with the web server. For example, if you point your browser to wikipedia.org, your browser uses HTTP to say: "Hello Wikipedia, please give me the files (mostly HTML files but potentially also image files etc.) for your title page." Wikipedia then sends the file(s) to the browser, again through the HTTP protocol.

11.1.6 JavaScript and Ajax

Some webpages are difficult to scrape, especially if they use JavaScript and possibly AJAX. **JavaScript** is a programming language that your browser can execute. It is a whole programming language, so it is very different than HTML (which is a markup language). The difference between JavaScript and HTML is that JavaScript is dynamic, you can have for-loops, if/else statements, and so on, while with HTML it is much more static in the sense that it tells the browser the content of the website *once*, and then that's it.

So the basic idea with JavaScript is that you can make a the website more interactive by changing it on the fly **without having to reload the whole website** (i.e. without loading another HTML file from scratch from the server). For example, Facebook makes heavy use of JavaScript and has written famous JavaScript libraries such as React. And

while JavaScript is already updating the website in the browser on the fly, it sometimes needs to exchange data with the web server (e.g. one of the computers run by Facebook). This is done through **Ajax**, which is a technique to use JavaScript to asynchronously **retrieve data in the background** without interfering with the display or behavior of the existing webpage. A common Ajax technique is to use the **XMLHttpRequest** object (or **XHR** in short) to transfer data between the browser and a web server. It is important to understand that **Ajax is not a new technology and not a new programming language. Ajax is simply a different way to use JavaScript in the web browser to communicate with a web server.**

As you can imagine, scraping webpages that use JavaScript and/or Ajax can seem difficult if you do not have any means to execute that JavaScript code. (For example, you cannot straightforwardly use the Requests library.) All you would get is some HTML that looks very differently than the HTML you would see rendered in your browser. The reason is that **the JavaScript running in your browser updates and changes the HTML** (strictly speaking, it changes the document object model, or **DOM** in short, see also Section 11.1.3 starting on page 67). So without running JavaScript (and packages such as Requests cannot run JavaScript), it seems you are out of luck.

Luckily there are (at least) three solutions to this problem:

- 1. Don't just grab the HTML. Take a remote-controlled browser, point it towards the webpage, let it run the JavaScript (inside the browser), and after the webpage is fully loaded, collect the resulting HTML from the browser. This is the main idea of **Selenium** and a few other similar projects such as Watir (for Ruby) or CasperJS (for JavaScript). This approach can also be useful if you need to **fill out some forms or press some buttons** on the webpage to get to your data (no matter whether the webpage uses JavaScript or not).
- 2. Open your browser such as Chrome or Firefox (no Selenium needed here) and take a look at the XHR communication between your browser and the web server (F12 in Chrome or the Firebug extension in Firefox). Most likely you will see some JSON files (or XML files or other file formats in some cases) being sent back and forth. Most likely the XHR communication will be over HTTP, but it could also be another protocol. The great thing is that often you will find that the Ajax interface is in fact really an undocumented API. An API is a so-called "application programming interface," in this case an interface to programmatically access the database behind the web server. So instead of going through a lot of painful HTML parsing, you can simply use this API to directly grab all the data you need. In fact, due to this undocumented API, scraping data from Ajax may actually make your life much easier in some cases. You can use the Requests library (maybe together with Beautiful Soup to parse the HTTP response and with the json package to deal with the data in JSON format) to accomplish this by sending data (usually JSON data, sometimes XML or other data) back and forth via XHR. This sounds great, but here is a word of caution: It really depends on the website whether this approach works well. Often times it does, but some websites interleave and nest their Ajax so much that it's difficult to understand what's going

on. In these cases it might be easier to just use Selenium.

3. The website has an official API. For example, Twitter uses JavaScript and Ajax, but you don't have to worry about that because Twitter provides a separate API where you can download the tweets directly. See Section 11.3 starting on page 73 for more details.

11.2 Web Scraping Examples

We begin by downloading a website from Wikipedia and saving it to a file. The code is really straightforward, thanks to the simplicity of the Requests package. The code below can be found in the file code-requests-scrape-wikipedia.py on the course website.

```
# This script shows how to use the requests package to scrape a page
# from Wikipedia.
import requests
# Get the response. It is always a good idea to set the 'timeout'
# argument in production code. Otherwise your program may hang
# indefinitely.
r = \setminus
    requests.get(
        'https://en.wikipedia.org/wiki/Natural_language_processing',
# The following code block is strictly speaking not necessary, but it
# helps you to better understand the response you got.
r.raise_for_status()
                                 # Ensure we notice bad responses.
r.status_code
r.headers['content-type']
r.encoding
                                 # 'print' gives nicer output for HTML.
print(r.text)
r.text.encode('utf-8')
                                 # Use specific encoding.
                    # Doesn't work in this example since no JSON data.
r.json()
# Finally we write the response content to file.
with open('data-wikipedia-NLP.html', mode='wb') as fd:
    fd.write(r.content)
```

The next thing we do is use the Beautiful Soup (bs4) library to parse the webpage downloaded by Requests. By "parsing" we mean to extract certain elements from the webpage. The code below can be found in the file code-bs4-read-wikipedia.py on the course website.

```
# This script shows how to download and parse a Wikipedia page using # Requests and Beautiful Soup. We also use the 're' module for regular # expressions. import re import requests from bs4 import BeautifulSoup r =
```

```
requests.get(
        'https://en.wikipedia.org/wiki/Natural_language_processing',
s = BeautifulSoup(r.text, 'lxml') # Use 'lxml' to parse the webpage.
print(s.prettify())
                                   # Take a look at the parsed webpage.
# Extract all the text from the webpage. THIS IS WHAT YOU NEED MOST
# OFTEN FOR NLP AND TEXT ANALYTICS, unless you need to extract only
# part of the webpage.
print(s.get_text())
# Ways to navigate the data structure.
s.title
s.title.name
s. title.string
s.title.parent.name
                  # First 'p' tag.
s.p
s.p.get_text()
                   \hbox{\it \# In our example, there's no 'class' HTML attribute.}
s.p['class']
                  # First 'a' tag.
s.a
# The difference between the 'find' and the 'find_all' methods is that
# the former only finds the FIRST child of this tag matching the given
# criterial, while the latter gets ALL of them.
s.find(id='footer')
s.find(style='clear:_both;')
# Extract all 'a' tags.
atags = s.find_all('a')
                                 # Find all '<a ... > ... < /a>' tags.
atags[3]
atags[3].name
atags[3].get('href') # Get the actual 'href' attribute (i.e. the URL).
# If we want to get all 'href' attributes, we loop over all 'a' tags.
{tag.get('href') for tag in s.find_all('a')}
# Find all tags whose names start with the letter 'b' (in this case
# 'body', 'b', and 'br').
{tag.name for tag in s.find_all(re.compile('^b'))}
# Find all tags whose name contains the letter 't'.
{tag.name for tag in s.find_all(re.compile('t'))}
# We can also pass a list to the 'find_all' method, in which case bs4
# allows a string match against any item in that list.
{tag.name for tag in s.find_all(['a', 'body'])}
# Find all the tags in the document, but none of the text
# strings. 'True' matches anything it can.
{tag.name for tag in s.find_all(True)}
```

The previous example using the Requests package works well on webpages that don't use a lot of JavaScript. However, when a webpage includes a significant amount of JavaScript, the Requests package might no longer work so well because it cannot easily execute the JavaScript to produce the final HTML version of the website. To solve this problem, we can use Selenium WebDriver (using the selenium-python package) in-

stead of Requests to obtain the data we need. After we have downloaded the desired HTML file using Selenium WebDriver, we can then extract further data from the HTML file using Beautiful Soup, similar to the previous example. Before running the following code, you need to download the chromedriver file from chromedriver.chromium.org. Make sure that the version of your chromedriver file matches the version of your Chrome browser, otherwise you will receive an error message. (Selenium WebDriver also works with browsers other then Chrome. We just use Chrome here because it is very widely used.) The code below can be found in the file code-selenium-download-google.py on the course website.

```
# This script shows how to navigate and download a website
# (i.e. Google in the code below) that potentially includes a lot of
# JavaScript. We use Selenium WebDriver for web browser
# automation. This code is only intended for academic purposes.
from selenium import webdriver
from selenium.webdriver.common.keys import Keys
from selenium.webdriver.common.by import By
import os
home = os.path.expanduser('~') # Home directory.
# You might have to update the following line, depending on where you
# have saved the 'chromedriver' file during installation.
driver = webdriver.Chrome(home + '/bin/chromedriver')
# Go to the Google website.
driver.get('https://www.google.com')
# Enter a search string.
e = \setminus
    driver.find_element(
        By .XPATH,
        '/html/body/div[1]/div[3]/form/div[1]/div[1]/div[1]/div/div[2]/input')
                           # Enter search string into Google.
e.send_keys('NLP_trends')
e.send_keys(Keys.ENTER)
                                # Hit the "Enter" key (in browser).
# Go back to Google homepage by clicking on "Google" logo.
driver.find_element(By.XPATH, '//*[@id="logo"]/span/img').click()
# Download the HTML of the webpage for further analysis. Keep in mind
# that this only downloads the HTML of the website, not any CSS or
# JavaScript. After you have downloaded the HTML, you can extract data
# from it, e.g. using Beautiful Soup (not shown here).
with open('data-google-page.html', 'w') as f:
    f.write(driver.page_source)
```

11.3 Mining Twitter

In the previous sections we have seen how to manually scrape content from a website. Twitter also has a website, so we could use these techniques on Twitter as well. However, it might be cumbersome due to the heavy usage of JavaScript and Ajax on Twitter.

Luckily we do not have to worry about this problem because Twitter provides an official API (the "Twitter API") we can use to programmatically download Twitter content. Furthermore, there are several high-quality packages available that allow us to interface to this Twitter API with relative ease (see also Section 9.2 starting on page 47).

Among these packages, Tweepy is one of the most popular. Tweepy uses the Twitter API, which is an interface provided by Twitter. You can use this interface to access Twitter using a program, e.g. a Python script written by you. To obtain access to the Twitter API, you first need to sign up for a (regular) Twitter account, in case you haven't already got one. Second, you need to sign up for a developer account.

There are different access levels to the Twitter API. When signing up for a developer account, you will by default get Essential access, which only allows you to connect to Twitter API v2, but not to Twitter API v1.1. To be able to connect to Twitter API v1.1, you need to apply for Elevated, Elevated+, or Academic Research access. You can apply for this access from within the developer portal after you have signed up for a developer account.

Tweepy only partially supports the Twitter API v2, e.g. the streaming endpoints in the Twitter API v2 are not yet supported in Tweepy. We will therefore focus on Twitter API v1.1 in what follows. For example, this script uses the Twitter API v1.1 fields (e.g. user.screen_name), not the Twitter API v2 fields (e.g. includes.user.username).

After you have set up your Twitter developer account, you need to save the following information given to you by Twitter when you create a "Project" and an "App" on the Developer Portal:

- API Key and API Key Secret: These are the user name and password representing your App
- Access Token and Access Token Secret: These specify the Twitter account the request is made on behalf of

For easy reference, you can save them in a file that you can read into Python whenever you need to. In this example, I save them in a file called code-API-key-tweepy.py:

```
consumer_key = '...'  # Here you enter the API Key.
consumer_secret = '...'  # Here you enter the API Key Secret.
access_token = '...'  # Here you enter the Access Token.
access_token_secret = '...'  # Here you enter the Access Token Secret.
```

You can later read this file into Python by running

```
exec(open('code-API-key-tweepy.py').read())
```

If you have saved this file in a different directory, you need to add the location of the file before the file name. In the following example, the file is saved in the parent directory (represented by ".."), so we prepend ".." to the filename:

```
exec(open('../code-API-key-tweepy.py').read())
```

If you would like to obtain a better overview of the data provided by Twitter, you can take a look at Twitter's data dictionary. Specifically, the documentation about the user object contains information about the various fields such as user.screen_name or user.followers_count that we'll use in some of the Tweepy examples in the section.

In order to connect to Twitter, you supply all of your authentication material to Tweepy, which then handles the connection to Twitter for you (see the file code-tweepy-020-setting-u on the course website):

```
# This script shows how you can set up your Twitter API login
# information.
import tweepy

# Load Twitter API authentication data.
exec(open('../code-API-key-tweepy.py').read())

auth = tweepy.OAuthHandler(consumer_key, consumer_secret)
auth.set_access_token(access_token, access_token_secret)
api = tweepy.API(auth)

# Print the screen name / user name of your account.
print(api.verify_credentials().screen_name)
# Print the tweets in your stream.
public_tweets = api.home_timeline()
for tweet in public_tweets:
    print(tweet.text)
```

If you want to listen to what is happening on Twitter for a given keyword or hashtag, you can use the following code (see the file code-tweepy-040-streaming-api.py on the course website):

```
# This script listens on the Twitter stream for the keyword 'python'
# and prints select fields from each tweet, such as name, follower
# count, date, and the tweet itself.
import tweepy
# Load Twitter API authentication data.
exec(open('../code-API-key-tweepy.py').read())
# Define subclass of 'tweepy.Stream' to add logic to 'on_status'. See
# http://docs.tweepy.org/en/latest/extended_tweets.html.
class listener(tweepy.Stream):
    def on_status(self, status):
```

```
print(
               status.user.screen_name, ':',
                status.user.followers_count, ':',
                                                     # UTC time.
               status.created_at, ':')
           if hasattr(status, 'retweeted_status'): # Check if it's a retweet.
               try:
                    print(status.retweeted_status.extended_tweet['full_text'])
               except AttributeError:
                    print(status.retweeted_status.text)
           else:
               try:
                    print(status.extended_tweet['full_text'])
               except AttributeError:
                    print(status.text)
   # Create a Stream.
   mystream = \
       listener (
           consumer_key,
           consumer_secret ,
           access_token,
           access_token_secret)
   # Start a Stream.
   mystream.filter(track=['python'])
Sometimes you might just want to save the ongoing Twitter stream to a file. This is also
easy to do (see the file code-tweepy-060-stream-to-file.py on the course website):
   # This script listens to a keyword (e.g. 'python' in the example
   # below) and writes the whole Twitter stream to a file.
  import tweepy
   # Load Twitter API authentication data.
   exec(open('../code-API-key-tweepy.py').read())
   class listener (tweepy.Stream):
       def on_status(self, status):
           with open('data-streaming-tweets.txt', 'a') as f:
               f.write(str(status) + '\n\n')
       def on_request_error(self, status_code):
           print(status_code)
   mystream = \
       listener (
           consumer_key,
           consumer_secret ,
           access_token,
```

```
access_token_secret)
mystream.filter(track=['python'])
```

Alternatively, you can also stream only a selection of attributes of the tweet. The difference to the previous example is that here we are using on_status, which is usually what you need, unless you want to stream as much data as possible (in which you can consider using on_data). Another difference is that we are looking for mentions of stock index codes (e.g. \$SPX for the S&P 500) and related codes. This ensures that we are picking up the relevant finance Twitter stream. The following code can be found in the file code-tweepy-070-stream-select-fields-to-file.py on the course website.

```
# This script shows how to stream only selected fields to a file. The
# fields are separated by ': '(i.e. space, colon, space) for easier
# parsing later on.
import tweepy
# Load Twitter API authentication data.
exec(open('../code-API-key-tweepy.py').read())
# Define custom listener class that is a derived class from the
# 'tweepy.Stream' base class. It writes specific data fields from the
# Twitter stream to a file.
class listener (tweepy.Stream):
    def on_status(self, status):
        with open('data-streaming-tweets.txt', 'a') as f:
            if hasattr(status, 'retweeted_status'): # Check if it 's a retweet.
                try:
                    mytweet = status.retweeted_status.extended_tweet['full_text']
                except AttributeError:
                    mytweet = status.retweeted_status.text
            else:
                try:
                    mytweet = status.extended_tweet['full_text']
                except AttributeError:
                    mytweet = status.text
                                # Write the data to file. UTC time.
                status.user.screen_name + '_: ' + \
                str(status.user.followers_count) + '_: ' + \
                str(status.created_at) + '_:_' + \
                 '_'.join(mytweet.split()) + '\n')
    def on_request_error(self, status_code):
        print(status_code)
# Instantiate an object of class 'listener' (which is a subclass of
# 'tweepy.Stream').
mystream = \
```

```
listener (
        consumer_key,
        consumer_secret ,
        access_token,
         access token secret)
# Filter the stream for certain keywords. Here we filter tweets that
# mention a select stock index/ETF, fixed income index/ETF, or a
# commodities index/ETF. More tickers and/or keywords could be added
# here.
mystream. filter (
    track=[
         '$SPX', '$SPY', '$ES',
'$DJI', '$DJIA', '$INDU', '$YM',
         '$NQ', '$NASDAQ', '$QQQ',
         '$TLT',
         '$GC', '$GLD',
         '$NG', '$WTI'])
```

Sometimes you just want to download all the tweets from a given user. You can do this with relative ease and then save the result to a CSV file that you can later process separately (see the file code-tweepy-080-download-all-tweets-from-user.py on the course website). If you are looking for interesting people to follow, you can google "top finance twitter accounts."

```
# Twitter only allows access to a users most recent 3240 tweets
# with this method.
import tweepy
import csv
# Twitter user you want to download.
screen_name = "StockCats"
# Load Twitter API authentication data.
exec(open('../code-API-key-tweepy.py').read())
# Authorize the Twitter API.
auth = tweepy.OAuthHandler(consumer_key, consumer_secret)
auth.set_access_token(access_token, access_token_secret)
api = tweepy.API(auth)
# Initialize a list to hold all the tweets.
alltweets = []
# Make initial request for most recent tweets (200 is the maximum
# allowed count).
new_tweets = \
    api.user_timeline(
        screen_name = screen_name,
```

```
count = 200)
# Save most recent tweets to the list.
alltweets.extend(new_tweets)
# Save the id of the oldest tweet less one.
oldest = alltweets[-1].id -1
# Keep grabbing tweets until there are no tweets left to grab.
while len(new_tweets) > 0:
    print("getting_tweets_before_%s" % oldest)
    # All subsequent requests use the max_id param to prevent
    # duplicates.
    new_tweets = \
        api.user_timeline(
            screen_name = screen_name,
            count = 200,
            max_id = oldest
    # Save most recent tweets.
    alltweets.extend(new_tweets)
    # Update the id of the oldest tweet less one.
    oldest = alltweets[-1].id -1
    print("...%s_tweets_downloaded_so_far" % len(alltweets))
# Use list comprehension to transform the tweets into a 2D array
# (list of row objects) that will populate the CSV file.
outtweets = \
    [[tweet.id_str, tweet.created_at, tweet.text.encode("utf-8")]
     for tweet in alltweets]
# Write to CSV file.
with open('data-%s-tweets.csv' % screen_name, 'w') as f:
    writer = csv.writer(f)
    writer.writerow(["id", "created_at", "text"]) # Column names.
    writer.writerows(outtweets)
```

Chapter 12

AI and Machine Learning for NLP

Without going into technical detail, this chapter provides a conceptual overview of artificial intelligence (AI) and machine learning. AI and machine learning have many subfields, with perhaps the two most well-known subfields being supervised learning and unsupervised learning. What distinguishes those two approaches is whether the machine learning model/algorithm can find things out by itself (unsupervised learning) or whether it needs initial guidance (supervised learning), e.g. in the form of labeled data. In addition to supervised and unsupervised learning, we also discuss semi-supervised learning and reinforcement learning.

From the outset, it is important to clarify some terminology. There is often confusion when people talk about artificial intelligence (AI), machine learning (ML), and deep learning. The best way to think about it is that AI is the most general topic, followed by ML, followed by deep learning. In other words, **deep learning is a subfield of ML, and ML is a subfield of AI**.

Another way of jokingly thinking about the difference between AI and machine learning is as follows: If it's written in Python, it's probably machine learning. If it's written in PowerPoint, it's probably AI.

Machine learning can be very important in NLP and text analytics. The usual application of ML is that once you have converted your text into a numeric representation (e.g. BoW, tf-idf, doc2vec, or others, see Section 14.5 starting on page 101), you need to do something with it. One thing would be to classify documents (e.g. whether they are "positive" or "negative" along some dimension such as the outlook of a stock) or you could cluster similar documents together (e.g. if you're trying to figure out the different topics discussed). Most or all of these tasks require machine learning.

12.1 A Brief History of NLP and AI

There are basically two different historical approaches to NLP, which happen to parallel the developments in artificial intelligence (AI). In fact, **one could consider NLP a subfield of AI.**

Before going into the details, it is important to keep in mind that there were **two** major "AI winters," one approximately 1974–80 and another one approximately 1987–

93. Daniel Crevier writes, "it is a chain reaction that begins with pessimism in the AI community, followed by pessimism in the press, followed by a severe cutback in funding, followed by the end of serious research." Basically it is a pattern of excessive hype and the following bust that has occurred in other emerging technologies as well, e.g. the railway mania or the dot-com bubble. The general point to be taken away is (a) not to get carried away if everyone around you is euphoric and (b) not to think all is lost when everyone around you is pessimistic. Of course this is easier said than done, but it is still worth bearing in mind.

In any case, in the early days of NLP and AI (roughly the time before the mid-1980s), people tried to develop a set of logical rules (if-then rules) that tried to help understand what was going on. Often these rules were **hand-written**. Logic programming languages such as **Prolog, ASP, and Datalog** were very popular in those areas and were used to formalize the logical rules. Another very popular programming language for AI was (and still is to some extent) the venerable **Lisp**, because it supports the implementation of software that computes with symbols very well. The strand of AI that uses **logical theory** is often also termed "**symbolic AI**."

It turned out that a problem with this approach to NLP and AI is that the rules often do not generalize (e.g. if you are originally analyzing the French language and want to instead analyze the English language, you need a whole new set of rules), are often fragile to small changes (e.g. if you give it a sentence it has never seen before it might not work even close to as expected), and often require a lot of understanding of the specific problem domain (e.g. what are the quirks of the French language vs. the quirks of the English language).

After the second AI winter another approach gained traction, so-called "subsymbolic AI." The basic idea is to take a statistical approach (i.e. machine learning) and let the data speak. Suppose for example you want to write a software that translates French to English. With symbolic-AI, you would write an elaborate set of rules to translate each sentence. On the other hand, sub-symbolic AI you would for example take a large set of documents that are issued simultaneously in both languages for the European Union, and run machine learning models over both of these sets without significant human intervention. It turns out that this second approach using sub-symbolic AI works much better and requires less work, as long as you have a sufficiently large set of data to learn from.

Nowadays when people talk about AI, they are almost always referring to machine learning. However, it is important to keep in mind that AI is strictly speaking a much broader field, even though most applications nowadays are from machine learning. An interesting area of AI that combines ideas from both symbolic and sub-symbolic AI is **inductive logic programming (ILP)**, which is a subfield of machine learning that uses logic programming to represent hypotheses, examples, and background knowledge.

12.2 Supervised Learning

The basic idea is to give the model/algorithm a few examples it can learn from, and afterwards the model can perform the task by itself. For example, you have two kinds of

text documents, one is a movie review, the other is a cooking recipe. You go through a few, say, 100 documents, read them, and tell the model for each of these 100 documents what it is (a review or a recipe). The model thus has 100 examples where it knows the input (the document text) and the desired output (whether it is a review or a recipe). Based on these examples (which are called a **training set**), the model then tries to "understand" what the relation between the input and the output is. This is called **model fitting**. Once it has learned a reasonably accurate representation thereof, it can then classify documents by itself into reviews or recipes. For example, if you give the model a new text it has never seen before, the model then can tell you (with a hopefully high degree of certainty) whether this document is a movie review or a cooking recipe.

Model fitting can include both parameter estimation and variable selection. **Parameter estimation** means finding out the model parameters that guarantee the best model fit. **Variable selection** means finding out which predictor variables actually matter and which ones don't (and "select" them, i.e. keep, only those variables that matter). For example, the word "and" might not contain a lot of information in a text document. If we think about this word as a predictor variable for whether the document is about a movie review or a cooking recipe, it might actually be a good idea to remove this word from the text document because it just introduces unwanted noise.

The key thing with supervised learning is that it needs a couple of examples to learn from (the training set). So it needs some form of initial guidance, which may or may not involve human input. For example, if you want to know whether a tweet was written in a happy mood, you might have to read a few tweets yourself to determine which ones seem to be happy and which ones are not. On the other hand, if you want to identify tweets that predict a steep downward move in the stock market, you simply select a few relevant tweets that were posted before a few down moves from the past and use those as a training set. No human interaction is required here (in the sense that nobody has to manually go through these tweets and manually read them).

How large should the training set be? There is no fixed rule for that, since it always depends on the specific application at hand and the specific algorithm you use. For example, naïve Bayes typically needs relatively few examples to learn from (in some cases as few as 100), while deep learning is often said to require hundreds of thousands or even millions of observations in the training set. In any case, a general rule is that **the larger the training set**, **the better the model fit**.

A key problem we might face is **overfitting**, which means that the model fits the training dataset very well, but as soon as you provide a new data point the model has not seen before (in the training set), it fails miserably. We discuss this problem further in Chapter 13 starting on page 91, but to get an intuitive understanding let's consider the following example. Suppose for example that you run a company whose sales are a linear function of GDP plus some small random fluctuations (noise). You could fit two models, a linear model and a polynomial model. The polynomial model would have a much better fit on the training set than the linear model. But if you look at a new sample that was not used for model fitting before, the linear model would perform better. The polynomial model in this case suffers from overfitting, i.e. it seems to work great on the training sample, but not on any new data points.

A simple but very popular and powerful model from supervised learning is the naive Bayes model. It answers the question of: Given a particular piece of data, how likely is a particular outcome? For illustration we use the multinomial naive Bayes classifier from sklearn, MultinomialNB. This classifier normally requires integer feature counts, so it is very much suitable for BoW. For tf-idf it might also work (even though it is not specifically tailored for floats), but for tf-idf you might also want to try support vector machines (SVM) or linear models. In any case, here we focus on integer inputs and multinomial naive Bayes. The code below can be found in the file code-sklearn-naive-bayes.py on the course website.

```
# This script illustrates how to use the naive Bayes classifier from
# SciKit-Learn.
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import MultinomialNB
from sklearn import metrics
                                # To evaluate model performance.
# Generate random data to work on. We don't expect any great results
# from naive Bayes here because the input data is just random. In
# reality, if there is some structure in the data, naive Bayes should
# (hopefully) be able to pick it up.
X = np.random.randint(5, size=(6, 100))
y = np.array([1, 2, 3, 3, 2, 1])
# Split up the data into training and testing.
count_train , count_test , y_train , y_test = \
    train_test_split(X, y, test_size=0.33, random_state=42)
# Initialize our class. If the result is suboptimal, you can change
# the 'alpha' parameter.
nb = MultinomialNB()
                                # MultinomialNP(alpha=0.5)
# Determine internal parameters based on dataset. Pass training count
# vectorizer first, and training labels second.
nb.fit(count_train, y_train)
# Get the class labels.
nb.classes_
# Log-probability of features occurring, given a class.
nb.coef_[0]
# Make predictions of the label for test data.
pred = nb.predict(count_test)
# Test accuracy of predictions.
metrics.accuracy_score(y_test, pred)
# The confusion matrix shows correct and incorrect labels. 'labels'
# can be used to reorder the resulting matrix. On the ouput array, the
# main diagonal shows the true scores, i.e. the labels that have been
# correctly predicted. The true labels correspond to the rows while
# the predicted labels correspond to the columns. For example, if you
\# have in the second row "sci-fi" and in the first column "action"
# (e.g. if you're looking at movie reviews), then the entry in the
# second row, first column shows you the number of movie reviews that
```

```
# are about sci-fi movies but were incorrectly classified as reviews
# of action movies.
metrics.confusion_matrix(y_test, pred, labels=[0, 1])
```

12.3 Unsupervised Learning

Unsupervised learning tries to extract information from the data completely without any human supervision or input. A classic example is **clustering**, where you have a set of objects and you're trying to group similar objects together. For example, you could have a set of newspaper articles, and the clustering algorithm would try to put newspaper articles together into groups such that similar articles are in the same group. The key thing is that the algorithm looks at the data and tries to identify the groups by itself. Importantly, you do not specify these groups in advance, so for example you do not tell the algorithm that you want to group the articles into politics vs. business. The groups found by the clustering algorithm should in the end hopefully somehow be interpretable, but the interpretation might be different than what you would expect from the outset. For example, maybe instead of politics vs. business, the algorithm might find it more convincing to group the articles into what could be interpreted as art vs. science.

Another key are of unsupervised learning is **anomaly detection**. The basic idea is to find observations that do not conform to expected patterns in a dataset. An example from NLP is to find errors in text. An example from finance is to identify bank fraud.

We illustrate unsupervised learning using k-means clustering, which is one of the most popular unsupervised learning algorithms. The basic idea is that the algorithm partitions the data into several groups (so-called "clusters") without using any training data. Each cluster should contain similar data. However, one has to specify the number of clusters before running the algorithm. It is therefore sometimes useful to experiment with different numbers of groups. The code below can be found in the file code-sklearn-k-means-clustering.py on the course website.

```
# This script illustrates how to use k-means clustering in
\# SciKit-Learn.
from sklearn.cluster import KMeans
import numpy as np
# We create a numpy array with some data. Each row corresponds to an
# observation.
X = \setminus
    np.array(
        [[1, 2],
         [1, 4],
         [1, 0],
         [10, 2],
         [10, 4],
         [10, 0]
# Perform k-means clustering of the data into two clusters.
kmeans = KMeans(n_clusters=2, random_state=0).fit(X)
```

```
# Show the labels, i.e. for each observation indicate the group
# membership.
kmeans.labels_
# Ifyou have two new observations (i.e. previously unseen by the
# algorithm), to which group would they be assigned?
kmeans.predict(
      [[0, 0],
      [12, 3]])
# Show the centers of the clusters as determined by k-means.
kmeans.cluster_centers_
```

12.4 Semi-Supervised Learning

Sometimes you have a relatively small amount of labeled data and a large amount of unlabeled data. In this case, you might use techniques from semi-supervised learning. The basic idea is that you want to get a better performance than either discarding the unlabeled data and running supervised learning, or by discarding the labels and running unsupervised learning.

12.5 Reinforcement Learning

What is reinforcement learning? You can think of it as a little robot who is in a room where there are lots of balls of different colors. The robot needs to find a red ball. It can perform certain actions such as walking around, stretching out its arm, and tightening his fingers to grab something (i.e. hopefully a red ball). Whenever it performs an action, the robot interacts with the environment and gets a reward based on that interaction (e.g. if it found the red ball, its batteries get recharged). Also, after performing an action, the little robot observes how its state changed. For example, if it made a step forward, it observes that it is now standing in a different position in the room. Through trial and error, the little robot learns what to do and what not to do, e.g. bump against a wall or fall down. In a nutshell, this is what reinforcement learning is all about.

As you can see, reinforcement learning is fuzzier than supervised learning. Although there is some reward, correct input/output pairs are never presented and sub-optimal actions are never explicitly corrected. The basic idea is that sometimes, due to the complexity of many situations encountered in real life, exact methods become infeasible, so we need to find a balance between exploration and exploitation. Exploration means to take a look at uncharted territory (e.g. the little robot walks over to another corner of the room he hasn't examined before), while exploitation in this context means to take advantage of current knowledge (e.g. it has found a way to walk around on this side of the room without bumping into its walls).

There are also related questions as to what payoff the little robot gets and the strategies it should use. Well, it gets rewarded when it finds a red ball and picks it up. But there are many red balls scattered throughout the room and the little robot should try to

maximize the expected sum of *all* balls it picks up. So how should it go about to pick up the balls? Should it just go in a straight line, which might give the highest payoff in the short run? Or should it circle around the room, which might initially have a lower payoff but in the long run performs better. My point is that cutting corners might work in the short run, but not in the long run, and this fact should somehow be picked up by the little robot's reinforcement learning. It is really non-trivial how the robot's actions relate to its overall payoff, the expected sum of all balls picked up.

Another point about the payoff is that there are two kinds of payoffs. The first one is the payoff that can be "seen" by the robot. If it picks up a red ball, it gets rewarded. But there is more to that. Let's say there is one red ball left on the other end of the room and the little robot makes a step towards it. There is no immediate reward, but it is definitely (and literally) a step in the right direction, because it brings the little robot closer to finally grabbing the ball. This is a different kind of payoff because the robot is doing the right thing, even though it does not directly result in a red ball in its hand. So my point is that it is difficult to specify a concrete payoff function that gives our little robot the correct information right after each action it takes. This is what I mean when I say that it is difficult or impossible to fully specify a payoff function that rewards or punishes each decision taken by the little robot.

Related to this are also strategic contexts, where there are two or more players. For example, you could put another little robot into that room and let both robots compete to find most red balls. How should our little robot react now? Should it just try to be faster? Or should it try to push the other robot off its feet?

Competitive self-play expands on this idea. The basic idea is that in order to become better and improve, you let the little robot play against a copy of itself. The good thing is that playing against itself has just the right difficulty level. It's not too difficult (so that the little robot doesn't always loose and doesn't learn anything) and not too easy (so that the little robot doesn't always win and doesn't learn anything either). There have been big advances recently, e.g. with AlphaGo, where GU Li (a very famous Go player) said that "AlphaGo's self play games are incredible—we can learn many things from them."

How does this apply to text analytics and NLP in finance? The basic idea of reinforcement learning is that you're trying to find a solution to a very complex optimization problem. Self-play is not directly important in NLP, but it means that through exploration and exploitation you may be able to find words or combination of words that better capture what you're looking for. For example, if you are using Twitter to predict the stock market, you might not only look at the words, but you might also want to look at the person who is writing the tweets and his position in the network of followers. The relations here can be very complicated (and noisy due to the stock market's reaction), so this is a very complex optimization problem that you might be able to solve approximately with reinforcement learning.

One key challenge is that the settings of reinforcement learning are usually experimental and not observational. This means in the example above that you can clean up the room, put the little robot back into its starting position, and start all over again. However, in finance, many things are not that "simple." Instead, if we take the example of the stock market, we can only observe from the past what happened in the stock market.

We cannot just observe for example what would have happened if the financial crisis of 2007–2008 would not have occurred. Another problem is that, unless you are a big player (e.g. a large mutual fund or hedge fund), your actions really don't matter much in terms of influencing the stock market's behavior. (Actually a nice application of reinforcement learning might be the actions of a large asset manager such a hedge fund and how its stock trading decisions might influence other players in the markets.) So after all, the applications of reinforcement learning might be more limited in a financial context.

Another problem is that reinforcement learning is often formulated as a Markov decision process. The basic idea is that as long as you know today's state of nature, you don't need to know what happened yesterday. As long as the little robot knows where it stands and where the balls are, in that moment, it doesn't matter whether it got there by walking forwards or backwards in its previous move. But in financial markets, a longer history often matters. If a stock has been going up for the last year, chances are higher that it will keep going up in the future (this is called "momentum"). So a Markov decision process might not yield the optimal results because it is misspecified. The questions is of course whether this misspecification is sufficiently large to cause concern. One could argue that financial markets are relatively efficient, so according to the efficient markets hypothesis, all you need to know is today's stock price because it reflects all available information. My point is that there is some debate whether financial markets are Markovian or not, and therefore it's reasonable to assume that the degree of misspecification of a Markov process is not too large (otherwise there would be no debate).

So not all hope is lost. Depending on how much data the algorithm needs, you can still let it learn over time. For example, you give all the data available on Monday evening, let it make its decision (e.g. which stocks to buy or sell), and the reward would be how much money it made on Tuesday (the following day). You then move the agent forward to give it all the data on Tuesday; rinse and repeat. The only limitation here might be the amount of data the algorithm needs. If you're looking at daily data and we assume we have 100 years worth of data (assuming we're not using more recent data sources such as Twitter), we only have 25,500 data points (100 times 255, the number of trading days per year). If this amount of data is sufficient, reinforcement learning could potentially work.

So when should you use reinforcement learning in text analytics and NLP in finance? While I think it's a very interesting concept, you should first try more traditional machine learning methods. The basic idea is that if your problem is sufficiently simple that you can give it correct input/output pairs and if your problem space is not exceedingly complex, it might be better to apply techniques from supervised learning. Only if your problem is too complex and/or it is difficult to give a precise utility/payoff function should you consider using reinforcement learning.

12.6 Standardized Variables

When using machine learning models, it is often important to transform the input and/or output variables in order to boost the performance of the model. A seemingly trivial but often effective method is to "standardize" some variables (sometimes also called calcu-

lating the "z-score" or the "standard score"). Often machine learning algorithms work better if they are provided with standardized variables, as their statistical properties fit the algorithm better. The basic idea is to "center" the variables so that they have mean zero:

$$z = x - \mu$$
,

where *x* is the raw score and μ is the mean of the sample or population.

Additionally, you can scale a variable so that it has a standard deviation of one:

$$z = \frac{x - \mu}{\sigma}$$

where σ is the standard deviation of the sample or population.

There are several ways how to do this in Python, one of them is using the function scale() from the Scikit-Learn package. In the following example we think of each *column* as containing realizations of a random variable.

```
>>> from sklearn import preprocessing
>>> import numpy as np
>>> X_train = np.array([[ 1., -1., 2.],
                        [2., 0., 0.],
                        [0., 1., -1.]
>>> X_std = preprocessing.scale(X_train)
>>> X_std
array([[ 0. ..., -1.22..., 1.34...],
       [1.22..., 0. ..., -0.27...],
       [-1.22..., 1.22..., -1.07...]
>>> X_std.mean(axis=0)
>>> np.mean(X_std, axis=0) # Check the mean of each column.
array([0., 0., 0.])
>>> X_std.std(axis=0)
>>> np.std(X_std, axis=0) # Check the standard deviation of each column.
array([1., 1., 1.])
```

12.7 Most Popular Machine Learning Models

Below is a list of some of the most popular ML models in use today. We might not go into all of them in detail, but if you're looking for inspiration how to further analyze your text data, feel free to delve further into some of these topics. In the list below I have **highlighted in bold** some of the models that I find **most exciting and practical**. If you are wondering how to implement these models, **see Sections 9.10, 9.11, and 9.12 starting on page 53 for a list of Python packages.**

12.7.1 Supervised Learning

• Naive Bayes is a very robust supervised learning method. It does not always give the very best results, but it is usually surprisingly close to the best available

method. For text classification it is often a good idea to start with naive Bayes as it easy to use and usually gives good results in a wide variety of scenarios.

- Ordinary least squares (OLS) regression
- Logistic regression
- Support vector machines (SVMs)
- *k*-nearest neighbors (*k*-NN or KNN, approximate nearest neighbor)
- Classification and regression trees (CART)
- Ensemble methods: Bayesian averaging, bagging (e.g. **random forests**), boosting (e.g. AdaBoost, **LightGBM**, **XGBoost**). LightGBM and XGBoost are very popular these days, e.g. on Kaggle competitions. For example, LightGBM has won the M5 Forecasting Competition. It is important to keep in mind that XGBoost often does not work well if the training set is too small relative to the number of predictor variables. In NLP applications, this means for example that the number of observations in your training set should be larger than the number of terms. To achieve this, you can for example, remove terms that occur in fewer than 0.5% of all documents in order to reduce the number of terms. Furthermore, to be on the safe side, you should always compare the performance of XGBoost to other methods such as naive Bayes or deep learning.

12.7.2 Unsupervised learning

- Apriori algorithm
- Clustering algorithms: Centroid-based (e.g. *k*-means clustering), connectivity-based (e.g. hierarchical clustering), density-based (e.g. DBSCAN or OPTICS), probabilistic (e.g. Gaussian mixture models), neural nets/deep learning (e.g. self-organizing map). Note that often it is useful to perform dimensionality reduction before running a clustering algorithm!
- Dimensionality reduction: Singular value decomposition (**SVD**), principal component analysis (PCA)
- Independent component analysis (ICA)

12.7.3 Neural Networks and Deep Learning

- Neural networks and deep learning can be supervised, semi-supervised, or unsupervised.
- A neural network is a collection of neurons that are connected with each other. Neurons are typically aligned in layers. There is typically an input layer to receive external data, followed by zero or more hidden layers, followed by the output layer to produce the final result.

- Traditionally, people were looking at a special class of neural networks where there are typically one or two layers between input and output. I call them "old-school" neural networks, and they are a special case of neural networks.
- In contrast, there is a more recent special case of neural network referred to as "deep learning." These networks typically have many layers between input and output.
- The famous word2vec for example is a shallow neural network, i.e. it is not deep learning as it only has two layers. For more details on word2vec see Section 19.8 starting on page 141.
- Especially **convolutional neural networks** (**CNN**) or **recurrent neural networks** (**RNN**), maybe as LSTM RNN, can be useful for text analytics and NLP.
- Note that there is a lot of progress for LSTM RNNs, with new and potentially better solutions appearing such as Attention-based models (e.g. Transformer or hierarchical neural attention encoders).
- CNNs are often used for text classification.
- RNNs or LSTM RNNs are often used for natural language generation or machine translation.
- RNNs should not be confused with recursive neural networks. RNNs can be a special case of a recursive neural network with a linear chain.

12.7.4 Reinforcement Learning

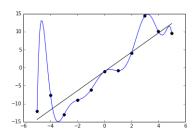
- We have discussed reinforcement learning in Section 12.5 starting on page 84.
- Reinforcement learning can be combined with deep learning. This is called deep reinforcement learning (DRL).
- Reinforcement learning is typically thought of as being distinct from supervised and unsupervised learning. It is not supervised because it does not rely on a labeled training dataset. It is not unsupervised because there is an expected reward influencing the reinforcement learning process.

Chapter 13

Overfitting and What to Do About It

Overfitting can be a big problem in NLP and machine learning in general. It occurs when you fit a model too closely to a given training dataset and that fitted model then works very badly on any new data it hasn't seen before (in the training set). We have already talked about an overfitting example on page 81.

The basic idea is that there is some "true" model, but you can't examine that model directly and instead can only see it through noisy observations of that model. For example, suppose the true model is linear as in the following figure, but all you can see are the data points:



What you could do with those data points is to fit a few models and hope that these models somehow capture the underlying "true" model.

It would be tempting to fit for example a linear model and a polynomial model to the data. As you can see, the polynomial model fits the training set's data points very well, while the linear model does not. So only by looking at the training set, it seems that you should go with the polynomial model.

Of course, if you later get new observations that have not been included in the training set, you will find out that the linear model fits much better than the polynomial model on these new data points. You thus realize that you have **overfitted the polynomial model**, because it does not work very well out-of-sample. By **out-of-sample** we refer to a model's performance on new data not in the original training set/sample.

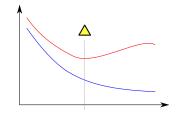
In general, overfitting can occur if you fit a model that is too general or if you have very limited training data (your training set is too small). The tradeoff is really between the true (but unknown) model and the noise. Ultimately you want to extract information about the true model and you want to ignore the noise as much as possible, but if you overfit, you end up doing the opposite, i.e. putting too much emphasis on the noise.

So how do you avoid overfitting? The basic idea is to split the dataset you have available for fitting the model into two or even three non-overlapping subsets. The terminology generally used can sometimes be a bit confusing because the terms are not always consistently applied. In any case, we use the following **conventions** in these lecture notes:

- If we are dealing with all **three sets**, we call then "training set," "validation set," and "test set."
- If we omit the optional last set and just deal with the first **two sets**, we call them "training set" and "validation set," or alternatively we call them "training set" and "test set."

The basic idea is to have three independent samples from the same population. The first dataset, the **training set** is used for model fitting as before. The second dataset, the validation set is used to validate your model in the sense that you check how well the fitted model performs on the new data, e.g. by looking at a statistic that summarizes the error of the model on the validation set. For example, you could look at various measures of goodness of fit or different loss functions for classification, such as mean squared error (MSE), hinge loss, logistic loss, or cross entropy loss. At this point in time you can iterate, for example, you can fit different models (e.g. in the example above you could fit polynomials with different degrees). As you increase the model complexity (e.g. use polynomials with higher degrees), you would typically see a decrease in the training error (on the training set), which initially is mirrored by a decrease in the validation error (on the validation set). However, as the complexity of your model increases, at some point you would start overfitting the model, which means that the validation error would start to increase (while the training error keeps decreasing). This is the point where you stop increasing the model complexity because you have found the model that performs best out-of-sample.

The figure below shows on the x-axis the model complexity, on the y-axis the error. The decreasing blue line is the training error, while the u-shaped red line is the validation error. The best predictive model is obtained at the minimum of the validation error.



Finally, to get a better understanding for how the performance of your "best" model will be like in reality, you use the third dataset, the **test set**. You can again compute the MSE or related measures. The basic idea is to compute a measure that tells us how the model is likely going to perform in reality on datasets we have not seen before. The reason why you don't want to use the validation set is that it is crucial to avoid using

any dataset that has previously been used for model building (such as the training or validation set). Otherwise you will not get an unbiased estimate of your model's true out-of-sample performance.

One way to think about this whole procedure is that **the model has parameters and hyperparameters**. Let's consider again for illustration the polynomial model, which can be written as follows:

$$\sum_{i=0}^{n} a_i x^i = a_0 + a_1 x + a_2 x^2 + \ldots + a_n x^n.$$

In this example, $\{a_0, a_1, \ldots, a_n\}$ are model parameters while n is a hyperparameter. This is an example of a model with a single hyperparameter, but there are other models with no hyperparameters (e.g. linear regression) or with several hyperparameters. In any case, the key characteristic of a hyperparameter is that it is set prior to the commencement of the model fitting. So for example, if you fit a quadratic model on the data, it means you fix n=2 before the model fitting, and then fit the model on the training set to estimate $\{a_0, a_1, \ldots, a_n\}$. The iterative procedure described above is really about pinning down the hyperparameter(s) of your model, i.e. you go back and forth between the training set and validation set until you have found the hyperparameter(s) with the best out-of-sample performance.

The polynomial model is useful for illustration, but in machine learning you often use other models, e.g. LASSO adds a regularization hyperparameter to OLS regression, or an artificial neural network for deep learning has the number of hidden units as a hyperparameter. In any case, the principles discussed for avoiding overfitting and finding the best hyperparameter(s) stay the same.

In terms of code implementation, it is often not necessary to "manually" split up a given dataset, as the Sklearn package already has this capability. The code below can be found in the file code-sklearn-train-test-split.py on the course website.

```
# This script shows how to split up your data into training and
# testing subsets.
import numpy as np
from sklearn.model_selection import train_test_split
# Create data that is for illustration evenly-spaced.
X, y = np.arange(10).reshape((5, 2)), range(5)
X
list(y)
# Split up the data. 'test_size' is the proportion of the dataset to
# be included in the test split. 'random_state' is the seed of the
# random number generator (so that you get reproducible results when
# you run the code the next time).
X_{train}, X_{test}, y_{train}, y_{test} = 
    train_test_split(X, y, test_size=0.33, random_state=42)
# Take a look at the training set.
X_train
y_train
# Take a look at the test set.
X_{test}
```

```
y_test
# Alternative way to split up the data. Do NOT shuffle the data before
# splitting.
train_test_split(y, shuffle=False)
```

Chapter 14

Main Concepts of Text Analytics and NLP

14.1 Difference Between Text Analytics and NLP

A quick note about terminology. You might be confused about the difference between NLP and text analytics (or text mining as it is sometimes called). There is in fact a great deal of overlap between both. Furthermore, the definitions of both are often a bit fuzzy. So in the end there is some room for interpretation.

Broadly speaking, NLP deals more with understanding semantics using the text's structure, e.g. the lexical relations among words. Text analytics on the other hand is more about turning text into data and running further analyses on this data. Text analytics has more of a "let the data speak" philosophy, often using machine learning techniques on collections of words. I think a useful memory hook is to think of text analytics as "text mining," because it uses "data mining" (basically machine learning) techniques on text. Another way of thinking about NLP versus text analytics is that NLP deals more with the understanding of single words (e.g. in the context of a sentence), while text analytics deals more with understanding whole text documents. As mentioned before, depending on the specific application, there can be substantial overlap between NLP and text analytics, so please take these explanations with a grain of salt.

For example, NLP deals with finding out whether a word is a noun, verb, adjective, adverb, or it figures out the gender of a word, or whether a word is singular or plural. Or you can use NLP to split a document into sentences or extract entities via namedentity recognition (NER), e.g. names of persons, organizations, locations, times and dates, quantities, monetary values, percentages, etc. On the other hand, text analytics often does not try to dig so deeply in terms of the semantics of each word and is more concerned with finding statistical patterns in the text. For example, you can find out how similar two text documents are to each other and/or group similar text documents together. Or you try to find the topic that a text document deals with, e.g. whether a newspaper article is about the weather or about business. Or you could perform sentiment analysis and for example figure out whether a twitter post has positive or negative tone.

In fact, **oftentimes NLP and text analytics build on top of each other.** For example, it is possible to tag each word in a text document (e.g. add a suffix indicating whether it is a noun or verb), and then use this tagged text as input for text analytics. This can be important, e.g. in the following sentence "dogs" is a verb: "The sailor dogs the hatch" (meaning he locks the hatch using a "dog," which is a tool to prevent movement). Or for example, if a word is negated ("I am NOT sad"), it can be converted by NLP to an antonym ("I am happy") and then fed into text analytics. Another example where the same word has different meanings (and where NLP can be useful to look at the semantics) is "accident" which usually has a negative connotation as in "I got in an accident." However, this word does not always have to have a negative meaning, e.g. "I met my wife by accident."

To avoid dealing with terminological differences that are not very relevant for our purposes, I will for the remainder of this course refer to "text analytics," "text mining," and "NLP" synonymously in the sense that all of them somehow try to "understand" the content of textual data. It will be clear from the context specifically what we are trying to accomplish in each step. Frankly, in many discussions in the media or elsewhere the terms are already used interchangeably, partly because people don't know the difference and partly because when talking about applications, it usually doesn't make much sense to split hairs about terminology.

14.2 Unstructured Data

Nowadays more than 80% of all data being created is unstructured. By "unstructured" I mean that it does not have a predefined data model or is not organized in a predefined manner. For our purposes, we can split up unstructured data into textual and non-textual data. (For a concrete list of potential data sources see Chapter 6 starting on page 33.)

Examples of non-textual unstructured data include:

- Metadata
- Health records (some of which actually may contain text as well)
- Audio
- Video
- Images, e.g. satellite image data
- Foot and car traffic (e.g. from cellphone data or other data sources)
- Ship locations

Examples of textual data (textual data is by definition unstructured), i.e. this is the kind of data we want to deal with in these lecture notes:

• Books

- Journals
- Newspapers, newswires, press releases
- Product reviews
- Legal and business documents, e.g. contracts, regulatory company filings, analyst reports
- Transcripts of conversations and meetings, e.g. conference calls, customer interactions with call centers, initial doctor-patient interviews, warranty claims, Federal Reserve Board meetings
- Open-ended survey responses, e.g. in marketing
- Word processor documents
- Web pages, e.g. company websites, blogs, message boards
- Social media footprints from Facebook, Twitter, LinkedIn, etc.
- Email messages
- Messages from WhatsApp, WeChat, etc.

In summary, we can see that unstructured data includes text, but also goes way beyond that. While analyzing unstructured data in general can be lots of fun, we will focus in this course on text only as it is already a big area in itself.

14.3 Limitations and Potential Pitfalls

Of course, it is important to keep in mind that **text analytics is not an "exact" science** in the sense that there is always room for interpretation. For example, if you use two different analysis methods on the same text document, you will receive two different answers. Furthermore, even the preprocessing steps can make a difference in the answers you obtain. Therefore it is always important to keep in mind that **working with textual data is messy**. It therefore of utmost importance to:

- 1. Obtain data of the highest possible quality.
- 2. Clean and preprocess the data with great care and attention to detail.
- 3. Conduct your analysis in a sensible way and use the right tools for the job.

Even if you have a valuable input and mess up the preprocessing and the analysis of textual data, you will end up with "garbage out." So a thorough understanding of your textual data is of absolutely crucial importance. If you are looking for a quick fix and are cutting corners, the analysis performed will more often than not be of low value added.

Another important item to keep in mind is the number of documents you have and how big each document is. For example, if you have downloaded millions of websites, you typically have relatively short documents, but you have lots of them. On the other hand, if you are analyzing books, you might have only two lengthy books (i.e. documents), but each one of them is very long. Depending on which case you are in, you might want to use different approaches to analyzing text, so it is important to keep in mind that **there** is no "magic" text analysis method that works well in all cases.

Related to this point is that it's always important to have enough data so that whatever algorithm you use actually works reliably. Even the best algorithm won't help you if you don't have enough data (or if your data is of low quality).

14.4 Typical Preprocessing Steps

Without going into technical details, I would like to delve a bit here into common preprocessing steps for text analytics. **Not all of these steps are necessary at all times; depending on the application you may omit some of them.** Furthermore, some of these steps depend on the language you are analyzing, e.g. stemming might be valid for English but not for Chinese.

- Stop word removal: This means we exclude some terms, e.g. "the," "a," "is," "at," "which," or "on" (so-called "**stop words**"). These are words that do not contribute a lot to a deeper understanding of the text in consideration as they do not have significant meaning. You can see programming examples of stop word removals in Section 15.4 starting on page 114 and in Section 19.5 starting on page 134.
- Stemming and lemmatization: Here you combine different grammatical forms of the same words, e.g. "travel," "traveling," and "traveled." Again the basic idea is to remove noisy data (in this case the suffix) that are of limited use in understanding the text. The difference between stemming and lemmatization is that stemming often uses a crude heuristic process, while lemmatization tries to do things properly with the use of a vocabulary and morphological analysis of words. For example, when considering the token "saw," stemming might just return "s" whereas lemmatization would attempt to return either "saw" or "see" depending on whether the token is used as a noun or verb. The following example in Python illustrates lemmatization (using the NLTK package; you could also use Pattern, which also has fancy lemmatization).

```
from nltk.stem import PorterStemmer
ps = PorterStemmer()
ps.stem('program')  # program
ps.stem('programs')  # program
ps.stem('programer')  # program
ps.stem('programing')  # program
ps.stem('programers')  # program
from nltk.stem import WordNetLemmatizer
```

```
wnl = WordNetLemmatizer()
wnl.lemmatize('dogs')  # dog
wnl.lemmatize('churches')  # church
wnl.lemmatize('aardwolves')  # aardwolf
wnl.lemmatize('abaci')  # abacus
wnl.lemmatize('hardrock')  # hardrock
```

• Lower case conversion means you convert all words to lowercase. This can be important, e.g. some words appear at the beginning of a sentence and are thus capitalized, but their content is the same as their lower case version.

```
>>> 'Walk' == 'walk' # To Python those strings are not the same!
False
>>> x = 'Example string.'
>>> x.lower() # Convert to lower case.
'example string.'
```

• Synonyms: For example "begin" and "commence" have nearly the same meaning and therefore could be treated the same. You could thus replace all synonyms, e.g. you would replace "commence" and "start" by "begin." For a given word (such as "begin" in the code below), you can find all its synonyms as follows:

```
from nltk.corpus import wordnet
syns = []
for s in wordnet.synsets('begin'):
    for lem in s.lemmas():
        syns.append(lem.name())
print(syns)
```

• Special words: For example "Microsoft Windows" has nothing to do with the common use of the term "Windows." Or "New York" should really be treated as a single term instead of the two separate words "New" and "York." If you end up having a lot of these kinds of words in your specific application, you can use named entity recognition (NER) as discussed in Chapter 17 starting on page 123. Alternatively you might consider concatenating them e.g. to "MicrosoftWindows" and "NewYork" to make sure they are treated correctly. Of course, this involves some hand-coding and in many applications might after all not be necessary. But it might make an important difference in some specific applications.

```
>>> s = 'New York and California'
>>> s.replace('New York', 'NewYork')
'NewYork and California'
```

• Part of speech tagging (see the example code in Chapter 17 starting on page 123): Each sentence is analyzed and tags are appended to nouns, verbs, etc. For example the sentence "And now for something completely different" becomes:

```
And/CC now/RB for/IN something/NN completely/RB different/JJ
```

CC is a coordinating conjunction, RB is an adverb, IN is a preposition, NN is a noun, and JJ is an adjective.

- Reduce the number of words in your text corpus.
 - Discard all one- or two-letter words, e.g. "a" or "to."
 - Exclude numbers.
 - For example, you could remove words that occur only in 0.5% or less of all documents. (The threshold level of 0.5% is arbitrary and can be chosen differently depending on your application.) So if a word does not occur in at least 99.5% of all documents, you remove it from your text corpus. To decide whether to remove a given word, you would cycle through all documents, check whether the word occurs in each document, and then divide the number of documents it occurred in by the total number of documents. If this number is less than 0.995, you would remove that word.
 - Alternatively you could for example only keep the 2000 most commonly-used words and discard the rest. The basic idea is that you can express yourself pretty well if you have a vocabulary of 2000 words (or even less), even though the English language for example has more than 100,000 words in total. Of course, whether you use 2000 as the cutoff depends on your specific application and you might want to use different cutoff levels.
 - Use a dictionary and only keep words that occur in this dictionary, i.e. remove a word if it does not have a matching entry in the dictionary.
- Convert the words into n-grams. Consider for example the sentence "I live in HK." If you set n = 2 (so-called bigrams), you will end up with: (I live), (live in), (in HK). So the basic idea is to not look at single words, but instead to look at **combinations of words** in order to capture more aspects of the language structure. The advantage of n-grams are that you can take care of word order, e.g. if the text says "not happy," you could capture the negation in a bigram. The disadvantage is that your vectors have a high dimensionality and tend to be very sparse. The following is an example of bigrams using the NLTK package:

```
from nltk.tokenize import word_tokenize
from nltk.util import ngrams
s = 'I live in HK.'
ngs = ngrams(word_tokenize(s), 2)
[' '.join(ng) for ng in ngs]
# Output: ['I live', 'live in', 'in HK', 'HK .']
```

An alternative implementation using TextBlob:

```
from textblob import TextBlob
ngs = TextBlob('I live in HK.').ngrams(n=2)
[' '.join(ng) for ng in ngs]
# Output: ['I live', 'live in', 'in HK', 'HK .']
```

In practice you might also consider using the ngram_range argument to Sklearn's CountVectorizer function. For an example usage of CountVectorizer, see Section 15.5 starting on page 115.

• Tokenization (see also Section 10.4 starting on page 63 for a programming example): The basic idea is to split up the whole text document into smaller parts, the so-called *tokens*. A token can be a single word, but it can also be an n-gram, a sentence, a paragraph, all hashtags (e.g. in a tweet), or you could use it to separate punctuation. Tokenization can be important to get a deeper understanding of words. Consider for example the following sentence:

```
I don't like Martin's gloves.
```

After tokenization, this sentence becomes (with tokens indicated in brackets for clarity):

```
(I) (do) (n't) (like) (Martin) ('s) (gloves) (.) You can clearly see the negation of "do" as well as the "s" to indicate possession.
```

• Whitespace elimination means that you remove excessive whitespace so that you end up having a contiguous sequence of words. Oftentimes you do not have to worry too much about whitespace elimination as it is done by the tokenizer. But if you would like to do it yourself you could use some of the following examples (the last example uses regular expressions, see Section 10.3 starting on page 61):

```
>>> x = ' Test of leading and trailing whitespace \n
>>> x.strip()
'Test of leading and trailing whitespace'
>>> x = ' Test of several whitespace \n '
>>> '.join(x.split())
'Test of several whitespace'
>>> import re
>>> re.sub('\s+', '', x)
' Test of several whitespace '
```

14.5 How Do We Go from Text to Numbers?

Ultimately we need numbers that we can use as input for further analysis, typically via machine learning. The key question is then, after preprocessing the text, how do we convert the text to numbers?

14.5.1 Binary Vector

The simplest way is to simply check whether each word is contained in the text document. If a word occurs in the document, the word gets a value of one, if it is absent it gets zero. So each document can be represented by a **binary vector**, where each vector element corresponds to a different word, indicating whether or not this word occurs in the document.

14.5.2 Bag-of-Words (BoW)

Another simple way is to count the words in each document, which is the **bag-of-words** text representation (see also Chapter 15 starting on page 113). This way, you can think of one text document being represented by a **vector containing the word counts**. And several documents are just a collection of several such vectors.

14.5.3 Word Weighting

Another way is to use a different **word weighting**. The basic idea is that instead of simply using the word counts, we give those words that are more important a higher weight while reducing the weight on the unimportant words.

A commonly-used weighting scheme is "**tf-idf**" (see also Chapter 16 starting on page 119). There are two elements to tf-idf. The first one is that the tf-idf value increases with the number of times a word appears in a document. This is basically the same idea as before in Section 14.5.2 where we were counting words using bag-of-words. The second one is that the tf-idf value decreases if the word appears very frequently in the corpus (the collection of all text documents). The basic idea is that if a word (for example "and") appears all the time throughout all documents, it probably is not that important and should receive a lower weight.

Although the main point of tf-idf is really just to weigh the words in a more informative way, you can also use it to **remove unimportant words**. One way to do this is to simply get rid of words with a very low tf-idf score.

14.5.4 Dimensionality Reduction

Another, maybe more "intelligent" way is to reduce the dimensionality of your word vectors with so-called **dimensionality reduction** techniques. The basic idea here is that you have a lot of words in each document, and you want to transform each document into another representation that does not use up so much space (so many words) but still contains approximately the same information content, or maybe even a more dense representation with unnecessary noise removed. We will go through a few important techniques in the following sections, specifically latent semantic analysis (LSA) in Section 14.5.5 as well as topic models (specifically LDA) in Section 14.5.6.

14.5.5 Latent Semantic Analysis (LSA)

What It's All About

There are many methods out there to achieve dimensionality reduction, one being **singular-value decomposition** (**SVD**). With SVD, you look into linear combinations of words that are arranged in such a way as to lose a minimum amount of information. SVD is closely related to principal component analysis (PCA), as SVD can be used to compute the PCA. (Alternatively, PCA can also be done using an eigenvalue decomposition.)

For the following discussion, it is useful to keep the following definition in mind: The matrix B is **orthogonal** if and only if $B^tB = BB^t = I$, where I is the identity matrix. This definition is equivalent to $B^t = B^{-1}$, i.e. a matrix is orthogonal if and only if its transpose is equal to its inverse. In the context of matrix notation, superscript t indicates the **transpose** of a matrix, i.e. B^t is the transpose of B.

Suppose your **term-document matrix** is given by the $m \times n$ matrix M, which means we have m terms and n documents. So the rows of M correspond to different terms while the columns of M correspond to different text documents. Using a mathematical result called "singular value decomposition" (SVD), it is possible to factorize M into

$$M = U\Sigma V^t$$
,

where:

- U is a $m \times m$ orthogonal matrix. Each **row** of U corresponds to a different **term**, and each **column** of U corresponds to a different **concept/topic**.
- Σ is a diagonal m × n matrix with non-negative numbers on the diagonal. Each diagonal element can be interpreted as corresponding to a different concept/topic in the term-document matrix.
- V^t is the transpose of an orthogonal $n \times n$ matrix V. Each **column** of V^t corresponds to a different **document**, and each **row** of V^t corresponds to a different **concept/topic**.

The diagonal entries of Σ are called the **singular values** of M. Without loss of generality, the singular values are commonly listed in descending order (in which case Σ is unique, even though U and V may not always be uniquely determined). The interpretation is that U captures information about the terms (spanning a term vector space), V^t captures information about the documents (spanning a document vector space), and both are linked to each other using the concepts/topics encoded in Σ .

In practice, you would not have to compute the SVD by hand. Instead, the computer program can create the SVD for you by computing the matrices U, Σ , and V. For an example application with programming code see Section 19.6 starting on page 138.

For **dimensionality reduction** (or "dimension reduction," as it is sometimes called), you set the smallest singular values to zero as follows:

• You create a modified $k \times k$ matrix $\hat{\Sigma}$ where you keep only the first k diagonal elements and discard the remaining elements.

- Values of k are usually chosen between 100 and 300 in text analytics applications. Keep in mind that this is a reasonable number as (i) the number of topics in a corpus is probably not much larger than 100 to 300, and (ii) in typical applications the dimensions of M are larger than 100 to 300.
- You also create the modified matrices $\hat{U} = (U_1, ..., U_k)$ and $\hat{V} = (V_1, ..., V_k)$, where $U_1, ..., U_k$ and $V_1, ..., V_k$ correspond to the first k columns of U and V, respectively.
- Then you compute the $m \times n$ matrix

$$\hat{M} := \hat{U} \hat{\Sigma} \hat{V}^t$$
.

• Note that the same \hat{M} can equivalently be computed by setting the diagonal entries below k in Σ to zero, calling this matrix $\hat{\Sigma}_0$, and computing $U\hat{\Sigma}_0V^t$, i.e. using just the original U and V. However, for computational reasons it is usually easier to use $\hat{U}\hat{\Sigma}\hat{V}^t$.

The factorization $\hat{U}\hat{\Sigma}\hat{V}^t$ is called the **truncated SVD**. The matrix \hat{M} still has the same dimensions $m \times n$ as M, but \hat{M} now has rank k. So \hat{M} is of lower rank than M (assuming that M has rank k, which corresponds to more than k elements on the diagonal of k being non-zero). Remember that the **rank of a matrix** is the dimension of the vector space spanned by its columns. In this sense, the dimension k of the vector space spanned by the columns of \hat{M} is smaller than the one spanned by M. That's why we call the calculations "dimensionality reduction."

Numerical Examples

Consider the following numerical examples. Suppose a 3×3 matrix M is given whose SVD is specified as follows by U, Σ , and V. Note how the singular values of Σ are shown in descending order on its diagonal.

$$U = \begin{pmatrix} -0.32 & 0.65 & -0.69 \\ -0.75 & -0.62 & -0.24 \\ -0.58 & 0.43 & 0.69 \end{pmatrix}$$

$$\Sigma = \begin{pmatrix} 15.75 & 0 & 0 \\ 0 & 5.11 & 0 \\ 0 & 0 & 3.27 \end{pmatrix}$$

$$V = \begin{pmatrix} -0.42 & 0.91 & 0.07 \\ -0.57 & -0.21 & -0.79 \\ -0.71 & -0.37 & 0.60 \end{pmatrix}$$

Given this SVD, we know that M satisfies

$$M = U\Sigma V^{t} = \begin{pmatrix} 4.98 & 3.96 & 1.00 \\ 2.02 & 8.02 & 9.09 \\ 5.99 & 2.96 & 7.03 \end{pmatrix}$$

For k=2, you calculate the truncated SVD as follows. Keep in mind that in the last part you need to transpose \hat{V} in order to calculate \hat{M} . Observe that \hat{M} is still kind of similar to the original M, although we have already removed some information.

$$\hat{U} = \begin{pmatrix} -0.32 & 0.65 \\ -0.75 & -0.62 \\ -0.58 & 0.43 \end{pmatrix}$$

$$\hat{\Sigma} = \begin{pmatrix} 15.75 & 0 \\ 0 & 5.11 \end{pmatrix}$$

$$\hat{V} = \begin{pmatrix} -0.42 & 0.91 \\ -0.57 & -0.21 \\ -0.71 & -0.37 \end{pmatrix}$$

$$\hat{M} = \hat{U}\hat{\Sigma}\hat{V}^t = \begin{pmatrix} 5.14 & 2.18 & 2.35 \\ 2.08 & 7.40 & 9.56 \\ 5.84 & 4.75 & 5.67 \end{pmatrix}$$

Or for k = 1 the truncated SVD is as follows. Observe that \hat{M} now becomes more dissimilar to the original M as we have removed more information, although some resemblance

to M still remains.

$$\hat{U} = \begin{pmatrix} -0.32 \\ -0.75 \\ -0.58 \end{pmatrix}$$

$$\hat{\Sigma} = \begin{pmatrix} 15.75 \end{pmatrix}$$

$$\hat{V} = \begin{pmatrix} -0.42 \\ -0.57 \\ -0.71 \end{pmatrix}$$

$$\hat{M} = \hat{U}\hat{\Sigma}\hat{V}^t = \begin{pmatrix} 2.12 & 2.87 & 3.58 \\ 4.96 & 6.73 & 8.39 \\ 3.84 & 5.21 & 6.49 \end{pmatrix}$$

Interpretation

You can interpret this new \hat{M} matrix as a new term-document matrix that now represents a latent semantic space containing the most important information about the documents in the corpus. Mathematically, you can think of SVD as an **initial rotation coupled** with scaling, or you could also think of it as a **projection into a lower-dimensional space**.

Application in Text Analytics

There are a few things to note about SVD. First of all, you can apply the SVD on a term-document matrix containing raw word counts (i.e. **BoW**) or on a term-document matrix containing weighted word counts, e.g. **tf-idf**. Second, it is important to realize that SVD is **unsupervised**, i.e. no human input is necessary for SVD to reduce the dimensionality of the text documents.

To illustrate SVD, let's assume for example that you are looking at customer reviews of cars. You might see that reviews using the term "gas-mileage" also use "economy" a lot. Or reviews using "reliability" also use the term "defects" a lot. So there are two dimensions, the first one being "gas-mileage/economy" and the second one being "reliability/defects." SVD would ideally pick up these two dimensions and realize that words within each dimension are referring to the same thing. It is kind of like using synonyms, but the difference is that this structure is picked up from the whole collection of text documents and no human input is necessary.

The basic idea of SVD is to pick up **latent semantic structure**, which means that there is some structure about the meaning of words hidden in the word counts that you crystallize with SVD. For this reason, any analysis that you perform after applying the SVD (the SVD is used to extract the latent semantics) is also called **latent semantic**

analysis (LSA) or latent semantic indexing (LSI). In addition to the following paragraphs, we discuss LSA in more detail in Section 19.6 starting on page 138.

For example, for LSA, you could look at **cosine similarity** measures to determine how similar certain documents are to each other (**document similarity**). Let's say you have identified a few newspaper articles that were published on days before a big drop in the stock market occurred. If all the sudden you find many other articles being published that are very similar according to the cosine similarity measure, you should get worried that another drop in stock prices might be around the corner.

Another application is to look at **term similarity**. It is similar to document similarity, but instead of looking at the rows of a document-term matrix you look at its columns (e.g. using cosine similarity). It can tell you how similar different terms are to each other. For example, you could use it to find terms with similar meaning.

Finally, you can also use LSA for **document retrieval**, similar to what Google does. You enter a query, which is viewed as a short document and projected into the latent semantic space generated by SVD. Then you can find the documents that are most similar to that query using document similarity, e.g. cosine similarity.

Keep in mind that **in order to work with document similarity measures such** as the cosine similarity, you actually do not need to use SVD. Instead, in principle you can apply the cosine similarity measure directly to a BoW or tf-idf term-document matrix (or document-term matrix). However, in that case it would not be called LSA or LSI because you are ignoring the latent semantic structure of the documents.

14.5.6 Topic Models

Another approach for dimensionality reduction is **topic modeling**. The basic idea is to look at co-occurrence patterns of terms corresponding to semantic topics in a collection of documents. Each document is thought of as a mixture of a number of topics (e.g. a "cats" topic and a "dogs" topic), and each word can be attributed to one of these topics with some probability. For example, "milk," "meow," and "kitten" would be cat-related with high probability, while "puppy," "bark," and "bone" would be dog-related with a high probability. It is also possible that a word is related to several topics, e.g. the word "salt" might be from a cooking-related topic (with, say 90% probability) or from a mining-related topic (with, say, 10% probability).

The most popular topic modeling approach is **latent Dirichlet allocation (LDA)**. Here we discuss the conceptual idea of LDA, while we show examples of programming code in Section 19.7 starting on page 140. (By the way, you should not confuse it with linear discriminant analysis, which is also abbreviated as "LDA.") LDA is an unsupervised machine learning model. LDA is a generalization of **probabilistic latent semantic analysis (PLSA)** (also known as **probabilistic latent semantic indexing (PLSI)**), which fulfills a similar purpose. Other relatively popular topic models are the **hierarchical Dirichlet process (HDP)** and **random projections/indexing**.

LDA (or topic modeling in general) is often associated with dimensionality reduction. The goal is similar to that of latent semantic analysis (LSA), which is to take BoW or tf-idf (which is typically high-dimensional) and convert it into another representation that

largely retains the information content but has a much lower dimension. The key benefit is that you try to extract and compress the relevant information and get rid of potential noise.

While both LDA and LSA can be used for dimensionality reduction, they have very different approaches. While LSA uses ideas from linear algebra (i.e. a matrix factorization), LDA on the other hand uses a probabilistic approach. The basic idea is to model the conditional probability that an author is using a given term, provided that he is writing about a specific topic. Because of its probabilistic nature, it is possible that a given word comes from several potential topics. For example, the word "milk" might be from a cowrelated topic or from a cat-related topic. **The dimensionality reduction then is based on the identified topics.** For example, you could keep the terms that have the highest probability of being generated conditioned on the topic. Or maybe more scientifically, you could represent a document not by its terms, but directly by its topic distribution. For example, you can represent document d by its topic distribution

$$p(z_i|d), \qquad i=1,\ldots,k,$$

where the z_i 's represent the k topics that are identified in the corpus. So you would convert each document d into a vector \hat{d} of length k:

$$\hat{d} := egin{pmatrix} p(z_1|d) \\ dots \\ p(z_k|d) \end{pmatrix}$$

The vector \hat{d} summarizes the probabilities that text document d covers the topics (z_1, \ldots, z_k) .

14.5.7 Vector Space Models and Word Embeddings

Another way to transform text to vectors are so-called **word embeddings**. The idea is to involve a mathematical embedding from a space with one dimension per word (i.e. a very high-dimensional space) to a continuous **vector space** with much lower dimensionality. The goal is to have a relatively dense representation to avoid issues with sparsity and to have a higher information denseness. **Words sharing common contexts should be located in close proximity to another in that space.**

A popular example is **word2vec**, which is an unsupervised model that transforms words based on their co-occurrence into a higher-dimensional vector space. For example, the vector for "king" minus the vector for "man" plus the vector for "women" should be close to the vector for "queen." Another popular model is **GloVe** which is similar to word2vec. For our purposes a more relevant model is **doc2vec**, which is based on word2vec but works on whole documents instead of words. Another popular way to encode sentences or short paragraphs is Google's **Universal Sentence Encoder**. We discuss word embeddings in more detail in Chapter 19 starting on page 131.

14.5.8 t-SNE

Finally, even if you end up with only a few topics (say, 20), it is still difficult to understand and visualize them. One way to further reduce dimensionality is the t-distributed stochastic neighbor embedding (**t-SNE**), which allows you to reduce the dimensions to, say, two, and then plot them for easier visualization and understanding.

For illustration, the following example is based on the Sklearn documentation. It takes four data points with three dimensions as input, and returns four data points transformed to two dimensions.

```
>>> import numpy as np
>>> from sklearn.manifold import TSNE
>>> X = np.array([[0, 0, 0], [0, 1, 1], [1, 0, 1], [1, 1, 1]])
>>> X
array([[0, 0, 0],
       [0, 1, 1],
       [1, 0, 1],
       [1, 1, 1]])
>>> TSNE(n_components=2).fit_transform(X)
array([[ 415.623
                          -0.99826646],
       [ 193.7694
                          96.35017
                                     ],
                     , -222.85101

√ 318.27267

         96.41912
                      , -125.502556 ]], dtype=float32)
```

14.6 "Terms" or "Words?"

In text analytics and NLP, you often hear people say "term" instead of "word." I think this can be a good or a bad thing, depending on the context. If you are just dealing with words, then by all means say "words."

On the other hand, sometimes you might for example be dealing with bigrams as in the example on page 100 in Section 14.4. In this case, "I live" would be a single term (consisting of two words) and you would feed this term into your further analysis. In this case, to avoid misunderstandings, it is clearly better to talk about "terms" rather than "words."

Another example is if you have transformed your document by singular-value decomposition (SVD), so the new (transformed) entries do not really correspond to single words any more. (Strictly speaking they correspond to linear combinations of other words.) In this case again it makes more sense to say "terms" to those transformed entries.

14.7 How to Analyze the Text After It Has Been Converted to Numbers

14.7.1 Representing Text as Numbers

There are a lot of things you can do after converting the documents to a numeric format (which we have described in Section 14.5 starting on page 101). We usually end up with a numeric vector representing a text document. Often each element of the vector corresponds to a specific word, e.g. in BoW or tf-idf. On the other hand, for vector space models such as doc2vec, a one-to-one relation between vector element and corresponding word does no longer exist.

The key thing to realize is that **each document is now represented by a vector**. And if we stack these vectors next to each other, we can represent the whole document collection (i.e. the corpus) using a **matrix**.

14.7.2 TDM, DTM, and Tidy Text

The matrix describing a text corpus can be represented in various ways. We will discuss three ways in this subsection. The terminology used in this section (e.g. "term-document matrix") assumes that there is a direct mapping between words ("terms") and matrix elements, e.g. as in BoW or tf-idf. Even so, from a mathematical point of view we can use all three matrix representations for a corpus from a vector space model such as doc2vec as well, even though in general vector space models are not designed to have have a direct mapping where each matrix element represents only one specific word.

The form of the matrix depends on whether we stack the vectors next to each other or on top of each other (after transposing them). If we stack them next to each other, we obtain a so-called **term-document matrix** (**TDM**) where the rows correspond to terms and the columns correspond to documents. On the other hand, if we transpose the term vectors and stack them on top of each other, we have a so-called **document-term matrix** (**DTM**) where the rows correspond to documents and the columns correspond to terms. Ultimately both matrices are equivalent to each other. For example, if you transpose a term-document matrix, you will obtain a document-term matrix (and vice versa).

Another, maybe more convenient way of representing a corpus is the **tidy text** format, popularized by the book Tidy Text Mining with R by Julia Silge and David Robinson and their tidytext R package. (The idea of tidy text data is not limited to R and can be used in other programming languages such as Python as well.) The basic idea is that for each document-word pair, you write down the word score (e.g. word count, tf-idf, etc.) next to it, like so:

Document				
	۱ -		-	
doc1		this		5
doc1		is		8
doc1		it		3

The tidy representation of text data can be very practical. There are several reasons for this. First, document-term matrices are often sparse (especially if you are using BoW), which means there are a lot of entries with zeros. Either you use special programming techniques (which can be cumbersome and sometimes slow) to deal with these sparse matrices, or you use a regular matrix representation and run out of computer memory very quickly (which is bad in general). The tidy text format representation is much more memory-friendly. Notice how **the word "is" does not occur in document doc2 and thus does not use up any space in the tidy text format**. Furthermore, it does not require any special programming techniques; after all, it's just a dense (i.e. non-sparse) table.

Second, it turns out that the tidy data format is actually relatively easy to program with. For example, in R there is a whole "tidyverse," which is a whole "universe" of software libraries with similar underlying philosophies about tidy data. "Tidy" programming principles can also be applied to Python as well, as these principles are mostly about the way we represent data, which is independent of the programming language used.

14.7.3 Processing the Numerical Text Representation

You can now do various things with this matrix (no matter in which form: TDM, DTM, tidy, or other). For example:

- You can compare the **similarity** of two documents (or two words) using the cosine similarity measure. For further explanations about cosine similarity (which can be used independently of LSA) please see the relevant part of Section 14.5.5 starting on page 103.
- You can find out which documents (or which words) form similar **clusters**. For example, if you are looking at newspaper articles, you might find three clusters (e.g. using *k*-means clustering), and upon closer inspection you might discover that these clusters roughly correspond to politics, business, and health. The point here is that a priori you don't know the clusters and learn about them from the data. For further information about clustering and unsupervised learning in general please see Chapter 12 starting on page 79.
- You can **classify** documents into different categories. For example, you could classify tweets from Twitter into "positive" or "negative" sentiment about the stock market. The difference to clustering is that with classification you know the categories (e.g. "positive" or "negative") you're interested in already from the beginning and, looking at the data, you just try to figure out which category each text document (e.g. the tweet) belongs to. For further information about classification and related techniques from supervised learning, please see Chapter 12 starting on page 79.

Chapter 15

Bag-of-Words (BoW)

The bag-of-words (BoW) model is probably the simplest way of converting text into numbers, i.e. into something the computer can more easily understand. We have already introduced this idea in Section 14.5.2 starting on page 102. The basic idea is to simply count the words in a given text document. If some words appear more frequently than others it might indicate that these words bear a particular significance for the meaning of this piece of text.

15.1 str.count and BoW

A simple word counter can be constructed using str.count, which we discussed in Section 10.1 starting on page 58. Here we assume that we already have converted a text document into a sequence of words (or "tokens") stored in a list.

```
>>> tokens = ['python', 'hello', 'python']
>>> dict([(tk, tokens.count(tk)) for tk in set(tokens)])
{'hello': 1, 'python': 2}
```

15.2 Dict and BoW

In this example we use a Python dict to store the word counts. We cycle through all words (or "tokens") and update the dictionary manually by increasing the word count.

This code results in the content of wc being

```
{'python': 2, 'hello': 1}
```

15.3 Defaultdict and BoW

This BoW example is similar to the one using a dict, but this time we use a defaultdict and initialize it with zero (i.e. a call to the int() function). We have mentioned the defaultdict class before on page 178.

```
from collections import defaultdict
tokens = ['python', 'hello', 'python']
wc = defaultdict(int)  # Initialize the word count.
for tk in tokens:
    wc[tk] += 1

This code results in the content of wc being
defaultdict(<class 'int'>, {'python': 2, 'hello': 1})
```

15.4 "Counter" Class and BoW

A simple way of calculating token frequencies is by using the Counter class from the collections module (see also the discussion on page 177). Recall that a Counter object has a similar structure as a dictionary. The code below can be found in the file code-nltk-collections-bag-of-words.py on the course website.

```
# This file shows simple examples how to calculate bag-of-words.
from nltk.tokenize import word_tokenize
from nltk.corpus import stopwords
from nltk.stem import WordNetLemmatizer
from collections import Counter
text = '''The cat is in the box.
          The cat likes the box.
          The box is over the cat. '''
# Simple example without any preprocessing.
c = Counter(word_tokenize(text))
                                 # View the counts.
c.values()
                                 # Only the word count numbers.
c.most_common(2)
                                 # The two most common words.
list(c.elements())
                                 # All the words.
# Convert to lowercase and only keep alphabetic (remove punctuation etc.)
tokens = [w for w in word_tokenize(text.lower()) if w.isalpha()]
# Keep words that are not stopwords (i.e. remove stopwords).
no_stops = [t for t in tokens if t not in stopwords.words('english')]
Counter(no_stops).most_common(2)
# Now we lemmatize the words (similar to stemming).
wnl = WordNetLemmatizer()
lemmatized = [wnl.lemmatize(t) for t in no_stops]
Counter(lemmatized).most_common(2)
```

```
# An alternative that is native to NLTK but actually an extension of
# the 'Counter' class. Another advantage is that it can plot easily
# using the matplotlib package.

from nltk.probability import FreqDist
fd = FreqDist(lemmatized)
fd # Looks like 'Counter'.
fd.plot(10) # Specify how many words to plot at most.
```

15.5 Scikit-learn Package and BoW

The scikit-learn (sklearn) package is the standard for machine learning in Python. However, it also has functionality for dealing with texts. The code can be found in the file code-sklearn-bag-of-words.py on the course website.

```
 \verb|# This code uses scikit-learn to calculate bag-of-words \\
# (BOW). 'CountVectorizer' implements both tokenization and occurrence
# counting in a single class.
from sklearn.feature_extraction.text import CountVectorizer
v = CountVectorizer(stop_words='english') # Vectorizer.
                                 # Minimal corpus for illustration.
    'The_sun_filled_the_sky_with_a_deep_red_flame.',
    'The_waves_rolled_along_the_shore_in_a_graceful,_gentle_rhythm',
    'The_winter_sun_touched_the_painting',
    'The_painting_was_a_field_of_flowers']
# Learn the vocabulary dictionary and return the document-term
# matrix. Tokenize and count word occurrences.
bow = v.fit_transform(c)
# Each term found by the analyzer during the fit is assigned a unique
# integer index corresponding to a column in the resulting
# matrix. This interpretation of the columns can be retrieved as
# follows.
                                 # Print the document-term matrix.
bow.toarray()
                                 # Same effect, shortcut command.
bow.A
                                 # Which term is in which column?
v.get_feature_names_out()
# Inverse mapping from feature name to column index.
v.vocabulary_.get('painting')
# Mapping of documents to BOW. Words that were not seen in the
# training corpus are ignored.
v.transform(['The_dry_painting.']).toarray()
```

15.6 Gensim Package and BoW

Gensim is a very powerful NLP package that can, among many other things, also do bagof-words. In the example below we use an in-memory corpus, i.e. the whole corpus is held in computer memory. On the other hand, to have a more memory-friendly implementation, we will discuss an alternative way to compute BoW in gensim in Section 19.5.1 starting on page 134.

The following code can be found in the file code-gensim-bag-of-words.py on the course website.

```
# Bag-of-words using gensim.
from gensim.corpora.dictionary import Dictionary
from nltk.tokenize import word_tokenize
# This is an example corpus consisting of movie reviews. You can think
# of each movie review as a separate text document.
cp = \setminus
    ['The_movie_was_about_a_spaceship_and_aliens._The_movie_is_wonderful!',
     'I_really_liked_the_movie._More_people_should_go_see_it.',
     `Awe some\_action\_scenes \,, \_but\_boring\_characters \,. \,\, ']
# Create tokenized corpus. Very basic preprocessing. Usually you would
# do more work here.
cp = [word_tokenize(doc.lower()) for doc in cp]
cp = [[token for token in doc if token.isalnum() and len(token) > 1]
      for doc in cp]
# Pass to gensim 'Dictionary' class. This assigns to each token
# (e.g. word) a unique integer ID. Later on we will just work with
# those IDs instead of the tokens directly because it is
# computationally easier to handle (there is a one-to-one mapping
# between both, so we are not losing any information). The reason why
# we use a dictionary is that it gives us a list of words we are
# interested in examining further. If a word is not in the dictionary
# but occurs in a document, it will be ignored by gensim.
d = Dictionary(cp)
d.token2id # Like dict(d); show mapping between tokens and their IDs.
d.token2id.get('awesome')
                                # What's the ID for 'awesome'?
d.get(0)
                                # Which token has ID=0?
d.dfs # In how many documents does each token appear? (Document frequency).
# For a single document, we can now calculate the token frequencies
# using the dictionary we just created. "Calculating token
# frequencies" means we're counting words.
d.doc2bow(cp[2])
# Next, using the dictionary we just created, we build a gensim
# corpus, which is just a bag-of-words representation of the original
# corpus. This is a nested list (a list of lists), where each list
# corresponds to a document. Inside each list we have tuples in the
# form
# (token_ID, token_frequency).
```

```
# So all we are really doing here is counting words.
cp = [d.doc2bow(doc) for doc in cp]
# This gensim corpus can now be saved, updated, and reused using tools
# from gensim. The dictionary can also be saved and updaed as well,
# e.g. if we need to add more words later on.
# Print the first three token IDs and their frequency counts from the
# first document.
cp[0][:3]
# For the first document, sort the tokens according to their
# frequency, with the most frequent tokens coming first.
sorted(
    cp[0],
                           # First document.
    key = lambda x: x[1], \# Sort by token\_frequency (second element).
                           # Most frequent first.
    reverse = True)
```

Chapter 16

tf-idf Weighting

Note that we have already introduced tf-idf from a conceptual point of view in Section 14.5.3 starting on page 102. In contrast, in this chapter we go into greater detail and demonstrate how to actually implement tf-idf using Python programming.

Let's suppose you're working in a bank and the term "loan" comes up all the time in all your text documents. In this case, "loan" might not be very meaningful since you're always dealing with loans. So you might want to down-weight these kinds of words. tf-idf can help in this situation. It stands for "term-frequency—inverse document frequency" and helps to put more weight on words that are seemingly more important (and less weight on words that are not, e.g. "loan" in the example above.) If a word appears very frequently in many of your documents (such as "loan" in the example above), it will receive a lower weighting in tf-idf.

Mathematically the idea of tf-idf translates to

$$tfidf(t,d,D) = tf(t,d) \times idf(t,D),$$

where $tf(t,d) = f_{t,d}$ is typically the number of times the term t occurs in document d (same as in bag-of-words) and

$$idf(t,D) = \log \frac{N}{|\{d \in D : t \in d\}|}$$

is the logarithmically scaled inverse fraction of the documents that contain the word, where N is the total number of documents in the corpus D (so N=|D|) and $|\{d\in D:t\in d\}|$ is the number of documents where the term t appears in. There are a couple of variants in the way tf-idf can be calculated, but the version presented here is the most important baseline case.

Taken together, you can see that if in the example above "loan" appears in many documents (so $|\{d \in D : t \in d\}|$ would be relatively close to N), it means that idf is closer to zero and thus tfidf is closer to zero as well. In total, "loan" would receive a lower weighting.

Let's consider a simple example. Suppose that tf = 8 and idf = 0.3. Then to compute the tf-idf value, you simply multiply both and obtain $8 \times 0.3 = 2.4$.

16.1 Scikit-learn and tf-idf

Here is another example showing the usage of SciKit-Learn (or sklearn for short) to compute tf-idf. Although sklearn is most often used for machine learning, it can also calculate tf-idf (or BoW as we have seen in Section 15.5). The code can be found in the file code-sklearn-tf-idf.py on the course website.

```
\# This code uses scikit—learn to calculate tf—idf. The code is very
# similar to the one used with 'CountVectorizer' (which give BoW).
from sklearn.feature_extraction.text import TfidfVectorizer
# Vectorizer. 'max_df' tells sklearn to ignore terms that have a
# document frequency higher than this threshold when building the
# vocabulary.
v = TfidfVectorizer(stop_words='english', max_df=0.9)
                                # Minimal corpus for illustration.
c = [
    'The_sun_filled_the_sky_with_a_deep_red_flame.',
    'The_waves_rolled_along_the_shore_in_a_graceful,_gentle_rhythm',
    'The_winter_sun_touched_the_painting',
    'The_painting_was_a_field_of_flowers']
# Learn the vocabulary dictionary and return the document-term
# matrix. Tokenize and count word occurrences.
tfidf = v.fit_transform(c)
# Each term found by the analyzer during the fit is assigned a unique
# integer index corresponding to a column in the resulting
# matrix. This interpretation of the columns can be retrieved as
# follows.
tfidf.toarray()
                                   # Print the document-term matrix.
tfidf.A
                                   # Same effect, shortcut command.
v.get_feature_names_out()
                                  # Which term is in which column?
# Inverse mapping from feature name to column index.
v.vocabulary_.get('painting')
\# Mapping of documents to their tf-idf scores. Words that were not
# seen in the training corpus are ignored.
v.transform(['The_dry_painting.']).toarray()
```

16.2 Gensim and tf-idf

Here we have a basic example that shows how you can calculate tf-idf using the gensim package. This code uses an in-memory corpus, meaning the whole corpus is held in computer memory. An alternative implementation using streaming data can be obtained by adapting the concepts presented in Section 19.5 starting on page 134. Streaming data allows for larger corpora to be processed, as the entire corpus is not held in computer memory.

The following code can be found in the file code-gensim-tf-idf.py on the course website.

```
# This script illustrates how to use tf-idf with gensim.
```

```
from nltk.tokenize import word_tokenize
from gensim.corpora.dictionary import Dictionary
from gensim.models.tfidfmodel import TfidfModel
corpus = ['The_movie_was_about_a_spaceship_and_aliens._It_is_wonderful!',
          'I_really_liked_the_movie._More_people_should_go_see_it.',
          'Awesome_action_scenes,_but_boring_characters.']
tokenized_corpus = [word_tokenize(doc.lower()) for doc in corpus]
d = Dictionary(tokenized_corpus)
bowcorpus = [d.doc2bow(doc) for doc in tokenized_corpus]
# All the above steps are standard, but now it gets interesting:
tfidf = TfidfModel(bowcorpus) # Create new TfidfModel from BoW corpus.
tfidf_weights = tfidf[bowcorpus[0]] # Weights of first document.
tfidf_weights[:5]
                                    # First five weights (unordered).
 # Print top five weighted words.
sorted_tfidf_weights = \
    sorted(
        tfidf_weights,
        key=lambda x: x[1],
        reverse=True)
for term_id, weight in sorted_tfidf_weights[:5]:
    print(d.get(term_id), weight)
```

Chapter 17

Named Entity Recognition (NER)

The purpose of named entity recognition (NER) can be used to identify important named entities in a text. Examples include people, places, and organizations. It can also deal with identifying other things such as dates. **NER can help answer: Who? What? When? Where?**

Besides its obvious purpose of extracting named entities, NER can also be used to increase the precision of other tasks such as classification. For example, instead of using a vector that represents the document (e.g. a BoW vector) as an input to the classifier, you can augment this vector by adding the extracted named entities occurring in the document. A very simple application for illustration: If you are trading the U.S. and UK markets, you could add an indicator variable that equals one if any entity related to New York (e.g. "New York," "NYSE," "Nasdaq," etc.) shows up in the document, and zero if any entity related to London (e.g. "London," "London Stock Exchange," "LIFFE," etc.) shows up. This concept can be generalized by adding a categorical variable with values corresponding to all possible named entities in the corpus, or various indicator variables, each representing the occurrence of a separate named entity.

Before looking at named entity extraction, we typically run **part-of-speech tagging**, **also called POS tagging**, **POS tagging**, **or POST**. (We have briefly mentioned POST in Section 14.4 starting on page 98.) POST allows for a better NER. What does POST do specifically? It tries to figure out whether a word is a proper noun, pronoun, adjective, verb, or another part of speech, based on the English grammar. For example, "NNP" in NLTK means "proper noun, singular."

Popular packages for performing NER:

- NLTK
- SpaCy
- StanfordNLP, which the recommended way to use Stanford CoreNLP (a Java library) in Python.
- Polyglot, especially if you are dealing with languages other than English.

17.1 NLTK and NER

Here is an illustration how it works in NLTK. The code can be found in the file code-nltk-NER.py on the course website. You will see when running the code that the MTR or the HK Space Museum are identified as organizations while Andy Hayler is identified as a person. The code can be found in the file code-nltk-NER.py on the course website.

```
# This file shows how to do named entity recognition (NER) using NLTK.
import nltk
sentence = '''In Hong Kong, I like to ride the MIR to visit the HK Space Museum
              and some restaurants rated well by Andy Hayler. ""
to_sentence = nltk.word_tokenize(sentence)
ta_sentence = nltk.pos_tag(to_sentence) # Tag the sentence for parts of speech.
ta_sentence[:3]
# Returns sentence as a tree, with named entities such as PERSON,
\# ORGANIZATION, etc.
nltk.ne_chunk(ta_sentence)
# Extract stems of the tree with 'NE' tags, i.e. we're getting all the
# named entities.
ner_sentence = nltk.ne_chunk(ta_sentence, binary=True) # Tags named entities as "NE
for chunk in ner_sentence:
    if hasattr(chunk, 'label') and chunk.label() == 'NE':
        print(chunk)
# If you have more than one sentences, you can adapte the above
# workflow as follows:
article = '''I like riding the MTR.
             And sometimes I visit the HK Space Museum.
             Andy Hayler rates restaurants '''
sentences = nltk.sent_tokenize(article)
token_sentences = [nltk.word_tokenize(sent) for sent in sentences]
pos_sentences = [nltk.pos_tag(sent) for sent in token_sentences]
# chunked_sentences = [nltk.ne_chunk(sent, binary=True) for sent in pos_sentences]
chunked_sentences = nltk.ne_chunk_sents(pos_sentences, binary=True) # Using a gener
for sent in chunked_sentences:
    for chunk in sent:
        if hasattr(chunk, 'label') and chunk.label() == 'NE':
            print(chunk)
# We can also plot a pie chart showing how often each named entity
# type appears in the text. For counting we use as usual a
# defaultdict.
from collections import defaultdict
import matplotlib.pyplot as plt
chunked_sentences = nltk.ne_chunk_sents(pos_sentences)
ner_categories = defaultdict(int)
for sent in chunked_sentences:
```

```
for chunk in sent:
    if hasattr(chunk, 'label'):
        ner_categories[chunk.label()] += 1

labels = list(ner_categories.keys())
values = [ner_categories.get(l) for l in labels]
plt.pie(
    values,
    labels=labels,
    autopct='%1.1f\%',  # Add percentages to chart.
    startangle=140)  # Rotate initial start angle.
plt.show()
```

17.2 SpaCy and NER

Yet another alternative to NLTK NER is SpaCy, a relatively young but fast-growing and popular Python package. In general, SpaCy is good for creating NLP pipelines to generate models and corpora. It has has extra libraries and tools, e.g. in the following link you can visualize named entities in your text online using displaCy.

- spaCy.io
- displaCy

You can do many things with spaCy, and here we are going to use it for NER. It has different entity types and often labels entities differently than NLTK. It can also deal better with informal corpora, e.g. tweets and chat messages. You can find the following code in the file code-spacy-NER.py on the course website.

```
# This script shows how to do named entity recognition (NER) with
\# spaCy.
import spacy
# Show info about model 'en'. This is a model trained on the English
# language.
spacy.info('en')
# nlp = spacy.load('en')
                               # Load the 'en' model.
# We set all the other options to 'False' since we only care about
# 'entity' here.
nlp = spacy.load('en', tagger=False, parser=False, matcher=False)
nlp.entity # Entity recognizer object, used to find entities in the text.
doc = \
    nlp(
        'Ottawa_is_the_capital_of_Canada_' +
        'and_the_residence_of_Prime_Minister_Justin_Trudeau.')
doc.ents
                                # Document attribute called 'ents'.
```

```
# Investigate labels for first entity using 'label_' attribute.
print(doc.ents[0], doc.ents[0].label_)
print(doc.ents[0].text, doc.ents[0].label_) # Same effect.
# Iterate over 'doc.ents' and print out the labels and text.
for ent in doc.ents:
    print(ent.label_, ent.text)
```

17.3 Polyglot and NER

There is also the Polyglot package. Its main advantage is that it can deal with many different languages. The following code can be found in code-polyglot-NER.py on the course website.

```
# This script shows named entity recognition (NER) with polyglot.
from polyglot.text import Text
text = '''Abraham Lincoln fue un politico y abogado estadounidense
          que ejercio como decimosexto presidente de los
          Estados Unidos de America '''
ptext = Text(text) # No need to specify language here; recognized automatically.
ptext.entities # 'entities' attribute; see a list of chunks (with label).
                               # Print each of the entities found.
for ent in ptext.entities:
    print(ent)
                             # Print the type of the (last) entity.
type(ent)
                             # Tag of (last) entity.
ent.tag
'los' in ent
                             # Check is 'los' is in the (last) entity.
'Abraham' in ent
                             # Is 'Abraham' in the (last) entity?
# List comprehension to get tuples. First tuple element is the entity
# tag, the second is the full string of the entity text (separate by
# space).
[(ent.tag, '..'.join(ent)) for ent in ptext.entities]
# The 'pos_tags' attribute queries all the tagged words.
for word, tag in ptext.pos_tags:
    print(word, tag)
```

Chapter 18

Sentiment Analysis

Sentiment in general refers to opinions, feelings, and emotions, and in a financial context to optimism and pessimism in financial markets. As such, sentiment is of subjective nature, but it is nonetheless important because it can drive human decision making in finance. A key example is that positive sentiment might put upward pressure on prices, resulting in stock returns that are at least temporarily higher.

18.1 TextBlob Sentiment Score

A very simple way to do sentiment analysis is through the TextBlob package, which is built using the NLTK and Pattern libraries. It aims to simplify text processing such as sentiment analysis, but also many other tasks such as part-of-speech tagging, noun phrase extraction, classification, and translation. TextBlob can calculate the polarity score, which captures how positive the text in a string is. It ranges from minus one (very negative) to one (very positive). It can also calculate the subjectivity score, which captures how subjective the string is. This score ranges from zero (very objective) to one (very subjective). By default, TextBlob uses the dictionary-based sentiment calculation from the Pattern library, although it can alternatively use machine learning (a Naive Bayes analyzer) that is trained on a dataset of movie reviews.

```
from textblob import TextBlob
s = TextBlob('I love NLP.').sentiment
s.polarity  # How positive the text is.
s.subjectivity  # How subjective the text is.
```

18.2 NLTK Vader

NLTK has Vader, which is very good at analyzing sentiment on social media. Vader has been integrated into NLTK based on code from vaderSentiment.

```
from nltk.sentiment.vader import SentimentIntensityAnalyzer
sid = SentimentIntensityAnalyzer()
sid.polarity_scores('I love NLP.')
```

18.3 afinn

Afinn does sentiment analysis using a dictionary. The dictionary itself is also available separately, see the Section 18.4 starting on page 128. The following example is taken from the Afinn documentation:

```
>>> from afinn import Afinn
>>> afinn = Afinn()
>>> afinn.score('This is utterly excellent!')
3.0
```

18.4 Dictionaries

There are various ways one can measure sentiment. The easiest way is to look at dictionaries (or "lexicons") containing certain vocabulary that is associated with positive or negative emotions. All you do is to count how many positive or negative words appear in a document relative to its length.

Furthermore you can divide the positive or negative score by the document length, i.e. the number of words contained in the document. The rationale is that if there are a lot of positive words in a document, it could be because the document has a positive content, or because it is simply a long document with lots of words in it (some of which will be positive). To eliminate this effect, you can divide by the total number of words in the document.

Here is a list of popular dictionaries:

- 1. SentiWordNet is a lexical resource for opinion mining based on WordNet
- 2. NRC Emotion Lexicon from Saif Mohammad and Peter Turney
- 3. Sentiment lexicon from Bing Liu and collaborators
- 4. AFINN lexicon of Finn Arup Nielsen
- 5. qdapDictionaries has various dictionaries, e.g. labMT for a happiness score (polarity)
- 6. Loughran-McDonald sentiment lexicon
- 7. Affective Norms for English Words (ANEW)

For some reason, the R community has been quite active in collecting some of these dictionaries and making them easily available, for example in the following R packages: tidytext, textdata, SentimentAnalysis, syuzhet, and quanteda.dictionaries. You can install these packages in R using install.packages, e.g. type install.packages('tidytext') to install the tidytext R package (you do not need to be root for that).

To import these dictionaries into Python, it's probably easiest to write the dictionary to a CSV file in R, then import this CSV file into Python. The following example imports the Loughran-McDonald sentiment dictionary into Python:

```
# In R:
library('tidytext')
DF = get_sentiments('loughran')
write.csv(DF, 'sentiment.csv', row.names = FALSE)
# In Python:
import pandas as pd
DF = pd.read_csv('sentiment.csv')
```

18.5 Machine Learning

Another way is based on supervised (machine) learning. Either you read a few text documents manually and assign them labels such as "positive" or "negative," or alternatively you look e.g. for market reactions (i.e. whether the market interprets something as "positive" or "negative"). You then feed these labels into a machine learning classifier of your choice (e.g. naive Bayes or others) and let it learn which text it should regard as "positive" or "negative." After a few observations for training, your machine learning classifier should then (hopefully) be able to reliably label hitherto unseen text documents as either "positive" or "negative."

The advantage of the dictionary approach is that it is really simple and therefore easy to understand and easy to interpret. People will not come charging at you and accuse you of using a "black box." Instead, even people who have no idea about text analytics and NLP can easily understand what you are doing. Using a dictionary approach can therefore be useful if your application requires convincing someone that what you are doing makes sense.

The advantage of the machine learning approach is that (depending on the quality of the training data) it might be more accurate. Another advantage is that it can deal better with different contexts. For example, the vocabulary used in financial applications might be quite different from applications in psychology. As a consequence, if you use dictionaries that are not specifically made for the context you're dealing with, you might misclassify some texts. Using the machine learning appraoch lets you avoid many of these issues, assuming that you can get hold of a good and suitable training set.

Chapter 19

LSA, Topic Models, and Word Embeddings in Gensim

19.1 Overview of Gensim

- Gensim is very versatile, e.g. we have used it before for bag-of-words in Section 15.6 starting on page 115 or for tf-idf in Section 16.2 starting on page 120.
 Please take a look at these earlier examples to refresh your understanding of how gensim represents tokens (i.e. words, usually) by integer IDs. You can also take a look at Gensim's introduction to refresh your memory on how documents are represented by vectors in Gensim.
- But Gensim can do much more than BoW and tf-idf. It is a very popular Python library for dealing with LSA, topic models, and word embeddings.
- In Gensim, a document is usually represented by the features extracted from it, not by its "surface" string form. This means that it is up to you how you get the features. This is intentional, since there are a lot of ways to process documents and the creator of Gensim decided not to constrain it in this regard.

19.2 In-Memory Corpus

An easy way to store a text corpus is a list of strings, with each list entry representing a different text document. The advantage is that this corpus is conceptually easy to understand. The disadvantage is that the whole corpus is held in computer memory, which can be a problem if the corpus grows big. We have discussed how to compute BoW using an in-memory corpus in Section 15.6 starting on page 115 or tf-idf in Section 16.2 starting on page 120. On the other hand, we discuss a more memory-efficient alternative using streaming data in Section 19.5 starting on page 134.

```
documents = \
    ["Human machine interface for lab abc computer applications",
    "A survey of user opinion of computer system response time",
```

```
"The EPS user interface management system",
"System and human system engineering testing of EPS",
"Relation of user perceived response time to error measurement",
"The generation of random binary unordered trees",
"The intersection graph of paths in trees",
"Graph minors IV Widths of trees and well quasi ordering",
"Graph minors A survey"]
```

19.3 Saving and Loading a Corpus

Often you have created a corpus and would like to save it to disk. Or you would like to load it from disk. The following code illustrates how this can be done in gensim. Here we assume the corpus is already available in vector form, i.e. there is a token ID (representing the word/roken) and a corresponding numerical value, e.g. from BoW or tf-idf. The code below can be found in the file code-gensim-060-save-corpus-to-file.py on the course website.

```
from gensim import corpora
```

```
# Create a very simple corpus (just a list) consisting of two
# documents. The first document has one word with token ID of 1 and
# value of 0.5 while the second document is empty.
corpus = [[(1, 0.5)], []]
# Save corpus in Matrix Market format. This is a relatively popular
# format.
corpora.MmCorpus.serialize('data-gensim-corpus.mm', corpus)
# Save corpus in SVMlight format.
corpora.SvmLightCorpus.serialize('data-gensim-corpus.svmlight', corpus)
# Save corpus in Blei's LDA-c format.
corpora.BleiCorpus.serialize('data-gensim-corpus.lda-c', corpus)
# Save corpus in GibbsLDA++ format
corpora.LowCorpus.serialize('data-gensim-corpus.low', corpus)
# You can also read the corpus back into Python. Here you load a
# corpus iterable from a Matrix Market file. We then print it in a
# couple of different ways. The first way just shows some meta
# information about the corpus (after all, it's a stream). The second
# way loads it entirely into memory and prints it. The third way
# prints it one document at a time in a memory-friendly way.
corpus = corpora.MmCorpus('data-gensim-corpus.mm') # An iterable.
print(corpus)
print(list(corpus))
```

```
for doc in corpus:
    print(doc)
```

19.4 Conversion of Corpora to/from NumPy/SciPy

Sometimes a corpus might be saved in a matrix using NumPy or SciPy (see also Section A.13 starting on page 183). It is possible to convert back and forth to/from the gensim corpus format. Here we again assume each document is already represented by token ID and value pairs. The code below can be found in the file code-gensim-080-numpy-scipy.py on the course website.

This script illustrates the conversion between numpy/scipy matrices # and gensim corpora.

```
import gensim
import numpy as np
# Use a random matrix as an example. This is in fact a dense matrix
# from numpy, which is not as efficient as a sparse matrix (which we
# will discuss further below). The basic idea during the conversion is
# that the numpy matrix has stored the word counts (or whatever vector
# space we're talking about here) such that a text document
# corresponds to a given COLUMN. On the other hand, when you print the
# gensim corpus, each text document corresponds to a given ROW (at
# least when you're printing it in the IPython command shell).
m_np = np.random.randint(10, size = [5, 2])
corpus = gensim.matutils.Dense2Corpus(m_np)
list(corpus)
for doc in corpus:
    print(doc)
# Here we convert the the corpus back into a dense numpy 2D array.
1 = \max(len(doc) \text{ for doc in corpus}) \# Generator epression.
gensim.matutils.corpus2dense(corpus, num_terms=1)
# Here we use a sparse matrix from SciPy. This can be much more memory
# efficient than a dense matrix, especially when many matrix elements
# are zero (as is often the case in text mining).
import scipy.sparse
m_{sp} = scipy.sparse.random(5, 2, density=0.5)
                # Take a look.
m_sp.todense()
                 # Take a look (same as above).
m sp.A
corpus = gensim.matutils.Sparse2Corpus(m_sp)
list (corpus)
for doc in corpus:
    print(doc)
```

```
# Convert corpus back to 'scipy.sparse.csc' matrix.gensim.matutils.corpus2csc(corpus)
```

19.5 Streaming Corpus

To avoid the problem of holding the whole corpus in memory, you can stream the corpus. This means that instead of holding it in memory, you only hold one document at a time in memory and process the corpus step by step. This approach uses **iterators**, which are **discussed in Section A.14** starting on page 187.

In the following example code we convert each document to lowercase and return a list of words in that document. Here we assume that data-gensim-mycorpus.txt is **a file holding one document per line** (see the course website for a copy of that file).

```
class MyCorpus(object): # Class that instantiates an iterable.
    def __iter__(self): # Define a generator function.
        for line in open('data-gensim-mycorpus.txt'):
            yield line.lower().split()
```

Now you can create a corpus object **without** loading the whole corpus into memory:

```
mycp = MyCorpus() # Create iterable by instantiating the 'MyCorpus' class.
```

If you want to take a look at the corpus, you can iterate over it and print each document, one at a time:

```
for doc in mycp:
    print(doc)
```

19.5.1 BoW Using Streaming Corpus

A more complete example can be found in the following script. The code assumes that all documents are contained in a single text file, with each line containing a separate text document. For simplicity, we only convert to lowercase and use a very simple tokenizer that splits on whitespace. (In production, you would use a for example NLTK to tokenize as discussed in Section 10.4 starting on page 63 or use gensim's tokenizer as illustrated in Section 19.5.2 starting on page 136.) Finally, we calculate BoW. All of this is done using iterators, so we can stream the whole corpus through memory without having to load the whole corpus all at once. The code below can be found in the file code-gensim-040-streaming.py on the course website.

```
# This script shows how to stream text documents using iterators. The # key advantage is that we do not need to hold all the documents # (i.e. the whole corpus) in memory. Instead, we process it chunk by # chunk using iterators.
```

```
# In the first part, we show how to generate and modify a gensim
# 'Dictionary' using iterators.
# In the second part, we show how to stream a corpus and calculate bag
# of words.
from gensim import corpora
# Construct dictionary WITHOUT loading all texts in memory. Instead,
# we chunk over the documents step by step using a generator
# expression. Once we're done with a document, the memory is freed.
dictionary = \
    corpora. Dictionary (
        line.lower().split() for line in open('data-gensim-mycorpus.txt'))
# Get the IDs in gensim that correspond to stopwords.
stoplist = set('for_a_of_the_and_to_in'.split())
stop_ids = \
    [dictionary.token2id[stopword] for stopword in stoplist
     if stopword in dictionary.token2id]
# Obtain the token IDs for words that appear only ONCE in the text
# corpus (these words will be removed later). 'iteritems' from the six
# library is used here to iterate over the keys and values of the
# dictionary 'dictionary.dfs' in a way that works both with Python 2
# and 3. (In Python 3 you would use 'dictionary.dfs.items()'.)
# 'dictionary.dfs' is a dict with the token ID and its document
# frequency (i.e. the frequency with which it occurs in the text
# corpus as a whole).
from six import iteritems
once ids = \setminus
    [tokenid for tokenid, docfreq in iteritems(dictionary.dfs)
     if docfreq == 1]
# Remove tokens that are stopwords or words that appear only once (or
# both).
dictionary.filter_tokens(stop_ids + once_ids)
# Remove gaps in ID sequence after words that were removed.
dictionary.compactify()
# Show how many words ("tokens") the dictionary contains, show their
# mappings to integer IDs, and show their document frequency (i.e. how
# often they appear in the text corpus).
print(dictionary)
print(dictionary.token2id)
print(dictionary.dfs)
```

```
# Using the 'dictionary' created above, we can also stream a corpus
# into gensim and calculate BoW. Here we define the '__iter__()'
# method as a generator function. As a consequence, an instance of
# 'MyCorpus' is an iterable. We iterate through a file under the
# assumption that each line holds one document. If this assumption is
# not satisfied (e.g. your file has a different format to hold the
# text documents), you can modify the definition of the '__iter__'
# method to fit your input format.
class MyCorpus(object):
    def __iter__(self):
                                # Define generator function.
        for line in open('data-gensim-mycorpus.txt'):
            yield dictionary.doc2bow(line.lower().split())
# Printing the corpus just shows the address of the object in
# memory. To see all the documents (represented as vectors using
# 'doc2bow' above), we iterate over the corpus. The important thing is
# that only a SINGLE vector resides in memory at a time!
                      # Create iterable (by instantiating the class).
mycp = MyCorpus()
print(mycp)
for vector in mycp:
    print(vector)
# Another way to print a corpus is to load it entirely into memory.
print(list(mycp))
```

19.5.2 BoW and Truncated SVD Using Streaming Corpus

Here is another example using iterators. In this example, the assumption on the data is that **each document is stored in a separate text file** ending with .txt. We also use gensim's tokenizer gensim.utils.tokenize to split up each document into separate words (tokens). We then calculate BoW and finally conclude with an example application of truncated singular value decomposition (SVD). The point of writing this script is to show how all of these steps (including truncated SVD) can be done by steaming data without overloading the memory. The code below can be found in the file code-gensim-50-streaming-with-iterators.py on the course website.

```
# This script shows how you can construct a streaming corpus and then
# pass it on to functionality in gensim. The basic idea is that the
# corpus is not held in memory all at once, but instead processed
# record-by-record. For the purpose of this script, we assume that one
# document corresponds to one file on disk. At the end of the script
# we show an illustration using singular value decomposition.
import gensim, os

def iter_documents(top_directory):
    """This is a generator function: Iterate over all documents, yielding
```

```
one tokenized document (=list of utf8 tokens) at a time. Each
    document corresponds to one file. The generator function finds all
    '.txt' files, no matter how deep under 'top_directory'.
    for root, dirs, files in os.walk(top_directory):
        for fname in filter (lambda fname: fname.endswith('.txt'), files):
            # Read each '.txt' document as one big string.
            document = open(os.path.join(root, fname)).read()
            # Break document into utf8 tokens.
            yield gensim.utils.tokenize(document, lower = True, errors = 'ignore')
# Here we print all tokens for all documents. 'iter_documents()'
# creates a generator iterator that yields one text document at a
# time. Keep in mind that 'gensim.utils.tokenize' itself returns a
# generator iterator for each document, so we have to iterate again
# through 'doc tokens'.
for doc_tokens in iter_documents('.'):
    print('\nNext_document:\n')
    for token in doc_tokens:
        print(token)
class TxtSubdirsCorpus(object):
    """This class instantiates an iterable: On each iteration, return
    bag-of-words vectors, one vector for each document. Process one
    document at a time using generators, never load the entire corpus
    into RAM.
    def __init__(self, top_dir): # Constructor method.
        self.top_dir = top_dir # Save the top-level directory name.
        \# Create a dictionary, which is a mapping for documents to
        # sparse vectors.
        self.dictionary = gensim.corpora.Dictionary(iter_documents(top_dir))
    def __iter__(self):
                               # Define a generator function.
        for doc_tokens in iter_documents(self.top_dir):
            # Transform tokens (strings) into a sparse bag-of-words
            # vector, one document at a time.
            yield self.dictionary.doc2bow(doc_tokens)
# Create the streamed corpus of sparse document vectors. 'corpus' is
# an iterable.
corpus = TxtSubdirsCorpus('.')
# Print the corpus vectors. Each vector is sparse and corresponds to
# the contents of one text document.
```

```
for vector in corpus:
    print(vector)
# Run truncated Singular Value Decomposition (SVD) on the streamed
# corpus. In this case we give a hint to the SVD algorithm to process
# the input stream in groups of 5000 vectors. This is called
# "chunking" or "mini batching." The return values are as follows: 'U'
# contains the left-singular vectors (encoded as a matrix where the
# columns correspond to documents), while 'Sigma' contains the
# singular values of the corpus (encoded as a vector).
from gensim.models.lsimodel import stochastic_svd as svd
U, Sigma = \setminus
    svd(
        corpus,
        rank=200,
        num_terms=len(corpus.dictionary),
        chunksize=5000)
```

19.6 Latent Semantic Analysis (LSA)

We have discussed LSA (and how it works together with SVD) before in Section 14.5.5 starting on page 103. Here we go into more details and show how it works in Gensim. Gensim internally uses the BLAS libraries for calculating the singular value decomposition (SVD), so if you run this code on a multi-core machine, BLAS takes care of parallelizing the computations to speed up the code.

A big advantage is the "online" feature (vs. "batch" feature) of LSA. Suppose the nature of the input stream of documents changes and there is **topic drift**. For example, you've been following the financial press and people were originally talking about financial companies (e.g. banks) but now prefer to talk about tech companies. In that case, LSA re-orients itself to reflect these changes in a relatively small amount of updates. This makes LSA more suitable for text analytics where topics can change dynamically. In contrast, LDA (see Section 19.7) can be much slower to react to these changes and therefore it might make sense to run LDA in batch mode if there is topic drift (i.e. run it on the whole corpus and there are not changes to the corpus afterwards).

The data is based on a random download of parts of the English Wikipedia. Please see the instructions in Chapter 6 in the "Wikipedia" bullet point on how you can download the data yourself from Wikipedia. You still need to run the following command (from the Linux command line) on it to convert the articles to plain text and store the results as sparse BoW or tf-idf vectors:

```
$ python3 -m gensim.scripts.make_wiki
```

As always, when working with data, you need to monitor your data quality and be careful not to fall into the garbage in, garbage out trap. Apparently there are lots of templates listed in the Wikipedia download. Furthermore, some websites seem to be

influenced by bots that import databases of cities, countries, etc. So if you would get some real work done with this Wikipedia data, **further preprocessing and cleaning would be required**.

In the code below, $1\sin[mm]$ (after converting it to a dense matrix with corpus2dense) corresponds to $U^{-1}M$ from the SVD given by $M = U\Sigma V^t$. And mm in the code below corresponds to the matrix M. (For a primer on SVD, see Section 14.5.5 starting on page 103.) This implies that $V^t = \Sigma^{-1}U^{-1}M$, or $V = (U^{-1}M)^t\Sigma^{-1} = 1\sin[mm]^t/\Sigma$, where we have used the fact that Σ is a diagonal matrix. This is the formula we use in the code below to compute V.

The code listed below for computing LSA with Gensim can be found in the file code-gensim-090-LSA.py on the course website.

```
# This script illustrates how to use latent semantic analysis (LSA)
# using the Gensim package. The data is a random extract from the
# English Wikipedia.
import logging, gensim
logging.basicConfig(
    format='%(asctime)s_:_%(levelname)s_:_%(message)s',
    level=logging.INFO)
# Load the ID to word mapping, i.e. the dictionary. This has been done
# in a previous step, when preparing the corpus with
# 'gensim.scripts.make_wiki'.
id2word = \
    gensim.corpora.Dictionary.load_from_text(
        '../data-wiki-en/wiki_en_wordids.txt.bz2')
# Load the corpus iterable.
mm = gensim.corpora.MmCorpus('../data-wiki-en/wiki_en_tfidf.mm')
# Take a look at basic information about the corpus.
print (mm)
# Compute LSA (or LSI as it is sometimes called) of the English
# Wikipedia.
lsi = \
    gensim.models.lsimodel.LsiModel(
        corpus=mm,
        id2word=id2word,
        num_topics=400)
# Print the words that contribute most (both positively and
# negatively) for each of the first ten topics.
lsi.print_topics(10)
# Recover the matrices of the SVD. See
# https://github.com/RaRe-Technologies/gensim/wiki/recipes-&-faq for
# details. Keep in mind that the matrix V is not stored explicitly by
# gensim because it may not fit into memory, as its shape is 'num_docs
# * num_topics'. However, you can manually compute it as shown below
# to get it as a 2-dimensional numpy array (i.e. a matrix).
```

```
U = lsi.projection.u
Sigma = lsi.projection.s
V = \
    gensim.matutils.corpus2dense(
        corpus=lsi[mm], # Use gensim's streaming 'lsi[corpus]' API.
        num_terms=len(lsi.projection.s)).T / \
    lsi.projection.s
```

19.7 Latent Dirichlet Allocation (LDA)

We have discussed LDA in Section 14.5.6, starting on page 107. The idea is that you can think of words in a given document coming from a set of topics. For example, if you see the word "meow," it is likely to come from a topic related to cats, while "bark" comes from a dogs-topic. The word "milk" might be from a cow-related topic (with, say 80% probability) or from a cat-related topic (with, say, 20% probability).

In this chapter we focus on gensim, but keep in mind that you can do LDA with other packages as well, such as lda or sklearn, specifically sklearn.decomposition.LatentDirichletAllocation.

You can run LDA in online or in batch mode. **Online mode** means that you have an input stream of training documents and update LDA sequentially based on the new documents observed. **Batch mode** on the other hand means that you go over the entire corpus when creading the LDA model and there are no additional documents encountered afterwards. In contrast to LSA (see Section 19.6), where online mode usually works fine, you need to be **careful when using LDA in online mode**. The problem is that if the properties of the input stream change (e.g. the documents were about financial companies, now they are about tech companies), the impact of later updates on the LDA model gradually diminishes. This means that the quality of the LDA's model fit to the data decreases and LDA effectively gets confused and increasingly unsuitable to work with the updated data. In that case, you might want to **run LDA in batch mode** where the entire training corpus is known beforehand and thus does not exhibit topic drift by definition. However, running LDA in batch mode might take a longer time than running it in online mode.

In the following example, the data is the same as in Section 19.6, please take a look there how to obtain the data and the caveats mentioned there about the data quality. The code below can be found in the file code-gensim-095-LDA.py on the course website.

```
# This script illustrates how to use latent Dirichlet allocation (LDA)
# using the Gensim package. The data is a random extract from the
# English Wikipedia.

import logging, gensim
logging.basicConfig(
    format='%(asctime)s_:_%(levelname)s_:_%(message)s',
    level=logging.INFO)
```

```
# Load the ID to word mapping, i.e. the dictionary. This has been done
# in a previous step, when preparing the corpus with
# 'gensim.scripts.make_wiki'.
id2word = \
    gensim.corpora.Dictionary.load_from_text(
        '../data-wiki-en/wiki_en_wordids.txt.bz2')
# Load the corpus iterable.
mm = gensim.corpora.MmCorpus('../data-wiki-en/wiki_en_tfidf.mm')
# Take a look basic information about the corpus.
print (mm)
# Extract 100 LDA topics, using 1 pass and updating once every 1 chunk
# (10,000 documents).
lda = \
    gensim.models.ldamodel.LdaModel(
        corpus=mm,
        id2word=id2word,
        num_topics=100,
        update_every=1,
        chunksize=10000,
        passes=1)
# Print the most contributing words for 10 randomly selected topics.
lda.print_topics(10)
# Extract 100 LDA topics, using 20 full passes, no online updates.
lda = \
    gensim.models.ldamodel.LdaModel(
        corpus=mm,
        id2word=id2word,
        num_topics=100,
        update_every=0,
        passes=20)
# A trained model can used be to transform new, unseen documents
\# (plain bag-of-words or tf-idf count vectors) into LDA topic
# distributions. Here 'doc_bow' would be a set of new documents.
doc_lda = lda[doc_bow]
```

19.8 Word2vec

Word2vec is a very popular way to **convert words into vectors** in a meaningful way. We have already discussed it briefly in Section 14.5.7 starting on page 108. The analysis is based on the **co-occurrence of words** in each text document. The algorithm learns automatically some inherent relations between the words. Each word is transferred into a vector in a larger vector space using unsupervised learning (so no human intervention is necessary, besides maybe tuning some parameters). The basic idea is that ideally you

can do calculations in this vector space as follows:

```
queen = king - man + woman.
```

However, one has to keep in mind that doc2vec really works on the word level, not the document level. So it is not directly comparable to BoW or tf-idf because those methods yield one vector per document, while word2vec yields one vector per word. There are also alternatives to word2vec, such GloVe (for a comparison see here), but they don't seem to perform that much better.

So how does word2vec actually work? The basic idea is that it chains two or three layers in a neural network together. There is an input layer, a hidden layer, and an output layer. (Word2vec is therefore not deep learning, see Section 12.7.3 starting on page 88.) There are two alternative ways this works, the first one being called **continuous bag of words (CBOW)**, which tries to use the word context to predict a so-called target word. For example, if you give it "I love" then it would (hopefully) predict that the next word is "NLP." The second way is called **skip-gram** and tries to predict a target context. For example, if you give it the word "love," it will (hopefully) predict that the word before is "I" and the following word is "NLP."

So you see that word2vec really looks at the co-occurrence of words. Specifically, it **takes word order into account**. It is similar to performing BoW of tf-idf on n-grams, but it may perform better (depending on the application) because it does not suffer from the high degree of sparsity and the high dimensionality that BoW on n-grams is plagued with.

The code below can be found in the file code-gensim-100-word2vec.py on the course website.

```
# This script contains a basic example on how to use word2vec in
# gensim.
# Import modules and set up logging.
import gensim, os, logging
logging.basicConfig(
    format='%(asctime)s_:_%(levelname)s_:_%(message)s',
    level=logging.INFO)
# word2vec expects a sequence of sentences as its input. In this
# example, each sentence is a list of words. This is fine, but
# everything is in RAM so if you have lots of sentences this could be
# a problem. Iterables to the rescue! In fact, all gensim requires is
# to get the sentences sequentially, e.g. using an iterable (see
# further below).
sentences = [['first', 'sentence'], ['second', 'sentence']]
# Train word2vec on the two sentences.
model = gensim.models.Word2Vec(sentences, min_count=1)
# Here we assume the input is on several files on disk, one sentence
# per line. Using an iterable, gensim can process each input file
```

```
# line-by-line. It yields one sentence after another.
class MySentences(object): # This class instantiates an iterable.
    def __init__(self, dirname): # Constructor method.
        self.dirname = dirname
    def __iter__(self): # Here we define a generator function.
       for fname in filter (lambda fname: fname.endswith('.txt'),
                            os.listdir(self.dirname)):
            for line in open(os.path.join(self.dirname, fname)):
                yield line.split() # Yield one sentence, split into words.
# Here we get a memory-friendly iterable and hand it over to gensim's
# word2vec. '.' refers to the current directory. Gensim will run
# several passes over the iterable. The first one is to collect words
# and frequencies to build an internal dictionary structure. The later
# ones are to train the neural model.
sentences = MySentences('.')
                              # Instantiate the class. Create iterable.
model = gensim.models.Word2Vec(sentences)
# If for some reason your input stream is non-repeatable, you can
# perform the steps mentioned above manually. This code is just for
# illustration, it will NOT work in this script as we have not defined
# the 1-pass generators here.
model = gensim.models.Word2Vec(iter=1) # Empty model, no training yet.
model.build_vocab(some_sentences) # Can be a non-repeatable, 1-pass generator.
model.train(other_sentences) # Can be a non-repeatable, 1-pass generator.
\# If you want to fine-tune the training, you can change some
# parameters.
model = \
   gensim.models.Word2Vec(
        sentences,
                         # Prune infrequent words. Default is 10.
       min_count=10,
                        # Size of neural net layers. Default is 100.
        size = 200,
                        # Workers for parallelization. Default is 1.
        workers=4)
# You can save and load the model to/from file.
model.save('mymodel')
new_model = gensim.models.Word2Vec.load('mymodel')
new_model.train(more_sentences) # You can continue to train it with more sentences.
# Originally, word2vec was released by Google and was written in
# C. You can read from the format used by that implementation as well.
model = Word2Vec.load_word2vec_format('/tmp/vectors.txt', binary=False)
# Using gzipped/bz2 input works too, no need to unzip.
model = Word2Vec.load_word2vec_format('/tmp/vectors.bin.gz', binary=True)
```

```
# You can find the most similar words. Here the output might be
# something like "[('queen', 0.50882536)]".
model.most_similar(
    positive = ['woman', 'king'],
    negative=['man'],
    topn=1)
# You can check which word does not match. Here the output might be
model.doesnt_match("breakfast_cereal_dinner_lunch";.split())
# You can also check the similarity between different words. Here the
# output might be a number such as 0.74.
model.similarity('woman', 'man')
# If you need the raw output vectors.
                                # Raw NumPy vector of a word.
model['computer']
# If you want them en-masse as a two-dimensional NumPy matrix.
model.syn()
```

19.9 Doc2vec

Doc2vec is an extension of word2vec. We have already discussed it briefly in Section 14.5.7 staring on page 108. The basic idea is to extend word2vec (which works on the word level) to work on larger blocks of texts, such as sentences, paragraphs, or entire documents. The goal is that similarly to BoW and tf-idf, you want to **generate one vector per document**. Similar to word2vec, doc2vec takes word order into account.

As the name suggests, doc2vec is related to word2vec. One could extend word2vec in various ways. One simple way would be to simply take the average over all word vectors that occur in a document. However, this might be a bit crude, so what doc2vec does is to add additional features (variables) to the word2vec neural network representing document IDs.

The script below shows basic usage; for a more complete example see this Jupyter Notebook. The code below can be found in the file code-gensim-120-doc2vec.py on the course website.

```
# This script illustrates doc2vec in gensim. Actually this is more of
# a collection of code snippets, not really a runnable script. In this
# example, we assume that we're interested in sentence—level data (not
# document—level data; in any case, the code would stay very similar
# if you update it to work on whole documents.)

# The input to Doc2Vec is an iterator of 'LabeledSentence'
# objects. Each such object represents a single sentence, and consists
# of two simple lists—— list of words and a list of labels. WHY DO WE
# NEED LABELS? Because labels in doc2vec act the same way as words in
# word2vec.
sentence = \
```

```
LabeledSentence(
                                # Or use 'TaggedDocument'.
        words=['some', 'words', 'here'],
        labels = [ 'SENT_1'])
# Here is an example to read text from a file with one sentence per
# line, using the following class as training data. In principle, one
# could have SEVERAL labels per sentence, but the most common
# application is probably to have ONE label per sentence, as shown
# here.
class LabeledLineSentence(object):
    def __init__(self, filename): # Constructor method.
        self.filename = filename
    def __iter__(self):
                                # __iter__() method implemented as generator function
        for uid, line in enumerate(open(filename)):
            yield LabeledSentence(words=line.split(), labels=['SENT_%s' % uid])
# You can manually control the learning rate as follows, which might
# in some cases yield better results. Or you could randomize the order
# of the input sentences.
model = Doc2Vec(alpha=0.025, min alpha=0.025) # Use fixed learning rate.
model.build_vocab(sentences)
for epoch in range(10):
    model.train(sentences)
                                  # Decrease the learning rate.
    model.alpha = 0.002
    model.min_alpha = model.alpha # Fix the learning rate, no decay.
# You can save and load Doc2Vec instances the usual way.
model = Doc2Vec(sentences)
# Store the model to mmap-able files (can map files into memory).
model.save('my_model.doc2vec')
# Load the model back.
model_loaded = Doc2Vec.load('my_model.doc2vec')
# Since labels in doc2vec act in the same way as words in word2vec. So
# to get the most similar words/sentences to the first sentence
# (i.e. label 'SENT_0'), you could type:
print(model.most_similar('SENT_0'))
print(model['SENT_0'])
```

Chapter 20

Summarizing, Displaying, and Visualizing Your Results

Once you have finished your analysis of the textual data, it is important to present it in a way that other people can understand what you are doing and act on it. In other words, you need to summarize, display, and visualize your results. Of course, it is often an **iterative process** where you go back and forth between analyzing, summarizing, and presenting your work.

It would be possible to have a whole course just on these topics, so we will necessarily have to be brief here. However, it is still possible to point you into the right direction, which is often the most important part of this journey. We break down this big topic into three different parts:

- Summarizing your data in Section 20.1
- Plotting your results in Section 20.2
- Creating web apps in Section 20.3

Importantly, in this section we are going to focus on the top packages only. If you want to get an **overview of related packages**, see Section 9 starting on page 47 for Python-related packages.

20.1 Summarizing Your Data

Once you have obtained your data from the textual analysis steps, you often want to summarize it. Another thing you often need to do is to **merge it with other data, e.g. stock market data or accounting/fundamental data**. For example, you are testing whether there are some users on Twitter whose textual output consistently predicts market returns in the next day or week, you need to add further data about stock or stock index returns and ultimately you need to bring all that data together.

In the Python world, the pandas package rules (for more details **see Appendix B starting on page 203**). Originally pandas was written by Wes McKinney while he was

working at the hedge fund AQR. He was trying to address some of the frustrations people had when using R (although my guess is that these people were not using data.table in R at that time, otherwise they might not have been so frustrated). In any case, **pandas was the big bang event for data science in Python** and it is fair to say that it was a total game-changer. Before pandas, people were mostly using Python in areas other than data science. Pandas changed all that, and it made great strides because a lot of infrastructure in Python (mainly for scientific computing) had already been built with the NumPy and SciPy packages. Pandas took a lot of inspiration from R. For example, R has data.frames while pandas has DataFrames. The pandas syntax is also relatively similar to working in R with data.frames. In fact, switching between pandas and R is relatively easy because many of the concepts are the same or similar.

If your data is larger than memory, but not yet huge as in big data, Python is in the sweet spot with dask. The dask package builds upon pandas and allows it to split up the data and process it in smaller chunk. It thus avoids running out of memory. Importantly, it achieves this largely transparently, so you don't have to worry about how it actually splits up the data. Furthermore, it tries to stay consistent with the pandas syntax, so if you know how to use pandas, you also know how to use dask automatically.

20.2 Plotting

This course is not mainly focused on plotting. However, sometimes it is necessary to create a graphical representation of your data. Below are pointers to some of the popular plotting packages in Python.

Python has various plotting packages available.

- 1. For an overview of the most popular visualization packages in Python, please take a look at Section 9 starting on page 47.
- 2. For plotting with the Pandas package (which wraps Matplotlib for plotting), see Section B.5 starting on page 213.
- 3. In the current section we provide a brief introduction to Matplotlib.

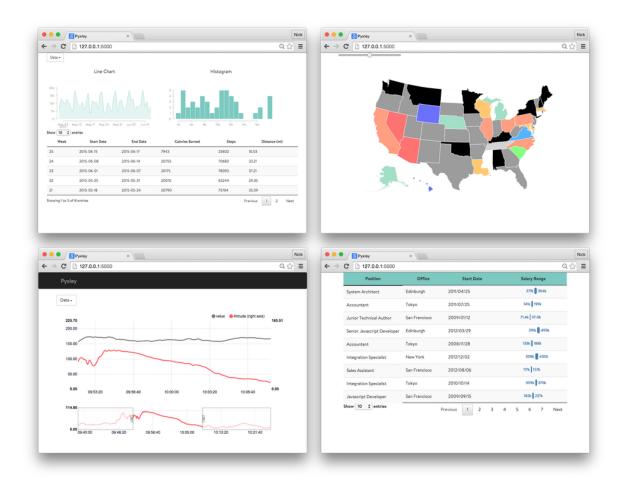
Matplotlib historically often has been a good place to start, although there are many other plotting packages available by now. For example, to plot a histogram using Matplotlib, you can simply type:

```
import matplotlib.pyplot as plt
import numpy as np
plt.hist(np.random.randn(1000))
plt.show()
```

20.3 Creating Web Apps

Often it makes sense to create a dashboard, which is a web-based analytics app, where you can showcase your work. In Python there are many options available, the most

popular ones being Dash and Pyxley, although there are many others as discussed in Section 9.9 starting on page 53. A dashboard can be very useful, e.g. if you have a client meeting and would like to demonstrate and visualize the functionality of your code. People can look at charts and diagrams and interact with them in real-time. A few example dashboards can be seen below:



Appendix A

Gentle Introduction to Python

Where does the name "Python" come from? Originally it nothing to do with the animal of the same name. Instead, it is named after the BBC show "Monty Python's Flying Circus," which is a British sketch comedy series from the late 1960s and early 1970s. The Python language was created by Guido van Rossum, a Dutch programmer who now lives and works in the U.S. He is the "benevolent dictator for life" (BDFL) of the Python programming language, a title given to him by the Python community.

Python is one of the most widely-used programming languages for text analytics and NLP. Of course you can do NLP in any programming language, but not all of them offer a large ecosystem of NLP-related libraries. Furthermore, for languages other than Python, the community might not always be very strong for NLP-related tasks, e.g. you might not be able to find a lot of answers on StackOverflow.

In this chapter will gently go through the basics of Python programming. If you would learn more about **text processing** in Python, you can visit Chapter 10 starting on page 55.

A.1 Python Philosophy

At the time when Python came out in 1991, another language called Perl was relatively popular. One of Perls programming mottoes was "there is more than one way to do it." This is fine, and Perl is still a great language, but one problem with this motto was that it sometimes confused newbies and made the language maybe less accessible to some people. Python on the other hand had a very different philosophy. One important Python motto is, "there should be one— and preferably only one—obvious way to do it." This really rang a bell with many people and Python has been and still is growing at an amazing speed, having overtaken Perl in popularity long ago.

You can read the Zen of Python directly here (you can also read it in Python by typing import this). There is a lot of truth and wisdom hidden in there that benefits most software developers, no matter which language they program in.

Beautiful is better than ugly. Explicit is better than implicit.

Simple is better than complex.

Complex is better than complicated. Flat is better than nested. Sparse is better than dense. Readability counts. Special cases aren't special enough to break the rules. Although practicality beats purity. Errors should never pass silently. Unless explicitly silenced. In the face of ambiguity, refuse the temptation to guess.

There should be one -- and preferably only one -- obvious way to do it.

Although that way may not be obvious at first unless you're Dutch.

Now is better than never.

Although never is often better than *right* now.

If the implementation is hard to explain, it's a bad idea.

If the implementation is easy to explain, it may be a good idea.

Namespaces are one honking great idea -- let's do more of those!

A.2 Why Are We Using Python?

Because it's a language that has a few key features that makes it very suitable to what we are doing in this course.

- 1. Easy to learn: Python is a high-level programming language.
 - It is dynamically typed. This means that you do not have to do a lot of bookkeeping such as in lower-level languages such as C, C++, or Java. For example, you can define new variables on the fly and you don't have to specify their type.
 - You also don't have to worry about memory management because a garbage collector is included.
 - It's interpreted. It means you don't have to compile a bunch of files like in C or C++, so you don't have to worry about linking files and loading them. Furthermore, you have a terminal where you try things out interactively. You type something and immediately you can see the result.

So Python tries to do a lot for your behind the scenes. Of course, this comes at a price. Python is slower than low-level languages. So the tradeoff is that you can write programs faster than in, say, C, but your programs tend to run slower. Still, for many tasks the speed of Python is just fine, so it's a great programming language to learn.

Furthermore, Python is a simple and minimalistic language. Reading a Python program almost feels like reading English. It allows you to concentrate on the solution rather than the language.

- 2. Powerful: Python is a full-fledged programming language originally designed by Guido van Rossum. Due to it being a high-level programming language, you can get things coded up very quickly.
- 3. Community: Even a great programming language cannot really thrive unless it's backed by a great community. (Of course, great programming languages often foster a great community, but it is not necessarily always the case.) A great community means that people share their ideas and experiences in blog posts, contribute to discussions in online forums, and share and answer questions on sites such as Stack-Overflow.com. This means that if you're looking for something, instead of studying a potentially obscure manual, you simply google whatever question is on your mind and you will most likely find an answer very quickly. This can be a tremendous productivity boost for your programming.
- 4. Documentation: This is the minimum requirement for any great programming language, but I view documentation in a larger context. It should not only encompass the language itself, but also the whole ecosystem of software packages around that language. Specifically, it should be relatively easy to create professional-looking documentation if you're writing your own software so as to encourage best practices and sharing of information. Even a great software package is not very useful if it is not well documented. And we all know that documentation is not necessarily the favorite hobby of most programmers. So it is good if there are state-of-the-art tools available such as Docutils and Sphinx that facilitate documentation and make it easier.
- 5. Libraries and ecosystem: Even if you have a well-designed programming language, if it doesn't come with "batteries included" (i.e. a set of useful add-on libraries), it might be difficult to get things done efficiently and quickly. In the worst case, you might have to re-implement what is already available in other programming languages. Consider for example Haskell. It is a beautiful language, but at the time of this writing it is behind Python or R in terms of its add-on libraries for data science. Even if you had the time to re-implement some of the Python or R packages in Haskell, it would likely be a painful, slow, and error-prone process. Take for example BLAS, which originated as a Fortran library in 1979. All it does is do common linear algebra operations such as vector addition or matrix multiplication. Sounds easy enough. But if you would implement it yourself, you would quickly run into many subtle problems. Among others, there are subtle numerical issues that a naive implementation almost certainly overlooks such as numerical stability when inverting a matrix or round-off errors when solving systems of linear equations. In many cases these issues might not show up, but when they do, they can lead to false conclusions and ultimately to wrong decisions, which, as we all know, can be very costly in finance.

Especially in the areas of text analytics, NLP, data science, and machine is where Python shines. It is a wealth of add-on libraries that are state-of-the-art and have

an active developer and user community. Besides Python being easy to use, this is the main reason why we are using Python in this course.

A.3 Software Development

When you develop your software, it is typically a good idea to follow these steps:

- 1. Analyze the problem you're dealing with.
- 2. Design a potential solution or approach to deal with your problem.
- 3. Implement it using software, i.e. start writing code. Begin with a simple version that gets the core job done.
- 4. Test and debug the previous step.
- 5. Deploy what your software to make sure it works as expected.
- 6. Maintain and refine your software.
- 7. Add new features (start again with the implementation part and repeat the steps thereafter as often as necessary)

A.4 A First Tour of Python

You can start Python by typing python3 in your terminal and you can quit by pressing [ctrl+d]. After starting Python, you can then print something, e.g.

```
print('I love NLP.')
```

You can also save this code into a file, e.g. testing.py (use any IDE or editor you like mentioned in Sections 4.7 and 4.8). Then run it from the Unix command line (make sure that you're in the same directory, otherwise change into it by using cd mydirectory):

```
$ python3 testing.py
```

Now let's go back into the Python shell. If you have text that should not be run ("interpreted") by Python, you can add it as a **comment** by using the # symbol.

```
print('I love NLP.') # This is a comment.
# This is another comment.
print('I love text analytics.')
```

Why should you be using comments? Because they let you explain the **why**. Comments can help you explain what you're actually trying to accomplish in this part of the code, the assumptions you're making, and further details that might not be obvious from just reading your code. They are not only useful for your teammates to understand you, but also for you to understand yourself two weeks from now.

If you would like to learn more about a given function or statement, you can simply type the following command to **display the help page** and you can type q to exit the help.

```
help(print) # Exit by typing 'q'.
```

A **string** is a sequence of characters. As you can imagine, in this course, strings are particularly important, since they are the fundamental building blocks for text analytics and NLP. In Python, it doesn't matter whether you use single or double quotes.

```
'Hello world.' # Both give the same result.
```

Sometimes you want triple quotes in case you have long strings spanning several lines (so-called multi-line strings). You can use ' or " to create triple quotes (in the following example we use ').

```
'''I love NLP.
NLP is great.
'''
```

Multi-line strings can be useful, e.g. if you want to document the functions you are writing (using a so-called **docstring**).

```
def f(x):
    '''Simple function that just squares
    its input and returns it.
    '''
    return x**2
```

In Python there are many ways to construct strings from other information.

```
>>> x = 'NLP'
>>> y = 'text analytics'
>>> 'I love {}'.format(x)
'I love NLP'
>>> 'I love {} more than {}'.format(x, y)
'I love NLP more than text analytics'
>>> 'I love {0} more than {1}'.format(x, y)  # Same as before.
'I love NLP more than text analytics'
>>> 'I love {1} more than {0}'.format(x, y)  # Reversed!
'I love text analytics more than NLP'
```

The good thing about the format() method is that it doesn't really care what data type you use. You could also input a number and format() would convert it to a string.

```
>>> x = 8
>>> 'I love {}'.format(x)
'I love 8'
```

Alternatively you can also concatenate strings with the + operator, but if you input a number, you need to convert it first to avoid an error.

```
>>> x = 1
>>> 'One plus ' + x + ' is two.'  # Doesn't work.
Traceback (most recent call last):
   File "<stdin>", line 1, in <module>
TypeError: must be str, not int
>>> 'One plus ' + str(x) + ' is two.' # This does work!
'One plus 1 is two.'
```

In some strings you might have seen backslashes such as \', \\, \n or \t. The purpose of the backslash is to "protect" whatever comes after it.

```
'That's great' # Won't work as Python things string ends after "that".
```

To solve this, you have two options, one of them is protecting the single quote with a backslash, the other one is to alternate between single and double quotes.

```
'That's great'
"That's great"
```

For example, if you want a backslash in your string, you need to protect it as well. Guess with what? Yes, with another backslash!

```
>>> print('I want to write a \\telegram.')
I want to write a \telegram.
```

There are other ways the backslash is used (in general, these are all called **escape sequences**). For example, a newline is \n and a tab is \t. Further escape sequences are also used in so-called regular expressions.

```
>>> print('This is a\ntest of the newline!')
This is a
test of the newline!
>>> print('Here we add \t a tab!')
Here we add a tab!
```

If you want to disable this special preprocessing of escape sequences, you can use a raw string by adding r in front of the string. **It is usually a good idea to use raw strings when dealing with regular expressions** as you don't have to use so many backslashes to protect things.

```
>>> print(r'Here we add \t a tab!')
Here we add \t a tab!
>>> print('Here we add \\t a tab!') # Same effect, but need extra '\'.
Here we add \t a tab!
```

A.5 Basics

- Difference between a Python module and a Python package: A **module** is a Python source file, while a **package** is a directory of Python modules. For further details see Section A.11 starting on page 179.
- Manage the working directory. The working directory is the location on your hard drive where Python reads or writes files by default.

• If you have a command that goes beyond one line, you can break it up using a backslash ("\"). Make sure to add **no space after the backslash.** The following is a stylized example where we break up the assignment into two lines of code.

```
x = /
```

• Multiple assignment

```
x = y = z = 8 # x, y, z will all be 8.
x, y, z = 'hello', 8, 3.14 # Same as assigning separately.
```

• pprint is useful for pretty-printing of data structures. The following example is based on the official Python documentation:

VALID:
hello_world
helloWorld

A.6 Elementary Python

A.6.1 Identifiers and Variables

Identifiers can be variable names, function names, class names, and so on. You can basically use any name you like as long as it doesn't start with a number or special symbol such as > and doesn't have spaces or dashes in it.

```
hello_world8
# INVALID:
8hello
>hello
with spaces in between
hello-world
Assigning things to variables is very straightforward in Python.
>>> x = 8
                 # Add three.
>>> x = x + 3
>>> print(x)
11
                 # Shorter: add one.
>>> x += 1
>>> print(x)
12
You can delete identifiers using del:
>>> x = 10
>>> x
10
>>> del x
>>> x
                     # It's gone.
Traceback (most recent call last):
  File "<stdin>", line 1, in <module>
NameError: name 'x' is not defined
```

Sometimes you have a **very long line of code and you want to break it into several lines for better readability**. Backslash to the rescue! Whenever you're not finished typing your command but want a newline to make your code more readable, just add a backslash at the end. But be careful, **after the backslash should not be any whitespace**, **otherwise it won't work**.

```
x = \
    'This is a really long' + \
    'string and I want to break' + \
    'out the code across' + \
    'several lines.'
```

A.6.2 Whitespace (Indentation) Has Special Meaning in Python

Speaking of whitespace, in Python code you might notice that there are very few curly brackets like in other languages. Here are two examples of a function definition, one in R, the other in Python.

```
# Function definition in R.
f = function(x) {
    y = x^2
    return(y)
}
# Function definition in Python.
def f(x):
    y = x**2
    return y
```

As you can see, what really matters in Python is whitespace, or in this specific case, indentation. **In Python, indentation has a special meaning** and the best practice is to use four spaces for indentation. For example, in R you have curly braces around the function definition so whitespace is irrelevant and the following code would work in R. In Python however the following code would fail to work because whitespace is part of the syntax.

```
# Would work in R (although it's ugly).
f = function(x) {
y = x^2
return(y)
}
# Would NOT work in Python as whitespace has meaning.
def f(x):
y = x**2
return y
```

A.6.3 Arithmetic, Comparison, and Logic with Operators

Operators generally behave like functions, but they usually have a special syntax. For example, you could write a function called plus and calculate plus (5, 2) which would yield 7, but I guess we can all agree that writing 5 + 2 is much easier and more direct.

The most important **operators** in Python are as follows:

```
+, -, *, / : basic arithmetics
**, //, % : power, divide and floor, modulo
<, <=, >, >=, ==, != : less/greater, equal to, not equal to
not, and, or : boolean operators
```

For example:

```
>>> 8 ** 2
              #8 to the power of 2.
64
>>> 7 // 2
               # Divide and floor.
>>> 7 % 2
              # Modulo.
>>> 8 < 10
True
>>> 8 == 8
True
>>> 8 != 10
True
>>> True and False
False
>>> True or False
True
>>> not True
False
```

Sometimes you want to assign a calculation back to the variable. In this case there are some **shortcuts for using operators**. The basic idea is simply that you add the operator you want to apply in front of the equality sign.

```
>>> x = 8

>>> x += 2

>>> print(x)

10

>>> x *= 3

>>> print(x)

30
```

A.6.4 if, for, while

if, for, and while are three key control flow statements in Python. They allow you to tell Python to do different things depending on different situations.

```
my_preference = 'yes to NLP'
if my_preference == 'yes to NLP':
    print('I love NLP')
else:
    print('I love text analytics')
```

This the basic if statement, but you can also make it more versatile by adding a few other conditions with one or several elif. But keep in mind that else and elif are optional.

Another way, if you only want to return some simple values in the if statement, you can also use a dictionary.

```
dictionary = {'Larry': 'larry@wall.org', 'Spammer': 'spammer@hotmail.com'}
g = input('Larry or Spammer?')
dictionary[g]  # Error if not in 'dictionary'.
dictionary.get(g)  # Works always (returns 'None' otherwise).
```

The next thing we're going to cover are loops. There are two kinds of loops, and the first one is the while loop. The following loop prints x and decreases x by one each time it runs. At the beginning, it always checks whether x is still at least one (or greater) and if this condition is no longer fulfilled, it stops. So in total, the following loop just prints the numbers eight to one in a decreasing sequence.

```
x = 8
while x >= 1:
    print(x)
    x -= 1
```

Here's another example, where we check a condition each time inside the loop and set a variable ct that determines whether to continue the while loop. The condition is random. We draw a random number between 1 and 100, and if that number is less than 5 we let the loop stop. So with a 5% probability we will stop and we will continue with a 95% probability each time the loop is run. To better monitor how often the loop runs, we print "NLP" each time it runs.

```
import random
ct = True  # Variable that will determine whether to continue.
while ct:
    print('NLP')
    if random.randint(1, 100) < 5:
        ct = False</pre>
```

The next kind of loop is a for loop. It just iterates over a sequence of objects. In the following example, it just prints the numbers one to eight.

```
for i in range(1, 8):
    print(i)
```

In fact, range() returns a so-called iterator, whose main purpose in this case is to return one number at a time. If you want to get all numbers at the same time, you can use list(range(8)) for example.

If you want more control over loops, you can use the break statement. The following example will only print the numbers one to five, because the loop will stop if i is larger than five.

```
for i in range(1, 8):
    if i > 5:
        break
    print(i)
```

Here's an example of how break works in the while loop. The following loop is going to print out the numbers one to eight, because it breaks for any larger number.

```
x = 0
while True:
    x += 1
    if x > 8:
        break
    print(x)
```

Sometimes you want to skip the rest of the statements in the current loop block. continue lets you do exactly that. As you can see, the first print statement is executed each time, but the second one only the first three times. The reason is that after the first three times, continue is run, so the rest of the for loop is skipped.

```
>>> for i in range(1, 8):
... print('First print statement, i =', i)
... if i > 3:
... continue
... print('Second print statement, i =', i)
...
First print statement, i = 1
Second print statement, i = 1
First print statement, i = 2
Second print statement, i = 2
First print statement, i = 3
Second print statement, i = 3
First print statement, i = 3
First print statement, i = 4
First print statement, i = 5
```

```
First print statement, i = 6
First print statement, i = 7
```

You can use continue also in a while loop. The following code will have the same output as the previous for loop.

```
x = 0
while x < 7:
    x += 1
    print('First print statement, x =', x)
    if x > 3:
        continue
    print('Second print statement, x =', x)
```

A.6.5 Defining and Using Functions

You write functions because you have code that you want to reuse, maybe with different parameters. A simple function that takes its function argument x, squares it, and returns the squared value is given here:

```
def f(x):
    return x ** 2
```

You can check that it performs as expected:

```
>>> f(3)
9
>>> f(8)
64
```

If course, you can also do other things in functions such as print or do other calculations.

```
def f(x, y):
    print('Multiplying both arguments with each other yields', x * y)
    return x * y
```

Again it works as expected. In the first function call, the function prints the message and then returns the output value of 24, which is directly printed by Python. In the second call, we save the output value to a variable x and then take a look at that variable (which, as we find out, contains again 24).

```
>>> f(3, 8) Multiplying both arguments with each other yields 24 24 >>> x = f(3, 8)
```

```
Multiplying both arguments with each other yields 24 >>> \mathbf{x} 24
```

You can also omit the return statement, in which the function by default will return None.

```
def f(x, y):

print('Multiplying both arguments with each other yields', x * y)
```

You can check its output as follows:

```
>>> x = f(3, 8) Multiplying both arguments with each other yields 24 >>> x == None True
```

Unless you define a variable inside a function as a global variable (which is usually a bad idea, so we are not going to cover this), it will be treated as a **local variable**. It simply means that whatever happens inside the function stays inside the function and has no effect on anything outside the function. The following code illustrates what that means concretely.

```
def f(x):
    print('Local variable x =', x)
    x += 100
    print('Local variable x changed to', x)
```

If we run it as follows, notice that the variable x outside the function is not changed at all if we run the function. The reason is that **the variable** x **inside the function is a completely separate entity than the variable** x **outside the function**.

Sometimes you have functions that have many function arguments, for example:

```
def f(a, b, c, d, e, f):
    return a * b * c * d * e * f
```

Suppose that for some reason, most of these arguments are usually equal to one. In that case you can just specify the **default argument values**, which means you don't have to supply them each time.

```
def f(a = 1, b = 1, c = 1, d = 1, e = 1, f = 1):
return a * b * c * d * e * f
```

You can now run it, and if you don't specify an argument, it will be equal to one by default.

```
>>> f()
1
>>> f(3)
3
>>> f(3, 8)
24
```

Another thing that is useful when defining or using functions is to use **keyword** arguments.

```
def f(x = 10, y = 20, z = 30):
print('x =', x, ', y =', y, ', z = ', z)
```

Now you can use it as follows:

```
>>> f(1, 2, 3)  # Normal usage.

x = 1, y = 2, z = 3

>>> f(x = 1, y = 2, z = 3) # Specify keyword arguments.

x = 1, y = 2, z = 3

>>> f(z = 3, y = 2, x = 1) # Order doesn't matter any more!

x = 1, y = 2, z = 3

>>> f(z = 3, x = 1) # If omit one, use default values.

x = 1, y = 20, z = 3

>>> f(z = 3) # If omit several, use several defaults.

x = 10, y = 20, z = 3
```

As we have seen Section A.4 you can (and should) use a docstring for most of your function. It's simply a bit of documentation that lives right in the function definition and gives the user some idea of what the function does and what its purpose is.

```
def f(x):
    '''This is the docstring of this
    function. It describes what it does
    and why it does it.
    '''
    return x ** 2
```

You can now take a look at this docstring by typing help(f) or by typing print(f.__doc__).

A.7 Object Oriented Programming

Python is a multi-paradigm programming language. You can do object-oriented, imperative, functional, and procedural programming with Python. However, if one would have to put Python into one category only, it would probably be the object-oriented corner. We have already seen how you can assign values to variables and how to define functions. Object-oriented programming in Python takes this one step further and puts more structure on this.

A.7.1 Class Example 1

Let's look at a very simple introductory example. Here we define the simplest class we can think of, a class that does nothing. The pass statement in the class definition below does nothing. You can use it when a statement is syntactically required but you don't want any action to be performed. In this case, we just use it as a placeholder so that we can at least define the class c.

```
class c:
    pass # Do nothing
```

Having defined a class is a good start, but we need to instantiate it next. The basic idea is that a class is a blueprint for something, but we first need to **instantiate** it before we can do anything with it. We instantiate two objects of that class and then print them. Both objects conform to the blueprint given by the class (i.e. do nothing), and you can see from the print statement that they reside in different addresses in computer memory, so they are really two distinct objects. The reason why we have instantiated two objects is simply to show that the class is just a blueprint, and you can create many objects from that blueprint.

```
>>> o1 = c()
>>> o2 = c()
>>> print(o1)
<__main__.c object at 0x7fcad25a4438>
>>> print(o2)
<__main__.c object at 0x7fcad25a44e0>
```

A.7.2 Class Example 2

Let's create a bit of a more sensible class that actually does something.

Here we create only one instance of this class for illustration, although we could, as before, in principle create as many as we wish.

```
>>> o = c()  # Create object 'o', which is an instance of class 'c'.
>>> o.hello() # Call method 'hello()' of object 'o'.
Hello world.
```

You can see that you can call the **method** hello() of the class by calling it after the object, separated by a dot. We will explain later why self has to be specified as an argument to the hello() method. For now, all you need to know is that **any method** (i.e. a function defined as part of a class) has to have as its first argument self, even though it doesn't always seem to be used.

A.7.3 Class Example 3

A classic introductory example is if you are dealing with people. You want to represent a person that can say "hi" and that has an age. **A class is blueprint of an object**, so for example, to create a person object, you first need to provide the blueprint.

```
class person:
    def __init__(self, name, age):
        self.name = name
        self.age = age

    def say_hello(self):
        print('Hi, my name is', self.name)

    def say_age(self):
        print('I am {} years old'.format(self.age))
```

Remember that indentation is important in Python. When you are inputting this class in a text editor or directly into Python, make sure to add four spaces in the empty lines between the function (in fact, "method") definitions.

We will go through this class definition (the blueprint) in more detail shortly. But before we do that, let's quickly see how to use it. In what follows we will **instantiate** two objects created from this class. These are two different persons with their respective names and ages.

```
g = person('Guido', 25)
l = person('Larry', 30)
```

We can now let these persons tell us their names and ages using the say_hello() and say_age() **methods**.

```
>>> g.say_hello()
Hi, my name is Guido
```

```
>>> g.say_age()
I am 25 years old
>>> l.say_hello()
Hi, my name is Larry
>>> l.say_age()
I am 30 years old
```

You could also access the name and age variables directly. These variables that are part of a such an object (which is an instantiated class) are called **fields**.

```
>>> g.name
'Guido'
>>> g.age
25
>>> l.name
'Larry'
>>> l.age
30
```

In general, fields and methods are collectively called **attributes** of a class. Now let's go step by step through this class definition and explain every detail.

- 1. There are three function definitions in the class object. As they are part of a class, each of these functions is called a "**method**." The reason why we use a special name und why we don't call them "function" any more is because they are closely tied to the class we just have defined and only work on objects of this class.
- 2. What is "self"? It refers to the object that will be created using this class (remember, the class is just the blueprint). So why is it included in all the method definitions when you don't usually use it inside the method definitions (with the exception of __init__() which we'll talk about shortly)? There is a technical reason for this. Technically speaking, suppose you have created an instance g of the class person. Whenever you call g.method(x1, x2), what happens is that Python behind the scenes actually calls person.method(g, x1, x2). That's why the first argument of each method is always self. So in this call, self gets set to g, which is the instance of the class.
- 3. Putting aside <code>__init__()</code> for a moment, you can see that all the other methods (i.e. <code>say_hello()</code> and <code>say_age()</code>) are basically just function calls that are executed on the specific instance of the object. In fact, what happens behind the scenes is that instead of <code>g.say_hello()</code>, Python actually executes

```
person.say_hello(g)
```

4. The __init__() method is also sometimes called the **constructor method**. It is called when an object of that class is instantiated. In this example, the __init__() method sets the field's name and age of the instantiated object.

A.7.4 Class Variables and Object Variables

What we have seen so far are object variables, e.g. name and age. They are specific to the object instantiated, and they can have different values for each object. In contrast, sometimes you might want to have **class variables** that are the same for all objects that are instantiated from a given class.

Now let's create to instances of this class and let's see what happens to the class and object variables. The point of the following code is to show that the class variable is the same for all objects of this class.

```
>>> x = c(8)  # Create the first instance of class 'c'.
>>> x.object_var  # Check the object variable.
8
>>> x.class_var  # Check the class variable.
1
>>> y = c(88)  # Let's create another instance.
>>> y.object_var  # As expected.
88
>>> y.class_var  # Class variable has been increased.
2
>>> x.object_var  # Still the same.
8
>>> x.class_var  # Has increased as well, even though it's a different object.
2
```

If you would like to change the class variable, there's a small caveat to keep in mind. You cannot just "overwrite" the class variable of an object. The reason is that if you use an object variable that has the same name as class variable, it will hide the class variable. For example:

```
>>> x.class_var = 300
>>> x.class_var # What we are seeing now is the object variable of the same name.
300
>>> y.class_var # 'y' still displays the class variable.
2
```

If you want to avoid this, you can add a **class method** that takes care of dealing with the class variable.

Now we can use this class method whenever we want to update the class variable.

```
>>> a = c(8)
>>> b = c(88)
>>> a.class_var
2
>>> b.class_var
2
>>> c.update_class_var(200)
>>> a.class_var
200
>>> b.class_var
200
```

A.8 Inheritance

In finance you can think of a stock and a European call option on a stock as financial instruments. Both are traded on the market and thus both have a price. What is unique about the stock is that we should also keep track of its volatility. And what is unique about the European call is that it has a strike price. (We ignore for now that the call also depends on a few other parameter, including the volatility of the stock.) So let's capture this hierarchy using different classes that build on each other.

Specifically, we first define a class financial_instrument that has a price field. Then we define a stock class and a option class. Both of them build on financial_instrument and can add further fields (as in the example below) or in principle further methods as well. The reason why this can be a good idea is that we don't have to reinvent the wheel twice, i.e. we don't have to deal with price again, as financial_instrument already takes care of it. In this example, financial_instrument is called the **base class** or **superclass**, while stock and option are called the **derived classes** or **subclasses**.

```
class financial_instrument:  # Define base class.
   def __init__(self, price):
        self.price = price
```

```
class stock(financial_instrument): # Define first derived class.
    def __init__(self, price, vola):
        financial_instrument.__init__(self, price)
        self.vola = vola

class option(financial_instrument): # Define second derived class.
    def __init__(self, price, strike):
        financial_instrument.__init__(self, price)
        self.strike = strike
```

Now we can instantiate a stock and an option.

```
>>> s = stock(100, 0.20)
>>> o = option(5, 80)
>>> s.price
100
>>> s.vola
0.2
>>> o.price
5
>>> o.strike
80
```

A.9 Data Structures

Python has the following basic data types:

- Boolean: Takes values True or False.
- Numeric types: int for integers, float for floating point numbers, and complex for complex numbers.
- Stings: Immutable sequences of Unicode code points, e.g. 'hello world'. For more details see Chapter 10 starting on page 55. In fact, **a string is an iterable** (see also Section A.14.3 starting on page 189) because it is a sequence, see

```
from collections.abc import Iterable
isinstance('', Iterable) # True.
```

Python has four built-in data structures called **containers**. Containers are objects that hold an arbitrary number of other objects. We will discuss each one of them in turn. Their main purpose is to hold some data.

1. list

- 2. tuple
- 3. dictionary
- 4. set

A.9.1 List

A **list** is a sequence type, e.g. ['item_1', 'item_2', 'item_3']. While a list can be a heterogeneous collection (i.e. containing mixed data types, e.g. ['hello', 8]), it is most frequently used for **homogeneous** collections containing the same data types (e.g. ['hello', 'world']). Lists can be **nested**, i.e. they have another list as an item, e.g. ['hello', [1, 2, 3]]. Lists are **mutable**, so you can change/update them:

```
>>> 1 = [2, 4, 6, 8]
>>> 1[0] = 10
>>> 1
[10, 4, 6, 8]
```

In this course, the list is probably the most important data structure overall. They contain a sequence of objects. For example, we can use it to hold a whole corpus, i.e. a sequence of text documents. We can also use a list to hold the words in a single text document. It is a very powerful and versatile data structure.

```
>>> 1 = ['I', 'love', 'NLP']
>>> len(1)
                      # Length of list.
3
>>> for item in 1:
                      # Loop over the items of the list
        print(item)
. . .
Ι
love
NLP
>>> l.append('and text analytics') # Add new element to the end.
>>> 1.sort()
                                     # Sort the list.
['I', 'NLP', 'and text analytics', 'love']
>>> del 1[0]
                                     # Delete first element.
>>> 1
['NLP', 'and text analytics', 'love']
```

To manipulate lists, people often use a **list comprehension**. This is a shortcut for looping over the items of the list and doing something with them. For example, the following list comprehension creates a new list where _hello is added to each item, unless the item is 'NLP'.

```
>>> 1 = ['I', 'love', 'NLP']
>>> [item + '_hello' for item in 1 if item != 'NLP']
['I_hello', 'love_hello']
```

Another example illustrating a list comprehension:

```
>>> [(x, y) for x in [1, 2, 3] for y in [3, 1, 4] if x != y] [(1, 3), (1, 4), (2, 3), (2, 1), (2, 4), (3, 1), (3, 4)]
```

zip can be used to iterate through any two iterables, e.g. through two lists.

```
>>> alist = ['a1', 'a2', 'a3']
>>> blist = ['b1', 'b2', 'b3']
>>> for a, b in zip(alist, blist):
... print(a, b)
...
a1 b1
a2 b2
a3 b3
```

The following table shows useful list methods:

Method	Description
append()	Add element to end of list
extend()	Add list to another list (can also done with +)
<pre>insert()</pre>	Insert item at the defined index
remove()	Remove item from the list
pop()	Removes and returns an element at the given index
clear()	Removes all items from list
index()	Returns index of first matched item
count()	Counts how many times an element has occurred in a list
sort()	Sort list in ascending order
reverse()	Reverse order of items
copy()	Returns shallow copy of list

The followin table shows built-in functions that can be used for lists:

Function	Description
all()	Returns True is all elements are True or if list is empty.
any()	Returns True if at least one element is True. any([]) is False.
enumerate()	Adds a counter, so you can for example loop through the list and have an automatic counter.
len()	Length of the list (the number of items).
list()	Convert an iterable (tuple, string, set, dict) to a list.
max()	Largest item in the list.
min()	Smallest item in the list.
sorted()	Returns a new sorted list (does NOT sort the list itself), e.g. sorted(1, key=lambda x: x[1], reverse=True).
sum()	Sums all elements in the list.

A.9.2 Lists and References

Suppose you have a list saved in a variable and assign it to a new variable. What happens in Python is actually that **the assignment does not copy the list, but it copies the reference to the list**. A reference is just pointing to the memory location. So in the code below you have both x and y pointing to the same location in memory. If you change x, you also change y.

This is fine if that's what you're trying to accomplish, e.g. in order to use computer memory more efficiently. However, if you want to **copy the list**, you can use one of the following ways to create a **shallow copy**:

```
x = ['hello', 'world']
a = x.copy()  # Use 'copy()' method.
b = x[:]  # Slicing
c = list(x)  # Use built-in 'list()' function.
```

A.9.3 Tuple

Tuples on first sight look like lists, but they are usually used in a different way. While a **list** is usually used for objects that **appear in a sequence**, e.g. different readings in time from a sensor, with a **tuple** it's more about the **position of the elements**, e.g. in

the form of coordinates such as (longitude, latitude). If you switch two elements in a list, it usually does not make a big difference. But if you switch two elements in a tuple, it might matter a lot (depending on what kind of data it contains).

A key difference in how you use tuples vs. lists is that a **tuple has structure while a list has order**. For example, you could store information about a customer, e.g. name, gender, and age e.g. ('Bob', 'male', 32). Or you would use a tuple to store coordinates like (latitude, longitude) or locations in a book such as (page_number, line_number). On the other hand, you would use a list if you take a trip and record the city names you visit e.g. ['Hong Kong', 'London', 'New York'].

A tuple is a sequence type (like a list), but unlike a list it is **immutable** and thus can be used as a key in a dict (see Section A.9.4 starting on page 176). The immutability of a tuple makes sense, e.g. you would not want to swap longitude and latitude. However, for a list, mutability makes sense, e.g. if you change your travel plans visit the cities in a different order.

Here is some example usage of tuples, with comments explaining the code inline. Keep in mind that **indexing in Python starts at zero!** Also keep in mind that you can use a tuple recursively, i.e. it can contain elements of different types, e.g. it can have another tuple as an element.

```
>>> fi = ('stock', 'call option', 'put option') # Financial instruments.
>>> len(fi)
3
>>> new_fi = ('asian option', fi) # Recursive usage.
>>> new_fi
('asian option', ('stock', 'call option', 'put option'))
>>> new_fi[1]
('stock', 'call option', 'put option')
>>> new_fi[1][2]
'put option'
```

There is no tuple comprehension like a list comprehension, but if you really need it you can use a **generator expression** (see Section A.14.5) and convert it into a tuple again.

```
>>> tuple(i + 3 for i in (1, 2, 3, 4, 5) if i \ge 2) (5, 6, 7, 8)
```

A word of caution: Tuples are immutable in the sense that the membership of a tuple cannot be changed, but **mutable elements** (e.g. a list) of a tuple *can* be changed:

If you want to avoid this kind of behavior, you can use deepcopy on the tuple:

```
from copy import deepcopy
l = list()
t1 = (1, 1)
t2 = deepcopy(t1)
l.append(1)
t1  # Has been changed.
t2  # Unchanged, yay!
```

A.9.4 Dictionary

A dictionary is like a phone book. It has a bunch of **key-value pairs**, which are like name-number pairs in a telephone book. Here is a similar example, but with email addresses. They key thing is that a dictionary uses curly braces, the keys and values are separated by a colon, and key-value pairs are separated by commas.

```
d = \setminus
    {'Jeff': 'jeff@amazon.com',
     'Bill': 'bill@microsoft.com',
     'Larry': 'larry.page@google.com'}
Let's see what we can do with a dictionary.
>>> len(d)
                      # Number of key-value pairs.
>>> d['Jeff']
                     # Jeff's email address.
'jeff@amazon.com'
>>> del d['Bill']
                      # Delete Bill's entry.
>>> for key, value in d.items():
                                    # Use 'items' method.
        print('{} has email {}.'.format(key, value))
. . .
Jeff has email jeff@amazon.com.
Larry has email larry.page@google.com.
>>> d['Guido'] = 'guido@python.org'
                                       # Add Guido.
>>> if 'Larry' in d:
                                       # Check if it has a given key.
        print('yay')
. . .
yay
You can have dict comprehensions similar to list comprehensions.
>>> {number: number * 2 for number in [1, 2, 3]}
{1: 2, 2: 4, 3: 6}
>>> d
{'Jeff': 'jeff@amazon.com',
 'Larry': 'larry.page@google.com',
 'Guido': 'guido@python.org'}
>>> {key: value for (key, value) in d.items() if key != 'Jeff'}
{'Larry': 'larry.page@google.com', 'Guido': 'guido@python.org'}
```

If you want to **sort a dict**, you need to use the items() method as well. Keep in mind that a dict is not sorted (which means **you cannot generate a sorted dict**, unless you use OrderedDict), so the output is a sorted list.

A.9.5 Set

Sets in Python are modeled after **sets in mathematics**. You can uses sets to test for membership. Every element can **occur only once** in a set. Also the **order of the set elements does not matter**. (If the order matters, you could use a list instead.)

There are also **set comprehensions**.

```
>>> s = {1, 3, 5}
>>> {x ** 2 for x in s}
{1, 9, 25}
```

If you want to get the intersection between a set and list:

```
myset.intersection(mylist)
```

Difference between list and set: lists are very nice to sort and have order while sets are nice to use when you don't want duplicates and don't care about order.

A.9.6 Collections

The **collections** module provides **specialized container data types** supplying alternatives to dict, list, set, tuple. Some useful ones for our purposes:

• **Counter** can be used for counting. (See also Section 15.4 starting on page 114 for an example how to use Counter for bag-of-words.) The resulting Counter object has a similar structure as a dictionary (it is a dict subclass).

Keep in mind that most_common does not tell us whether there are more words with the same frequency.

• **defaultdict** is a subclass of dict, so it behaves basically like a dict. defaultdict(int) can be useful for counting. The basic idea is that it is very similar to a dict, but the difference is that it is initialized with a "default factory" (in this case the int function) for a previously nonexistent key. The int function simply returns zero. In other words, if a new key is added to the dict, it will be initialized with a value of zero. Alternatively we could as well write defaultdict(lambda: 0) which would have the same effect (it uses an anonymous function that always returns zero).

```
from collections import defaultdict
l = 'spam spam spam spam spam eggs spam'.split() # List of food.
d = defaultdict(int) # Default value of int is zero.
for food in 1:
    d[food] += 1 # Increment value by one.
```

Another more elaborate example for counting with defaultdict(int):

```
from collections import defaultdict
l = [
    ("Lucy", 1),
    ("Bob", 5),
    ("Jim", 40),
    ("Susan", 6),
    ("Lucy", 2),
    ("Bob", 30),
    ("Harold", 6)
]
d = defaultdict(int)
for name, count in 1:
    d[name] += count
```

• namedtuple is a tuple subclass that adds names to each field. For example, instead of the tuple ('Bob', 'male') you would have (name = 'Bob', gender = 'male') with a named tuple.

A.10 Slicing

Slicing refers to extracting subsets of python objects. (See also the examples on slicing and stride in Section 10.1 starting on page 55.) It is important to remember that **indexing in Python starts with zero**. So 1 [0] extracts the first item.

```
>>> 1 = ['hello', 'world', 'slicing', 'is', 'great']
>>> 1[0]  # First item.
'hello'
>>> 1[2]  # Third item.
'slicing'
>>> 1[-1]  # Last item.
'great'
>>> 1[-2]  # Second to last item.
'is'
```

You can also perform slicing or indexing with strings in Python.

```
>>> s = 'hello world'
>>> s[0]
                # First character.
'n,
>>> s[2:5]
                # Characters three to five.
'11o'
>>> s[2:]
                # Starting at third character.
'llo world'
>>> s[1:-1]
                # Everything except for first and last.
'ello worl'
>>> s[:]
                 # All of it (no change).
'hello world'
```

A.11 Modules and Packages

We have seen in Section A.6.5 starting on page 163 how to make code reusable by wrapping it in functions. Once you start writing more code, you're likely going to have more an more self-written functions, and it is going to get increasingly difficult to manage them.

This is where the concept of a **module** comes in. You can easily create a module by putting your function (or class or variable) names into a text file whose name ends with .py. Use any IDE or editor you like mentioned in Sections 4.7 and 4.8. Suppose for example you create a file named my_first_module.py that has the following content:

```
def sq(x):
    ''', Function that squares
    its input.
    ,,,
    return x ** 2
_version__ = '1.0'
You can then import and it as follows:
>>> import my_first_module
>>> my_first_module.sq(8)
>>> my_first_module.__version__
>>> help(my_first_module.sq)  # Display help, exit with 'q'.
You can specify a shortcut for the imported module. This can save a lot of typing.
>>> import my_first_module as mfm
>>> mfm.sq(8)
64
>>> mfm.__version__
1.0
```

If you don't always want to type my_first_module.py, you can just import everything except names that start with __ by using *. However, this practice is usually discouraged because there could be name clashes. For example, if there is another module that also has a function called sq, it will not be clear which function (from which module) is called when you type sq(8).

```
>>> from my_first_module import *
>>> sq(8)  # Works fine.
64
>>> __version__  # Not imported.
Traceback (most recent call last):
  File "<stdin>", line 1, in <module>
NameError: name '__version__' is not defined
```

Alternatively, you can also import a few select functions, classes, or variables by explicitly specifying them, separated by commas:

```
>>> from my_first_module import sq, __version__
>>> sq(8)
64
>>> __version__
'1.0'
```

Depending on whether you import your module or run the module file standalone by itself, you sometimes might want your code to behave differently. Consider for example the file investigate_module_name.py with the following content.

```
print('__name__ has value', __name__)
```

If you import the module, it will have the following output:

```
>>> import investigate_module_name
__name__ has value investigate_module_name
```

On the other hand, if you run the following code on the Unix shell (i.e. you're running the module file standalone, all by itself, and don't import it), you will get a different output:

```
$ python3 investigate_module_name.py
__name__ has value __main__
```

So the thing to notice here is that __name__ has a different value depending on how you run your module. You can use this to make your module behave differently depending on how it is run.

For illustration, in the following example, we import our module the usual way, but if it is run in standalone, it will call one of its functions with a specific value. Suppose you have the following content in file mymodule.py.

```
def sq(x):
    return x ** 2

if __name__ == '__main__':
    print('Eight squared is equal to', sq(8))
```

You now can use this module the usual way from Python when you import it. Note that it does not print our sq(8).

```
>>> import mymodule as mm
>>> mm.sq(12)
144
```

And when you run it standalone, it will simply print sq(8).

```
$ python3 mymodule.py
Eight squared is equal to 64
```

If you want to see what's inside a module, you can use the built-in dir() function. There are other things in there as well (such as __name__), but you can also see your self-written sq function in there. In general, dir() shows all functions, classes, and variables in the module.

```
>>> import mymodule as mm
>>> dir(mm)
['__builtins__', '__cached__', '__doc__', '__file__',
'__loader__', '__name__', '__package__', '__spec__', 'sq']
```

You can also use dir() without an argument to see the names in the current scope. In the following code, we look at all names in the current scope, then we add a variable x, take a look whether it's listed in the current scope, delete it with del, and check that it's gone.

There are also packages in Python. A package is essentially a collection of modules, organized hierarchically in folders. Each folder contains a <code>__init__.py</code> file, telling Python that this folder is special because it contains Python modules.

In terms of using packages, you can import them the same way as modules. We do not further go into writing packages, as most likely it will suffice if you write modules for this course.

A.12 Saving Objects and Data to File

You can save any Python object to a file using **pickle**. The following example is from the official Python documentation.

We also discuss the pickle file format in Section A.13.3 starting on page 186 when discussing Pandas and DataFrames.

A.13 Matrices and Tabular Data in Python

- Python does not have matrices and support for tabular data (i.e. DataFrames) baked into the core language. This means that you need to use add-on packages. This is not necessarily a bad thing, but it illustrates the historically different focus Python has compared to, for example, the R programming language.
- In some simple cases you could actually represent a matrix as a list of list (without using any packages). But as you can imagine, you will quickly see the limitations of this approach once you need to run more complicated computations. In any case, here is an example how you can transpose such a matrix using list comprehensions.

```
>>> matrix = [
...     [1, 2, 3, 4],
...     [5, 6, 7, 8],
...     [9, 10, 11, 12],
... ]
>>> [[row[i] for row in matrix] for i in range(4)]
[[1, 5, 9], [2, 6, 10], [3, 7, 11], [4, 8, 12]]
```

A.13.1 NumPy and SciPy

- In Python you typically need add-on packages if you would like to deal with matrices and/or tabular data: NumPy, SciPy (sometimes), and Pandas (discussed later in Section A.13.2 starting on page 184).
- NumPy is usually imported as follows:

```
import numpy as np
```

• Often you don't import SciPy as a whole; instead you selectively import certain names from it (example follows).

```
from scipy import special, optimize # Just an example to import two names.
```

- For matrices you typically use NumPy (numpy.array) or SciPy (especially for sparse matrices, e.g. scipy.sparse.csc_matrix).
- NumPy stores data in **arrays**, which might be one-dimensional (a vector), two-dimensional (a matrix), or *n*-dimensional (ndarray) in general. For example, here we create a 2-dimensional array of size 2x3, composed of 4-byte integer elements. Keep in mind when indexing the array that indices in Python start at zero (not at one like in R). dtype is the data type of the array.

```
>>> x = np.array([[1, 2, 3], [4, 5, 6]], np.int32)
>>> x
array([[1, 2, 3],
       [4, 5, 6]], dtype=int32)
>>> x.T
                                  # Transpose.
array([[1, 4],
       [2, 5],
       [3, 6]], dtype=int32)
>>> x[:, 0]
                                  # First column.
array([1, 4], dtype=int32)
>>> x[1, :]
                                  # Second row.
array([4, 5, 6], dtype=int32)
>>> type(x)
<class 'numpy.ndarray'>
>>> x.shape
(2, 3)
>>> x.dtype
dtype('int32')
```

• Scipy has several different kinds of sparse matrices, e.g. see scipy.sparse. If you have a sparse matrix M, you can take a look at it by converting it to a dense matrix/array:

```
M.toarray() # Use the 'toarray' method.
M.A # Access the '.A' attribute.
```

- If you have worked before with other software such as MATLAB, don't get confused by the way NumPy stores its data. MATLAB stores data column by column ("Fortran order"), while NumPy by default stores them row by row ("C order"). This does not affect indexing, but may affect performance. For example, in MATLAB, an *efficient* loop will be over columns, e.g. for n = 1:10 a(:, n) end, while in NumPy it's preferable to iterate over rows, e.g. for n in range(10): a[n, :] (note that n is in the first position, not the last).
- Ranges of numbers:

```
>>> np.arange(0, 1, 0.1)
array([ 0. , 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9])
```

A.13.2 Pandas

- See also Appendix B starting on page 203 for a more detailed introduction of Pandas.
- Typically Pandas is loaded as follows:

```
import pandas as pd
```

- For tabular data comprised of different data types (e.g. numbers and names such as company names), you typically want to use a Pandas DataFrame. In case you have used R before, it is basically the same idea as a data.frame in R.
- Pandas has a **series**, which is a one-dimensional labeled array, a **DataFrame**, which is a two-dimensional table with row and column labels, and a **Panel**, which is a labeled three-dimensional data structure (e.g. think for example of a dataset containing time series of sales and net income for IBM and Apple).
- Pandas is built on top of NumPy.
- There are **two main differences between NumPy and Pandas**. The first is that Pandas' data structures have labels, e.g. the DataFrame has row and column labels, while a two-dimensional NumPy array does not. Second, Pandas data structures can contain different data types, e.g. the first column in a Pandas data frame might contain numeric data while the second column contains character strings. On the other hand a NumPy array can only contain a single data type.
- Create a DataFrame:

```
>>> pd.DataFrame(data=[4, 5, 6, 7], index=range(0, 4), columns=['A'])
    A
0    4
1    5
2    6
3    7
```

• Another example uses a **dict as an input**. We use the columns argument to specify the column order. Note that the dict has a different order than the columns in the DataFrame.

• You can look at the head and tail of a DataFrame.

```
df.head()
df.tail()
```

• You can look at the column names by accessing the columns attribute.

```
df.columns
```

A.13.3 Writing/Loading a DataFrame To/From a File

• You can save and load a DataFrame to a file. The simplest format is commaseparate values (CSV), which also lets you easily interchange the data with other programs.

```
df.to_csv('example-data.csv')  # Save data to a CSV file.
df = pd.read_csv('example-data.csv')  # Load from CSV file.
```

• One problem with CSV is that it does not retain information on metadata. For example, just by looking at the CSV file you can only guess what the original data types of the columns are. So CSV is good if you want to exchange your data with other programs, but if you just want to **save and reload your data in Python**, you can **pickle** it (which is a way to serialize the object and save it to disk, see also Section A.12 starting on page 182).

```
df.to_pickle('example-data.pkl')  # Save DataFrame 'df' to file.
df = pd.read_pickle('example-data.pkl')  # Load DataFrame 'df' from file.
```

• If you need to exchange DataFrame data with R (or even if you use it only in Python and want a different way to serialize the data), you can save it in the **feather format**:

```
df.to_feather('example-data.feather')
df = pd.read_feather('example-data.feather')
In R, you can write and read the data as follows:
library('feather')
write_feather(df, 'example-data.feather')
df = read_feather('example-data.feather')
```

• If your Pandas data is very large, you can also store it to the **HDF5 format** via the **PyTables** package. This format is good if you have data exceeding the size of your computer memory. In this case, HDF5 is a way to store it efficiently and retrieve the parts of the data that you need to work on. It is similar in spirit to a database.

```
store['df'] = df  # Save it. Equivalent to 'store.put('df', df)'
store['df']  # Load it. Equivalent to 'store.get('df')'
```

A.14 Iterators in Python

A.14.1 Overview

- Iterators pervade and unify Python. They are a powerful programming concept because they let you do **data streaming**. The basic idea is that you do not need to load the whole dataset into memory (which might be impossible if your dataset is very large). Instead, you process your data in small chunks one by one, thus potentially **saving a lot of memory**. Furthermore, you can **create long**, **data-driven pipelines** as you can feed generators as inputs to other generators. (A generator is a way of creating an iterator in Python.)
- The reason why we discuss iterators is that we need them when discussing the gensim package in Chapter 19 starting on page 131. The gensim package is very popular for doing text analytics in Python. There's also a nice discussion about iterators written by gensim's author Radim Rehurek, upon which this section is partially built.
- Iterators can be pretty powerful, but at the beginning they might sometimes seem a bit confusing. I'll try to explain them clearly here in these lecture notes.
- Be careful: When you google around, there are lots of explanations out there about this concept, but **some of them have their terminology mixed up**, which will only confuse you, especially if you are trying to learn about iterators for the first time. If in doubt, refer to the official Python documentation or just follow along here in these lecture notes. Explanations from Python's glossary:
 - iterator
 - iterable
 - generator
 - generator iterator
 - generator expression
 - sequence
- To avoid confusion, I strongly suggest **not to use "iterable" as a verb**, e.g. don't say: "Is this object iterable?" Instead, **"iterable" should be used as a noun**, e.g. say: "This object is an iterable." The reason is that an "iterable" has a very specific meaning in Python as we will see in Section A.14.3 starting on page 189.
- itertools is a popular module containing a number of iterator building blocks.
- We focus on Python 3 in this course. However, in case you need to deal with iterators and would like to write code that works on both Python 2 and 3, you can use the six library. The **six library** helps with writing code that is **compatible with both Python 2.5+ and Python 3**. For example, it has an iteritems method that

returns an iterator over a dictionary's items. In the code below, iteritems(d) creates an iterator that works with both Python 2 and 3. In Python 2, d.iteritems() creates an iterator. In Python 3, on the other hand, d.items() creates an iterable. In the following example, all three pieces of code below have the same output.

```
d = {'Jack': '#4098', 'Guido': '#4127'}  # Create a dictionary.
# Works in Python 2.
[key + ' has ' + value for key, value in d.iteritems()]
# Works in Python 3.
[key + ' has ' + value for key, value in d.items()]
# Following code works in Python 2 & 3 using the six library.
from six import iteritems
[key + ' has ' + value for key, value in iteritems(d)]
```

A.14.2 Iterators

- An **iterator** is an object that represents a stream of data. You can move from one item to the next by calling the next() function. Once there are no further items produced by the iterator, a StopIteration exception is raised; the iterator is exhausted, i.e. empty. This shows that **you can pass through an iterator once only**. In other words, an iterator is good for one pass over the set of values. If you wanted to cycle through several times, you would have to go back to the beginning and create another iterator from the list.
- Let's create an iterator from a list to illustrate some basic concepts.

```
>>> it = iter([8, 10, 12])  # Create iterator from list.
>>> next(it)
8
>>> next(it)
10
>>> next(it)
12
>>> next(it)

Traceback (most recent call last):
  File "<stdin>", line 1, in <module>
StopIteration
```

• Strictly speaking, each iterator has a __next__() method, which is implicitly called when using the built-in next() function. The example above could have equivalently been written as follows

```
it = iter([8, 10, 12])  # Create iterator from list.
it.__next__()
it.__next__()
```

```
it.__next__()
it.__next__()
```

• Likewise, the built-in iter() function actually calls the object's __iter__() method. The example above could have equivalently been written as follows.

```
it = [8, 10, 12].__iter__()
next(it)
next(it)
next(it)
```

• Instead of basing an iterator on a list, you can use many other objects in Python to generate iterators. We will dwell on this in the following Section A.14.3.

A.14.3 Iterables

• So what does a list have to do with an iterator? A list is not an iterator (you cannot call next() on it), but it can be used to generate an iterator. It is therefore by convention called an "iterable." An **iterable** is an object capable of returning its members one at a time. You can think of an **iterable as something that you can use to create an iterator** by calling the iter() function on the iterable. Specifically, an iterable is an object that manages a single pass over a sequence type (such as list, str, tuple) or some non-sequence type (such as dict, file objects, or a class you define with an __iter__() method).

```
from collections.abc import Iterable, Iterator
l = [8, 10, 12]  # A list is an iterable, but not an iterator.
isinstance(l, Iterable)  # True. A list is an iterable.
isinstance(l, Iterator)  # False. A list is NOT an iterator.
it = iter(l)  # Create iterator.
isinstance(it, Iterable)  # True. An iterator is an iterable.
isinstance(it, Iterator)  # True. An iterator is an iterator (duh).
```

- As we have just seen, **any iterator is also an iterable**. Why is that the case? It is because when you apply the iter() function on an iterator, the iterator will always return itself. So in this sense, it can generate an iterator (i.e. itself) and thus by definition is an iterable.
- Recap: All iterators are also iterables. But not all iterables are iterators. You
 can think of the set of all iterables being a strict superset of the set of all
 iterators. In symbols:

```
\{\text{iterators}\} \subseteq \{\text{iterables}\}
```

• In this context, it is important to understand how the for loop works internally. Let's consider for illustration the following example, which simply prints the numbers 8, 10, and 12 based on my_list (which, as we know, is an iterable).

```
my_list = [8, 10, 12]
for x in my_list:
    print(x)
```

What happens behind the scenes (done automatically by the for loop) is that **an iterator is created** by calling iter(my_list). This iterator has a __next__() method, and each time this method is called, its return value is assigned to x and the loop is executed. This goes on until a StopIteration exception is raised when calling __next__(), in which case the iterator is empty and the loop exits.

- So the for loop in Python internally uses an iterator!
- As we have noted before, an iterator is also an iterable, so let's see **what happens when we use an iterator in a for loop**. The first time, everything works as expected. The for loop calls iter(my_it), which returns itself (i.e. my_it) and then uses next() to get each successive item and assigns it to x. However, the next time we run the loop, nothing happens! The reason is simply that the iterator is exhausted, so there are no items left in the iterator. If we want to get some output, we need to get new iterator (e.g. run my_it=iter(my_list) again).

```
>>> my_list = [8, 10, 12]
>>> my_it = iter(my_list)
                             # Create iterator.
>>> for x in my_it:
                             # Works as expected.
        print(x)
. . .
8
10
                             # No output! Iterator is exhausted!
>>> for x in my_it:
        print(x)
>>> my_it = iter(my_list)
                             # Need to get a new iterator ...
>>> for x in my_it:
                             # ... to get some output.
        print(x)
. . .
8
10
12
```

• You can have a similar example when you want to convert the iterator back to a list. The second time, the iterator is exhausted and you get an empty list.

class my_iterator:

```
>>> my_list = [8, 10, 12]
>>> my_it = iter(my_list)  # Create iterator.
>>> list(my_it)  # Works as expected.
[8, 10, 12]
>>> list(my_it)  # Empty list because iterator is exhausted!
[]
>>> list(my_it)  # ... still empty!
[]
```

• Let's create an iterator manually. As mentioned, an iterator is what we ultimately care about because **an iterator represents a stream of data**. For illustration, we implement an iterator that is similar to range() (which is an immutable sequence type and as such it is an iterable), with the difference that our implementation creates an iterator directly (unlike range(), our implementation does not create an iterable). As mentioned in Section A.14.2, it needs a __iter__() method, which simply returns itself. It also needs a __next__() method, which yields the next item (or throws a StopIteration exception at the end). The first thing we do therefore is to define a class with those two methods (and an additional __init__() constructor method to initialize some variables when the class is instantiated).

```
def __init__(self, n): # Initialize some variables.
    self.i = 0
    self.n = n

def __iter__(self): # An iterator needs to return itself here.
    return self

def __next__(self): # This method yields the next item.
    if self.i < self.n:
        self.i += 1
        return self.i - 1
    else:
        raise StopIteration() # Raise StopIteration exception when exhausted.</pre>
```

Second, we need to instantiate this class and assign it to a variable, which will contain the iterator.

```
it = my_iterator(3) # Create new iterator by instantiating the class.
```

Third, we can check that the iterator works as expected.

```
>>> next(it)
0
>>> next(it)
```

1 2

···

>>> for x in it:

... print(x)

```
1
>>> next(it)
2
>>> next(it)  # Iterator is exhausted.
Traceback (most recent call last):
   File "<stdin>", line 1, in <module>
   File "<stdin>", line 15, in __next__
StopIteration

Also, as expected, the iterator is exhausted after finishing a given for loop.
>>> it = my_iterator(3)  # Create iterator.
>>> for x in it:
...  print(x)
...
0
```

• How could we **turn this iterator into an iterable** similar to range()? Remember that, using its __iter__() method, an iterable returns an iterator. That's exactly what we are going to do in the following class definition.

Iterator is exhausted, no output.

```
class my_iterable:
    def __init__(self, n):
        self.n = n

def __iter__(self):  # Return an iterator here.
    return my_iterator(self.n)
```

Then all we have to do is instantiate this class to actually create an iterable.

```
itrble = my_iterable(3) # Create an iterable.
```

And we can check that it works as expected.

```
>>> it = iter(itrble)  # Create an iterator from the iterable.
>>> next(it)
0
>>> next(it)
1
```

```
>>> next(it)
2
>>> next(it)
Traceback (most recent call last):
   File "<stdin>", line 1, in <module>
   File "<stdin>", line 14, in __next__
StopIteration
```

Because itrble is an iterable (and not an iterator), we can now for example use it several times in a for loop. Each time another for loop starts, a new iterator is created for that loop, because the for loop internally calls iter(itrble).

```
>>> for x in itrble:  # Creates new iterator for this for loop.
... print(x)
...
0
1
2
>>> for x in itrble:  # Creates another iterator for this for loop.
... print(x)
...
0
1
2
```

A.14.4 Generators

- A generator is a tool for creating an iterator with relative ease. Instead of manually creating an iterator like we have done in Section A.14.3 starting on page 189, it is often easier to create an iterator from a generator.
- Anything that can be done with generators can be done with class-based iterators (e.g. as done in my_iterator in Section A.14.3). However, what makes generators so compact is that they create the __iter__() and __next__() methods automatically.
- Strictly speaking, when talking about a generator we usually mean a generator function, which creates a generator iterator object.
- Sometimes people also use the term *generator* to refer to a *generator iterator*, which depends on the context and admittedly is a bit of a sloppy terminology. If you want to use sloppy terminology, I think it is better to call a generator iterator simply "iterator" (because that's what it really is).
- Here's a simple example illustrating a generator function and a generator iterator. We begin by defining a generator function. This generator function determines how the iterator (which will later be derived from it) will behave.

```
def gf():  # Define a "generator function".
   yield 1
   yield 2
   yield 3
```

Next we create a generator iterator from the generator function. The key thing is that you are pausing each time there is a yield expression until the following call to next() is made. As you can see, at the end, the generator iterator is exhausted as calling next() results in an error, so you can only consume the values of a generator iterator once. This is exactly the same behavior as we have seen for iterators in Section A.14.2 starting on page 188.

```
>>> gi = gf()
                    # Create a "generator iterator."
>>> next(gi)
                    # Retrieve next item from the iterator.
1
>>> next(gi)
                    # Retrieve next item from the iterator.
>>> next(gi)
                    # Retrieve next item from the iterator.
>>> next(gi)
                     # Raises StopIteration since iterator is empty.
Traceback (most recent call last):
 File "<stdin>", line 1, in <module>
StopIteration
>>> gi = gf()
                    # Again create a "generator iterator".
>>> next(gi)
                    # Retrieve next item from the iterator.
1
>>> next(gi)
                    # Retrieve next item from the iterator.
>>> next(gi)
                    # Retrieve next item from the iterator.
>>> next(gi)
                     # Raises StopIteration since iterator is empty.
Traceback (most recent call last):
  File "<stdin>", line 1, in <module>
StopIteration
```

• For illustration, you can also use a generator function in a for loop. As usual, the for loop internally takes care of creating the iterator and calling next(). We have briefly discussed this in Section A.14.3 starting on page 189. As you can see, you can go through several for loops because, based on the generator function, a new generator iterator is created each time a new for loop is encountered. In other words, each time a generator function is called, it creates a new iterator.

```
>>> for value in gf():
... print(value)
```

```
1
2
3
>>> for value in gf():
... print(value)
...
1
2
3
```

• Here is another way to write and use a generator function. What we do here is put the yield expression into a for loop. Everything else, e.g. the creation of the generator iterator and its behavior, is the same as before.

```
>>> def gf():
                             # Define the generator function.
       for x in range(3):
            yield x**2
>>> gi = gf()
                             # Create a generator iterator.
>>> next(gi)
>>> next(gi)
>>> next(gi)
>>> next(gi)
Traceback (most recent call last):
  File "<stdin>", line 1, in <module>
StopIteration
>>> for value in gf():
                            # Using the generator function in a for loop.
        print(value)
. . .
0
1
4
```

A.14.5 Generator Expressions

- **Generator expressions** are shorthand ways to create generator iterators.
- A generator expression is equivalent to creating an anonymous generator function and calling it (thus creating a generator iterator).
- A generator expression is conceptually related to a list comprehension, but the key difference is that it conserves memory by creating the items one at a time, and only when they are needed.

• Here's an example, where we create a generator iterator gi using a generator expression:

```
>>> gi = (i*i for i in range(10)) # No calculations done at this stage!
>>> sum(gi) # Calculations done here using the generator iterator.
285
>>> sum(gi) # Generator iterator is empty.
0
```

• Often you would use generator expressions and the resulting calculations in one line of code. In this case, you can omit one set of brackets (this is just syntactic sugar to save some typing).

```
sum((i*i for i in range(10)))
sum(i*i for i in range(10))  # Equivalent. Syntactic sugar.
```

• Keep in mind that when you put a generator iterator into a for loop, it will be exhausted (i.e. empty) after the first run. This is the same behavior we saw with iterators in Section A.14.3 starting on page 189.

```
>>> gi = (word + '!' for word in 'I love NLP'.split())
>>> for val in gi:
...     print(val)
...
I!
love!
NLP!
>>> for val in gi:
...     print(val)
...
>>>
```

• On the other hand, if you put a generator expression into the for loop, it will supply a new generator iterator each time a new for loop is run.

```
>>> for val in (word + '!' for word in 'I love NLP'.split()):
...    print(val)
...
I!
love!
NLP!
>>> for val in (word + '!' for word in 'I love NLP'.split()):
...    print(val)
...
I!
love!
NLP!
```

A.14.6 Creating Iterables the Easy Way Using Generators

- This section is the culmination of our previous efforts in understanding iterators. Specifically, in this section we describe the probably easiest way to write a custom iterable.
- Remember the iterable my_iterable defined earlier in Section A.14.3 starting on page 189. Simplifying this example, we are going to use a generator function to create an iterable.
- What we did in Section A.14.3 was to define a class with an __iter__() method that returned an iterator. Specifically, it returned my_iterator which was our custom-built iterator defined separately before. Instead of going through all the trouble and defining a custom iterator, we can simply create one using a generator function.
- Specifically, we write __iter__() as a generator function, which (as we know from Section A.14.4) always returns a generator iterator.

```
class my_iterable2:
    def __init__(self, n):
        self.n = n

def __iter__(self):  # Generator function.
    for x in range(self.n):
        yield x**2
```

- We have thus written a class for an iterable, because a call to the __iter__() method creates a new iterator (by using a generator function).
- We can now instantiate this class in order to create an iterable.

```
itrble2 = my_iterable2(3)
```

• This iterable can be used in the usual way. Each time a new for loop starts, the iterable itrble2 is used to create a new iterator for that loop.

0 1 4

• Equivalently, instead of using a generator function, we can also **use a generator expression to create an iterable**. Specifically, the __iter__() method simply returns the call to an generator expression, for which we know from Section A.14.5 starting on page 195 that this is an iterator.

```
class my_iterable3:
    def __init__(self, n):
        self.n = n

def __iter__(self):
    # Return an iterator created from a generator expression.
    return (x**2 for x in range(self.n))
```

A.14.7 Summary of Iterables and Iterators

- An iterable is an object that is capable of returning its members one at a time. For example, lists or tuples are iterables.
- Iterables can be used in a for loop or when a sequence is needed, e.g. zip() or map(). See the following examples. Side note: zip() or map() return iterators, so to view their contents in the following examples, we convert their outputs back to lists.

```
>>> myiterable_a = [1, 2, 3]  # An iterable.
>>> myiterable_b = [4, 5, 6]  # Another iterable.
>>> for i in myiterable_a:
...  print(i)
...
1
2
3
>>> list(zip(myiterable_a, myiterable_b))
[(1, 4), (2, 5), (3, 6)]
>>> list(map(lambda x: 2 * x, myiterable_a))
[2, 4, 6]
```

- You can **use an iterable to create an iterator** by calling the iter() function on the iterable, e.g. iter([8, 10, 12]).
- You need an iterator because **an iterator is the object that gives you the actual stream of data** one at a time.

- Every iterator is also an iterable. When calling the iter() function on an iterator, the iterator will always return itself. It thus satisfies the requirement of an iterable, i.e. that calling the iter() function returns an iterator.
- A difference between an iterable and an iterator is that (unless the iterable is an iterator) you cannot call next() on an iterable, while you can always call next() on an iterator. For example, next([1, 2]) will not work, because the list [1, 2] is not an iterator.
- You can pass through an iterator only once. Afterwards, it is exhausted.
- If you use an iterable in the for loop (e.g. for i in [1, 3, 5]:), you can run the same for loop again and and again. On the other hand, if you use an iterator, the for loop will only work once, because the iterator is exhausted afterwards.
- A **custom iterable can be created** by defining and then instantiating a class that has an __iter__() method. This __iter__() method must return an iterator. See Section A.14.3 starting on page 189 and Section A.14.6 starting on page 197.
- Iterators can be created in several ways:
 - 1. Calling the iter() function on an iterable, e.g. iter([8, 10, 12]) or iter(range(8)). See Section A.14.2 starting on page 188
 - 2. Calling the iter() function on an iterator. (The iterator will just return itself.)
 - 3. Define and then instantiate a custom class that has __iter__() and __next__() methods. The __iter__(self) method returns self, i.e. it returns an iterator. This is necessary because every iterator has to be an iterable, and an iterable requires that iter() (or equivalently __iter__()) returns an iterator. The __next__() method returns the next item of the iterator. See Section A.14.3 starting on page 189.
 - 4. Define and then call a generator function, see Section A.14.4 starting on page 193.
 - 5. Use a generator expression, see Section A.14.5 starting on page 195.

A.15 Functional Programming

While Python is often used for object-oriented programming (see Section A.7 starting on page 166), it is also possible to write your programs using functional programming. The basic idea is to decompose a problem into a set of functions.

A.15.1 map

The map function goes through an iterable and applies a given function to each element. For example:

```
>>> mylist = ['I', 'love', 'NLP']
>>> result = map(len, mylist)  # Apply then 'len' function to 'mylist'.
>>> for x in result:
... print(x)
...
1
4
3
```

Another example illustrates a function with two arguments:

```
>>> def f(x, y):
... return x + y
...
>>> result = map(f, ['hello', 'world'], ['foo', 'bar'])
>>> for x in result:
... print(x)
...
hellofoo
worldbar
```

As discussed in Section B.3 starting on page 204, there is an apply() method for DataFrames in Pandas, which works in a similar fashion as map.

A.15.2 Anonymous Functions

Sometimes you just need to use a function once, in which case it doesn't make a lot of sense to give it a name. Thus, Python has anonymous functions, i.e. functions without a name. Consider the following example, where we define a function and just use it once.

```
>>> def myfunction(z):  # A waste of space, we just need it once!
... return z + 5
...
>>> result = map(myfunction, [10, 11, 12]) # Here is where we need it.
>>> for x in result:
... print(x)
...
15
16
17
```

Instead, we can omit giving a name to the function, and instead just use an anonymous function, indicated by lambda. The following code produces the same output as the previous code, but does it in a much more concise way.

```
>>> result = map(lambda z: z + 5, [10, 11, 12])
>>> for x in result:
```

```
... print(x)
...
15
16
17
```

A.15.3 filter

Sometimes you have an iterable and would like to remove some elements. For example, you have stock price returns and would only like to keep the ones that are 1% or higher.

Another example that uses an anonymous function to keep only the odd numbers. Remember that % refers to the modulo operation, i.e. it finds the remainder after division. For example, x % 2 divides x by two and returns the remainder. Below we check whether the remainder is not equal to zero, which means that the number x is odd. In this case, the anonymous function returns a True value, telling filter to keep the corresponding element of sequence.

```
>>> sequence = [2, 3, 4, 5, 6, 7]
>>> result = filter(lambda x: x % 2 != 0, sequence)
>>> for x in result:
... print(x)
...
3
5
7
```

A.15.4 reduce

Sometimes you would like to summarize an iterable (e.g. a list) with a single value. For example, you would like to sum all elements in a list, or you would like to compute the maximum. Oftentimes, there are already functions available in Python that do these

things, e.g. sum() and max(). Still, sometimes you would like have customized behavior, in which case reduce comes in handy.

In the following examples, we add all elements in the list and we also find the maximum. The purpose of these examples is to illustrate how reduce() works. The purpose is not to show how to do summation or finding the maximum, as these could be done more easily using sum() and max() as discussed above.

The way reduce() works is that it has a function of two arguments, and iteratively applies this function to an iterable (e.g. a list). In the summation example below, it would first compute 5+1=6, and then it would compute 6+8=14, which is the end result. Or in the case of the maximum, it would first compute $\max\{5,1\}=5$, and then it would compute $\max\{5,8\}=8$.

```
>>> import functools
>>> mylist = [5, 1, 8]
>>> def myaddition(x, y):
...    return x + y
...
>>> def mymax(x, y):
...    if x > y:
...       return x
...    else:
...       return y
...
>>> functools.reduce(myaddition, mylist)
14
>>> functools.reduce(mymax, mylist)
8
```

Equivalently, the code above could have been written more succinctly using anonymous functions:

```
>>> import functools
>>> mylist = [5, 1, 8]
>>> functools.reduce(lambda x, y: x + y, mylist)
14
>>> functools.reduce(lambda x, y: x if x > y else y, mylist)
8
```

One could also use the operator module, as it provides some commonly-used functions. For example, operator.add(x, y) does the same thing as the expression x + y.

```
>>> from functools import reduce
>>> from operator import add, mul
>>> mylist = [5, 1, 8]
>>> reduce(add, mylist)  # Addition.
14
>>> reduce(mul, mylist)  # Multiplication.
40
```

Appendix B

Data Munging with Pandas

B.1 Introduction to Pandas

In this section we take a closer look at the Pandas Python package. We have briefly covered it before:

- 1. Section A.13 starting on page 183, mostly by describing how it extends NumPy and how to save and load your data.
- 2. Section 20.1 starting on page 147, mostly from a conceptual and historical point of view.
- 3. In contrast, in this current chapter we go into more depth on how to use Pandas based on example data files containing stock and accounting data (available from the course website).

When Pandas was created, the key idea was to enable Python users to perform similar tasks as were already common in R. The most important element was to carry the data.frame concept from R (i.e. rectangular data where columns can be of different data types) over to Python. It is fair to say that Pandas is probably the cornerstone of data science in Python and the main reason why data science has caught on in Python so well in the past. The creator of Pandas is Wes McKinney, an MIT graduate who wrote the first Pandas versions while working at AQR, a famous quantitative hedge fund.

Pandas can be useful in at least three ways.

- 1. You have analyzed your textual data and have obtained some sort of scores or values based on the textual analysis, e.g. sentiment scores. You can then analyze them in Pandas.
- 2. Your textual data is saved as tidy text in a Pandas DataFrame and you would like to perform further analysis on this data. See Section 14.7.2 starting on page 110 for details about tidy text.
- 3. If you have financial data, e.g. stock market data from AlphaVantage (see also Section B.6 starting on page 214) or accounting data from Morningstar, you can merge them together and analyze them with your textual data.

B.2 Introductory Pandas Example

Here we illustrate basic pandas usage. The code below can be found in the file code-pandas-020-intro.py on the course website.

```
# This script gives a brief introduction on pandas in Python.
import pandas as pd
import numpy as np
df = \setminus
    pd.DataFrame(
        np.random.randn(6, 4), # 6x4 numpy array with random numbers.
        index=pd.date_range('2019-12-20', periods=6),
        columns=['A', 'B', 'C', 'D'])
df.head()
                                 # Top rows.
                                 # Bottom 3 rows.
df.tail(3)
df['A']
                                 # Look at column named 'A'.
df[0:3]
                                 # First three rows.
df.apply(lambda x: x.max() - x.min()) # Apply a function to the data.
# The key thing about a DataFrame is that its columns can hold data of
# different types.
df2 = \
    pd.DataFrame(
        \{ 'A' : 1., \}
          'B': pd.Timestamp('2019-12-20'),
          'C': pd. Series(1, index=list(range(4)), dtype='float32'),
          'D' : np.array([3] * 4, dtype='int32'),
          'E' : pd.Categorical(["test", "train", "test", "train"]),
          'F' : 'foo' })
df2.dtypes
                                 # Look at data types.
```

B.3 Basic Pandas Usage

Below we show how to slice and dice data saved as a CSV file. It is an excerpt from the Center for Research in Security Prices (CRSP) from the University of Chicago. The data file data-CRSP-extract.csv can be found on the course website. The variables included are:

- 1. TICKER: The ticker symbol of the stock.
- 2. date: The date in which the stock traded.
- 3. PRC: The stock price.
- 4. RET: The stock return.

- 5. VOL: Number of shares traded in millions.
- 6. SHROUT: Number of shares outstanding.

The beginning of the CSV file looks like this, so you can see that different columns are separated by commas:

```
TICKER, date, PRC, RET, VOL, SHROUT

AAPL, 2014-01-02, 553.13, -0.014064, 8338094, 892.447

AAPL, 2014-01-03, 540.97998, -0.021966, 13992006, 892.447

AAPL, 2014-01-06, 543.92999, 0.005453, 14820614, 892.447

AAPL, 2014-01-07, 540.03748, -0.007156, 11381939, 892.447
```

After importing into Pandas (see the following code example, in particular pd.read_csv), the beginning of the dataset looks like this:

```
TICKER date PRC RET VOL SHROUT

0 AAPL 2014-01-02 553.13000 -0.014064 8338094 892.447

1 AAPL 2014-01-03 540.97998 -0.021966 13992006 892.447

2 AAPL 2014-01-06 543.92999 0.005453 14820614 892.447

3 AAPL 2014-01-07 540.03748 -0.007156 11381939 892.447
```

The following code contains in-line comments that explain what's going on. The code below can be found in the file code-pandas-040-longer-intro.py on the course website and the data file we use is also on our course website.

```
import pandas as pd
import numpy as np
df = pd.read_csv('data-CRSP-extract.csv')
df.dtypes
df['date'] = pd.to_datetime(df['date'])
\# df['date'] = [time.date() for time in df['date']] \# Get rid of time.
df['date'].head() + np.timedelta64(1, 'D') # Add one day.
# Basic information about the DataFrame.
type(df)
list(df)
                        # Column names.
df.head()
                        # Top \ of \ df.
df.tail()
                        # Bottom of df.
                        # Data types of columns.
df.dtypes
df.shape
                        # Dimensions
                        # Number of rows.
len(df.index)
len (df. columns)
                       # Number of columns.
df.describe
                        # Basic summary statistics.
# Indexing and selecting data.
df[0:3]
                       # First three rows.
```

```
df.head(3)
                       # First three rows.
df.iloc[:3]
                       # First three rows.
                       # Last row.
df[-1:]
df.tail(1)
                       # Last row.
df.TICKER
                       # Same as df['TICKER'], returns a Series.
df[['TICKER']]
                       # Returns DataFrame. Inner brackets are a list.
df[['TICKER', 'PRC']] # DataFrame with two columns.
# Apply a function to each column of the DataFrame. (If you would like
# to apply a function to each row of a DataFrame, use the 'axis=1'
# parameter.) As the functions below expect numeric types, we only
# apply them to the price (PRC) and stock return (RET) columns for
# illustration.
df[['PRC', 'RET']].apply(np.max) # Maximum of each column.
df[['PRC', 'RET']].apply(lambda x: x + 3) # Add three to each column.
# Querying the data (selecting rows that satisfy some condition(s)).
df[df.TICKER == 'FB']
                                         # Extract Facebook.
df.query("TICKER == 'FB'")
                                         # Extract Facebook.
df[(df.TICKER == 'FB') \& (df.PRC > 120)] # Facebook with price > 120.
df.query("TICKER == 'FB' & PRC > 120") # Facebook with price > 120.
df[(df.TICKER == 'FB') \mid (df.PRC < 120)] \# Facebook or price < 120 (or both).
df.query("TICKER_==_'FB'_|PRC_<_120") # Facebook or price<120 (or both).
df[df.TICKER.isin(['AAPL', 'FB'])]
                                    # Apple and Facebook.
df.TICKER.unique()
                                       # All tickers.
df[['TICKER', 'SHROUT']].drop_duplicates() # Two columns w/ unique entries.
# Modify the data.
df.rename(columns={'VOL': 'Volume'}) # New DataFrame with renamed column.
df.rename(columns={'VOL': 'Volume', 'PRC': 'Price'}) # Two columns renamed.
df.assign(HIGH\_PRC = (df.PRC > 120))
                                     # New column created.
df.assign(abc = df.PRC + 8 * df.SHROUT) # New column.
df.drop('TICKER', axis = 1)
                               # New DataFrame without this column.
df.sort_values(by='PRC')
                                # Sort increasing by PRC.
df.sort_values(by='PRC', ascending=False) # Sort decreasing by PRC.
df.sort_values(by=['SHROUT', 'PRC'], ascending=[False, True])
# Aggregating (summarizing) the data.
df.PRC.max()
                                # Maximum price.
df.PRC.min()
                                # Minimum price.
df.agg('max')
                                # Max applied to all columns.
df.agg(['max', 'min'])
                                # Max and min applied to all columns.
                                # Different aggregations per column.
df.agg({
    'PRC': ['sum', 'max'],
    'VOL': ['max', 'min']})
# Group by ticker and then for each group calculate the maximum and
# the minimum on each column.
```

```
df.groupby('TICKER').agg(['max', 'min'])
# Maximum price for each ticker.
df.groupby('TICKER').agg({'PRC': 'max'})
# Maximum price for each ticker and each group of shares outstanding.
df.groupby(['TICKER', 'SHROUT']).agg({'PRC': 'max'})
# Count how many observations have a high price (i.e. PRC>120).
df.\
    assign(HIGH_PRC = (df.PRC > 120)). \setminus
    groupby('HIGH PRC').\
    agg({'HIGH_PRC': 'count'}).\
    rename(columns={'HIGH_PRC': 'Count'})
pd.DataFrame(
                                 # Summary statistics on subset.
    {'Max\_PRC': [df[0:10].PRC.max()], \setminus}
     'Min_PRC': [df[0:10].PRC.min()], \
     'Std_PRC': [np.std(df[0:10].PRC)]})
# Total trading volume for each ticker, assigned to a new column.
df['TVOL'] = df['VOL'].groupby(df['TICKER']).transform('sum')
# Removing columns.
del df['TVOL']
                               # Delete column.
df.head()
                                 # Column is gone.
\# df.drop('TVOL', axis=1, inplace=True) \# Alternative way to delete column.
\# df.drop(df.columns[[0, 1, 3]], axis=1, inplace=True) \# Delete by column number.
```

B.4 Merging Two Datasets

We can also merge two DataFrames. Merging (or "joining" as it is sometimes called) means bringing two Pandas DataFrames together. This can be useful not only if you have two DataFrames, but also when you are aggregating (i.e. summarizing) data.

B.4.1 Using merge() for Data Aggregation

In the following example we have daily CRSP data and would like to add a new column that shows the monthly trading volume. We thus **aggregate** the data in the sense that several data points (e.g. daily trading volume) are summarized into fewer data points (e.g. monthly trading volume). Afterwards we merge the monthly trading volume back to the original daily dataset.

Please take a look at the comments in the code that explain what's going on. The code below can be found in the file code-pandas-060-monthly-trading-volume.py on the course website and the data files we use are also on our course website.

```
# This script shows how to calculate monthly trading volume and merge
# it back to daily data.

import pandas as pd
import numpy as np
```

```
df = pd.read_csv('data-CRSP-extract.csv') # Import from CSV file.
df['date'] = pd.to_datetime(df['date']) # Convert to date (and time).
df = df[['TICKER', 'date', 'PRC', 'VOL']] # Just need a few columns.
# Calculate monthly trading volume. We first extract the columns we
# need from the original 'df' DataFrame, group it by ticker and month,
# and sum up the (monthly) volume.
mvol = df[['TICKER', 'date', 'VOL']]
mvol = mvol.groupby(['TICKER', pd.Grouper(key='date', freq='M')]).sum().reset_index
mvol.rename(columns={'VOL': 'MVOL'}, inplace=True) # Rename column.
# Take a look at the data. You can see that here we have ONE
# observation per ticker and per month.
mvol.tail()
# Here we create a column in each DataFrame that has the day removed,
# so we only have year and month. This is necessary for merging both
# DataFrames ('df' and 'mvol') together in a later step where we use
# the year-month information to link up both DataFrames.
mvol['mdate'] = mvol.date.dt.to_period('M') # Convert to monthly dates.
del mvol['date']
                                             # Not needed any more.
df['mdate'] = df.date.dt.to_period('M') # Convert to monthly dates.
# Convert year-month to string. Although not strictly necessary, it
# makes some things easier to handle, e.g. querying a DataFrame in the
# subsequent step.
df['mdate'] = df.mdate.astype(str)
mvol['mdate'] = mvol.mdate.astype(str)
# Here we select some subsets of both DataFrames. In production, you
# would SKIP this step. Here we use it only to better illustrate the
# following merge to have a small result we can actually look at and
# inspect.
df = df.query("TICKER_==_''AAPL'_&_date_>=_'2014-12-20'_&_date_<=_''2015-01-15'")
mvol = mvol[(mvol.TICKER == 'AAPL') & (mvol.mdate >= '2014-11') & (mvol.mdate <= '2014-11')
# Next we merge both DataFrames together using the ticker symbol
# ('TICKER') and a column representing the year-month ('mdate'). Here
# we are using a so-called "left join," which is for our purposes the
# most important kind of join. It means that we are adding data to the
# "left" DataFrame 'df' (instead of the "right" DataFrame 'mvol'). The
# data added comes from the "right" DataFrame 'mvol'. We use "left"
 \# \ and \ "right" \ because \ in \ the \ call \ to \ `pd.merge', \ the \ `df' \ DataFrame 
# comes first, so is on the "left" side, while 'mvol' is on the
# "right" side.
```

df_mvol = pd.merge(df, mvol, how='left', on=['TICKER', 'mdate'])

```
# Take a look at the two original DataFrames 'df' and 'mvol' and the
\# merged DataFrame 'df_mvol'. It will hopefully become clear what has
# happened during the merge, i.e. information about monthly trading
# volume from 'mvol' has been added to 'df'. Observe that if there is
# a row in 'mvol' with no match in 'df', then this row from 'mvol'
# will NOT be included in the merged DataFrame 'df_mvol'. For example,
# the rows from November and February 2014 are in 'mvol', but they are
# NOT added to 'df_mvol' because there is no matching row in 'df'.
df
mvol[[ 'TICKER', 'mdate', 'MVOL']]
df_mvol
# Finally we can delete the 'mdate' column as we don't need it any
del df_mvol['mdate']
# Here is our final result, with monthly trading volume added to each
# observation (i.e. to each row).
df_mvol
```

B.4.2 Merging Stock Market and Accounting Data

In the next example we have stock market data from CRSP and accounting data from Compustat. The variables in Compustat are too numerous to list here, but we can give an explanation of the most important ones used in the example below:

- 1. tic: The ticker symbol of the company's stock.
- 2. datadate: The reporting date. In the example below we have quarterly data. Keep in mind that the data might not actually be released on the reporting date. For example, a balance sheet dated December 31 might be released in January or February due to the time required to prepare adjusting entries, write footnotes, and perhaps be audited or reviewed by a CPA firm. To be on the safe side, we wait for three months before using these numbers and merge them to the stock market data.
- 3. atg: A company's total assets.

We want to add the latest accounting data to each day in CRSP. Here we use the pandasql package which allows us to join different datasets based on date ranges. It works by converting the Pandas DataFrame to SQLite, running your SQL queries, and converting back to DataFrame. You can install it as usual with the following command (see also Section 4.6 starting on page 27 how to install it using pip and virtualenv, which is the recommended way).

```
python3 -m pip install -U pandasql
```

Or alternatively, if you are using Anaconda, you can install it with

conda install -c anaconda pandasql

So without further ado, here is the code. We first import the data and then merge it using pandasql. The important thing is that we then make sure we only get the last observation for each ticker-date pair. The code below can be found in the file code-pandas-060-merging-data.py on the course website and the data files we use are also on our course website.

```
import pandas as pd
import numpy as np
from pandasql import sqldf
pysqldf = lambda q: sqldf(q, globals()) # Shortcut function.
# Import CRSP and convert the date strings to 'datetime64'.
crsp = pd.read_csv('data-CRSP-extract.csv')
crsp['date'] = pd.to_datetime(crsp['date'])
# Same for Compustat.
cs = pd.read_csv('data-Compustat-extract.csv')
cs['datadate'] = pd.to_datetime(cs['datadate'])
# Next we define the time window during which the accounting
# information from Compustat is valid. First, we wait for three months
# before using any Compustat data. The reason is that in real life,
# the data is often released with a lag. For example, if a company's
# balance sheet is as of December 31, 2014, it is often released in
# January or February 2015. To be on the safe side, we therefore add
# three months to 'datadate'. This date corresponds to the BEGINNING
# of the time window we will use when merging.
cs['datadate'] = cs['datadate'] + np.timedelta64(121, 'D')
# Second, we let the accounting data expire after about one year. This
# date corresponds to the END of the time window we will use when
# merging.
cs['datadate2'] = cs['datadate'] + np.timedelta64(380, 'D')
# Extract a subset of the data for better illustration of the
# following merge. 'PRC' is the stock price and 'atq' is total assets.
crsp = crsp[['TICKER', 'date', 'PRC']]
cs = cs[['tic', 'datadate', 'datadate2', 'atq']]
# Inspect the data. Our GOAL in the remaining code is to add Compustat
# data to CRSP such that 'date' (from CRSP) lies in the time window
# given by 'datadate' and 'datadate2' (from Compustat).
crsp.tail()
cs.tail()
# Here we use SQL syntax to merge the two DataFrames using a left join
# (i.e. we add data to CRSP from Compustat because the 'crsp'
```

```
# DataFrame is mentioned to the "left" side of the 'cs' DataFrame in
# the 'FROM ... ' part of the SQL query). The following merge could be
# done in a more memory-efficient way completely in SQL. But for
# simplicity we show this version first. 'SELECT * ' means to select
# all columns from the resulting merged data. In the 'ON...' part, in
# order to tell SQL the DataFrame and column it should use, we add the
# DataFrame name in front of the column, separated by a dot. For
# example, 'crsp.TICKER' means that we are referring to the 'TICKER'
# column from the 'crsp' DataFrame.
df1 = \
    pysqldf(
        'SELECT_*_' + \
        'FROM_crsp_LEFT_JOIN_cs_' + \
        'ON crsp.TICKER=cs.tic AND ' + \
        'crsp.date_BETWEEN_cs.datadate_AND_cs.datadate2')
# Clean up the result from the merge. PandaSQL converts 'datetime64'
# to 'str'. So we need to convert it back to 'datetime64' here.
df1['date'] = pd.to_datetime(df1['date'])
df1['datadate'] = pd.to_datetime(df1['datadate'])
df1['datadate2'] = pd.to_datetime(df1['datadate2'])
# Let's take a look at the resulting merge. The first thing you should
# notice is that 'date' is always between 'datadate' and
# 'datadate2'. So we were successful in adding data to CRSP based on
# the time window specified in Compustat through 'datadate' (the
# beginning of the time window) and 'datadate2' (the end of the
# window).
df1.tail()
# There's still one problem we need to solve. The merge has added ALL
# rows from Compustat that fit the merging criteria, i.e. whose time
# window contains a 'date' from CRSP. So for each observation
# originally in CRSP, we now have around FOUR observations. One for
# each quarter from Compustat whose one-year time window between
# 'datadate' and 'datadate2' matches a given 'date' from CRSP. If you
# look closely at the 'date' column, you will see that many dates are
# repeated.
df1.tail()
# You can look at these multiple matches from a different angle by
# counting how many matches from Compustat we have on average for each
\# Ticker-date combination. For illustration, we look at the 'size()'
# of each ticker-date group and save it in the column name
# 'counts'. This gives us the number of observations we have for each
# ticker and each date. Most of these values turn out to be four as we
# typically have four observations in Compustat per year (i.e. four
# quarters). To further summarize this, we then calculate the mean of
# the 'count' column for each ticker. (When entering the code below,
# make sure to add NO SPACE after the the backslashes '\', otherwise
# it won't work.)
```

```
df1.\
    groupby(['TICKER', 'date']).\
    size().
    to_frame(name='counts').\
    reset_index().
    groupby('TICKER').\
    agg({ 'counts ': 'mean'})
# To finally solve this problem of the multiple matches, we only keep
# the most recent match from Compustat and discard all the others. To
# do that, we first need to sort by 'datadate' to ensure we are
# picking the latest information from Compustat. (Keep in mind that
# the following code works because 'groupby()' preserves the order.)
# Then we grab the latest observation in each ticker-date group using
# 'tail()' and thus discard all the earlier observations.
df1 = df1.sort_values('datadate').groupby(['TICKER', 'date']).tail(1)
df1 = df1.sort_values(by=['TICKER', 'date']).reset_index(drop=True) # Cosmetics.
            # Inspect the data. Now there is only ONE date per ticker.
df1.tail()
del df1['datadate2']
# The approach above works, but it is a bit clumsy. First, it might
# require a lot of memory if your data size gets larger because the
# SQL query keeps a lot of data that we don't need in the end. Second,
# it would be more elegant if we can perform the whole merge directly
# in SQL without having to remove some data using Pandas at the
# end. We will show how to solve these two problems next.
# The following code gives the same result, but the whole merging is
# done in SQL and it is more memory-friendly in case your dataset is
# large. The reason is that SQL has some internal optimizations it can
# perform when executing the query. Here it is important to keep in
# mind the order of execution of the SQL statements: 1) FROM ... LEFT
# JOIN ... ON ... (specifies which DataFrames to use and the kind of
# join to perform), 2) GROUP BY (specifies the grouping), 3) SELECT
# (specifies which columns to select; '* means all columns and
# 'MAX(cs.datadate)' takes the maximum value of 'datadate' from the
# 'cs' DataFrame in each group and discards all other observations in
# the group).
df2 = \
    pysqldf(
        'SELECT_*, MAX(cs.datadate)_' + \
        'FROM_crsp_LEFT_JOIN_cs_' + \
        'ON_crsp.TICKER=cs.tic_AND_' + \
        'crsp.date_BETWEEN_cs.datadate_AND_cs.datadate2_' + \
        'GROUP_BY_crsp.TICKER, _crsp.date')
df2['date'] = pd.to_datetime(df2['date'])
df2['datadate'] = pd.to_datetime(df2['datadate'])
df2.drop(['MAX(cs.datadate)', 'datadate2'], axis=1, inplace=True) # Remove columns.
```

```
df2 = df2.sort_values(by=['TICKER', 'date']).reset_index(drop=True) # Cosmetics.
# Inspect the data. It should be same as with 'df1'.
df2.tail()
# Check if both DataFrames are the same. They should be the same as
# both ways of merging should yield the same results.
df2.equals(df1)
```

B.5 Plotting with Pandas

The nice thing about Pandas is that you can often create plots very easily, without using many add-on packages (besides the standard Matplotlib). What Pandas does is to wrap Matplotlib so that it is straightforward to plot the data you have in a DataFrame. We have briefly discussed Matplotlib in Section 20.2 starting on page 148. Here, however, we focus more specifically on plotting with the Pandas wrapper around Matplotlib.

In the following code we take the data from CRSP, calculate cumulative stock returns, and plot them. The code below can be found in the file code-pandas-100-plotting.py on the course website and the data files we use are also on our course website.

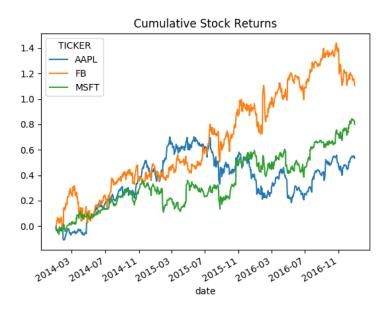
This script calculated cumulative stock returns and plots them.

```
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
# Here we import the data and select the columns we need.
df = pd.read_csv('data-CRSP-extract.csv')
df['date'] = pd.to_datetime(df['date']) # Convert string to 'datetime64'.
df = df[['TICKER', 'date', 'RET']] # Just ticker, date, and stock returns.
# Calculate the cumulative stock returns. They show by how much each
# stock price has gone up or down during our sample's time period.
df.RET = np.log(1 + df.RET) # Convert to log-returns so that we can sum them up.
df = df.sort_values(['TICKER', 'date']) # Ensure data is sorted before running 'cum
df['cum_RET'] = \
    df.\
    groupby('TICKER')['RET'].\
    apply(lambda x: x.cumsum()) # For each ticker, calculate cumulative sum.
df.cum_RET = np.exp(df.cum_RET) - 1 \# Convert log returns to linear returns.
del df['RET'] # Don't need raw returns any more for this application.
# Reshape the data to have each ticker in a separate column. The
# 'pivot()' method can save a lot of work if you compare it to
# reshaping the data yourself using for-loops! Keep in mind that here
# we specify 'date' as the index, which is useful for the subsequent
```

```
# plotting commands, as the date will automatically be put on the
# x-axis.
df = df.pivot(index='date', columns='TICKER', values='cum_RET')

# Plot the cumulative stock returns for all stocks (i.e. in this case,
# for all columns).
df.plot()
plt.title('Cumulative_Stock_Returns')
plt.show()
```

The code above should create the following plot, showing the cumulative returns of the stocks in our partial CRSP data extract. In case the lines are not easy to see in the printout, please take a look at the electronic version on the course website.



B.6 Downloading Equities Data from AlphaVantageInto Pandas

AlphaVantage is a data provider for financial market data. They provide the following data for free. All you need is to get a free API key from their websites.

- 1. Realtime and historical data in JSON and CSV format
- 2. Technical indicators
- 3. Bitcoin and other digital currencies

A such, it is a great alternative to Yahoo Finance or Google Finance, which had some problems recently. There are of course also other data providers out there who provide at least partially free data, for example

- 1. Yahoo Finance
- 2. Google Finance
- 3. AlphaVantage
- 4. Stoog
- 5. Quandl
- 6. BarChart
- 7. IEX

There are (at least) two ways to access AlphaVantage:

- 1. Since AlphaVantage provides the data as CSV (as well as JSON), you can directly import the relevant CSV files into Pandas. We will show such an example at the beginning of the following script.
- 2. Specific Python packages, such as alpha_vantage. Packages like these are usually just a thin wrapper around the AlphaVantage API. Their goal is to make it easier to get the data you want quickly. You can also directly specify the format you would like to data to receive in, e.g. JSON, CSV, or a Pandas DataFrame. As DataFrames are so practical, we will mostly focus on this data format.

The code below can be found in the file code-pandas-120-alphavantage.py on the course website.

```
# This script illustrates downloading financial data from
# AlphaVantage.
# Your AlphaVantage key. In order to download the data, you can get a
# key directly from the AlphaVantage website for free.
mykey = 'demo'
# Specify the stock ticker here. You can also use stock indices,
# e.g. IXIC (Nasdaq Composite), DJI (Dow Jones Industrial Average),
# INX (S&P 500) etc. You can also get data from stock markets around
# the world. For example, 'NSE:TITAN' gets the stock value of TITAN
# from the Indian Exchange NSE, or 'AI.PA' is for Air Liquide at the
# Paris stock exchange (PA).
myticker = 'MSFT'
import matplotlib.pyplot as plt
import pandas as pd
# Here we manually download some data from AlphaVantage, using only
# the Pandas package.
df = \setminus
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pd.read_csv(
        'https://www.alphavantage.co/query?' + \
        'function=TIME_SERIES_DAILY_ADJUSTED&' +\
        'symbol=' + myticker + \
        '&apikey=' + mykey + \
        '&datatype=csv')
# Take a look at the data we just downloaded. Keep in mind that
# AlphaVantage gives you the raw data sorted DESCENDINGLY based on the
# timestamp.
df.head()
df.sort_values(by='timestamp', inplace=True) # Sort by 'timestamp'.
df.reset_index(drop=True, inplace=True) # Reset the index to 0,1,2,3...
df.timestamp = pd.to_datetime(df.timestamp).dt.date # Convert to datetime and discar
\# Plot the stock price, adjusted for dividends and stock splits.
df.plot(x='timestamp', y='adjusted_close')
plt.title(myticker + '_Stock_Price')
plt.show()
# Calculate stock return.
df['L_close'] = df.adjusted_close.shift() # Lag the closing price.
df['RET'] = df.close / df.L_close - 1 # Calculate stock index return.
df.dropna(inplace=True)
                              # Drop missing data.
df.reset_index(drop=True, inplace=True) # Reset the index to 0,1,2,3...
# Cut returns into low (0), medium (1), and high (2) returns based on
# quantiles.
df['q_RET'] = pd.qcut(df.RET, 3, labels=False)
df[df.q_RET == 2].head()
                          # Show examples of high returns.
df[df.q_RET == 2].RET.mean() # Mean return of high quantile.
from pprint import pprint
                                # For pretty printing.
# Here we import a bunch of things from the 'alpha_vantage'
# package. Depending on what you want to download, you just need some
# of these imports.
from alpha_vantage.timeseries import TimeSeries
from alpha_vantage.techindicators import TechIndicators
from alpha_vantage.sectorperformance import SectorPerformances
from alpha_vantage.cryptocurrencies import CryptoCurrencies
from alpha_vantage.foreignexchange import ForeignExchange
# You can get the data in JSON format if you like. However, most of
# the time it is better to ask for a Pandas DataFrame as we do in the
# remaining examples.
ts = TimeSeries(key=mykey)
                               # In JSON by default.
```

```
data, meta_data = ts.get_daily_adjusted(symbol=myticker)
pprint(data)
# Here we request a Pandas DataFrame, which often is more convenient
# to work with.
ts = TimeSeries(key=mykey, output_format='pandas')
# Get object with data and another with the call's metadata. By
# default, it just returns a subset of the data. If you want the whole
# dataset, use 'outputsize='full'' as an additional function argument.
data, meta_data = ts.get_daily_adjusted(symbol=myticker)
# Plot the data.
data['5._adjusted_close'].plot()
plt.title('Times_Series_for_' + myticker)
plt.show()
# Bollinger Bands.
ti = TechIndicators(key=mykey, output_format='pandas')
data, meta_data = ti.get_bbands(symbol=myticker, time_period=20)
                                # Plot last year only.
data.tail(252).plot()
plt.title('BBbands_indicator_for_' + myticker)
plt.show()
# Sector performance.
sp = SectorPerformances(key=mykey, output_format='pandas')
data, meta_data = sp.get_sector()
data['Rank_G: Year_Performance'].plot(kind='bar')
plt.title('Sector_Performance_(Year)')
plt.tight_layout()
plt.grid()
plt.show()
# Crypto currencies.
cc = CryptoCurrencies(key=mykey, output_format='pandas')
data, meta_data = cc.get_digital_currency_daily(symbol='BTC', market='CNY')
data['4b._close_(USD)'].plot(logy=True)
plt.tight_layout()
plt.title('Daily_Value_for_Bitcoin_(BTC)')
plt.grid()
plt.show()
# Foreign exchange data is only available as JSON format (not in CSV
# or Pandas format). There's also no metadata in this call (so we just
# use a placeholder '_').
cc = ForeignExchange(key=mykey)
data, _ = cc.get_currency_exchange_rate(from_currency='EUR', to_currency='USD')
pprint(data)
```

The first part of the previous code prints the stock price from AlphaVantage similar to the following plot.

