

Course reader Basic Machine Learning for Bioinformatics 2021



Figure 1. Pretentious Machine Learning images on the covers of course readers, who can do without them? Taken from: <https://www.synaptica.com/machine-and-deep-learning/>

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Introduction

This course focusses on the very basics of Machine Learning (ML). This means that we'll implement linear regression (yes, that's ML), logistic regression (which, confusingly, is a classification method), a simple neural network, K-means clustering and hierarchical clustering, and Principal Component Analysis (PCA) ourselves using only basic functions and packages like Numpy and Matplotlib for plotting. Along the way we look at overfitting and underfitting and related bias and variance, look at how we measure model performance, and other things to keep in mind.

Machine Learning algorithms are implemented using Linear Algebra computations. Linear Algebra is a whole domain of mathematics with many interesting applications. If you are a studying bioinformatics, you have already come across its use in 2D and/or multi-D systems of ODEs. A proper course about Linear Algebra can take weeks, and we don't have that time. Therefore, for our purposes, Linear Algebra is just a nice way of manipulating numbers (such as multiplying the weights for each feature in a multivariate linear regression with the feature values for all training examples) in a form that's efficient for computation. The main ideas from Linear Algebra that are relevant for us will be introduced in a small crash course during the lectures, supplemented with suggestions for videos and other resources for further study.

We focus on the basics because it is important that you get a feel for what's really going on. There is an aura of magic around ML, and the excitement around AI has reached a fever pitch. True, advances like AlphaFold 2 and some of Google's and Facebook's convolutional neural nets are true marvels of engineering and a sight to behold. Yet at its core, we're just shuffling numbers around. That's why we work on implementing things ourselves: to get a feel for what is really happening.

It is likely that most of what you'll be doing during your studies and later on is applied ML: taking a problem from your domain of study, using an ML algorithm on it, and thereby solving said problem. Finding clusters (subtypes) in single cell cancer sequencing data, training a classifier to predict these different types of cancers based on biopsy sequencing data, that sort of thing. If you're going to train a neural network, you won't implement it yourself, you'll use Keras and TensorFlow. If you're making a Support Vector Machine, Random Forest, Linear Regressor, using PCA or a clustering of any kind, you're going to use Scikit-learn (or R equivalents). There are free, open-source packages for almost every popular ML algorithm, and you'd be foolish not to use them. For this reason, in the second week, we give a bird's eye view of Scikit-learn and Keras, so you'll have a basic familiarity and know enough to dive deeper yourself after the course.

The rest of the second week consists of a two-day ML project on Kaggle, where you're going to compete in groups to build the best classifier for a biological dataset using Scikit-learn, topped off with a short presentation about what you did. Finally, on the 23rd in the afternoon, there's a pen and paper exam that will probe your knowledge about what you've done.

Course structure

General

The first three days of week 1 cover supervised learning methods. The last two days of week 1 cover unsupervised learning methods. Week 2 switches from implementing the methods ourselves to using modern established ML libraries (Scikit-learn, Keras and TensorFlow) that you'll use for your applied ML projects. There's a two day project on Kaggle where you work in teams to get the best classification performance on a certain biological dataset and present your work afterwards. The pen and paper exam is on Thursday the 23rd in the afternoon.

Daily structure

We start at 9:00. Lectures and short practicals applying the lecture concepts are interleaved in the morning session. This means ~1 hour of lecture, followed by ~45-60 minutes of direct application in programming exercises, repeated about 2-3 times. We have lunch from ~12:15-13:00. The afternoon practical starts at 13:00 and runs until (at most) 17:00. This schedule is approximate: this is the first time the course is given so mismatches in actual timing versus planned timing are to be expected! Note also that sometimes there'll be 2 lectures rather than 3, or only 2 (somewhat longer) practicals.

Topics Week 1

- Day 1: Linear regression, gradient descent, introduction to linear algebra
- Day 2: Logistic regression, regularisation, ROC curve, introduction to neural networks (NNs)
- Day 3: NN Backpropagation algorithm, convolutional neural networks explained, guest speaker on deep learning in Oxford Nanopore sequencing (13:15-14:05).
- Day 4: K-means clustering, hierarchical clustering, deep dive into phylogenetics
- Day 5: Problems with high-dimensional data, Principal Component Analysis (PCA)

Topics Week 2

- Day 1: Working with scikit-learn, introduction to Keras and TensorFlow, project introduction and start
- Day 2: Working on the project
- Day 3: Working on the project, project presentation in the afternoon
- Day 4: Q&A session in the morning, exam in the afternoon

Before the course starts

Software requirements

Make sure you have a working installation of Python 3 (preferably the newest version, which is python 3.8 at the time of writing), and have installed [numpy](#), [matplotlib](#), [scikit-learn](#), [TensorFlow](#) (which automatically also installs [Keras](#)), [Biopython](#) (conda install -c conda-forge biopython) and [pandas-plink](#) (conda install -c conda-forge pandas-plink) in your working environment. Rather than manually setting this up, please download [Anaconda](#). It comes with many of the most-used packages pre-installed, a graphical user interface to manage packages, and many other tools. You might want to make an environment for the course specifically, but that is optional. Environments allow you to keep projects separate: rather than installing packages globally, you install them only for a specific environment, in this case the environment you make for this course. See [here](#) if you want to do that. **You do need to manually install Biopython and pandas-plink (see commands above).**

Downloading practical materials and slides

All practical material, slides, and this reader are on GitHub [here](#). Clone the git repository locally such that you can work with the files. **Do this by Sunday evening 12-12-2021 at the earliest: I will still update the materials this weekend!** If you do not know how to do this, do the following:

1. If on Windows, install [Git for Windows](#); if on Mac, install [Git for Mac](#); if on Linux (Ubuntu distro), run `sudo apt update`, followed by `sudo apt-get install git`. Follow all instructions for your respective case.
2. Go to the folder where you want the files to be downloaded. In Windows, you can open Git Bash there by right clicking (Figure 2). This opens a console. In all cases, you can type `git clone`, followed by <https://github.com/DieStok/Basic-Machine-Learning-for-Bioinformatics.git> in the console in the directory where you want the files. This will automatically download everything there.

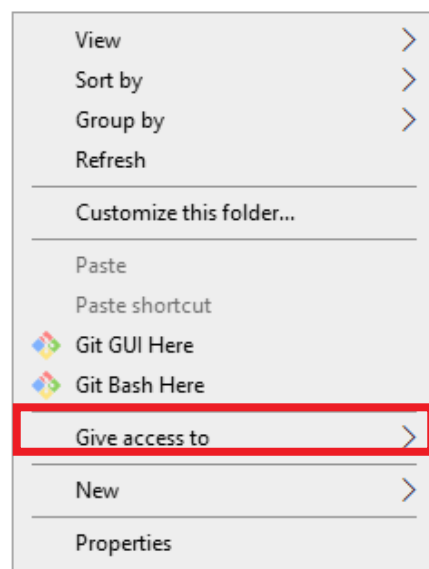


Figure 2. Opening Git Bash in a folder by right clicking in Windows.

3. Alternatively, if this doesn't work, you can download a .zip of all files manually, by clicking on the area indicated by the red rectangle in Figure 3.

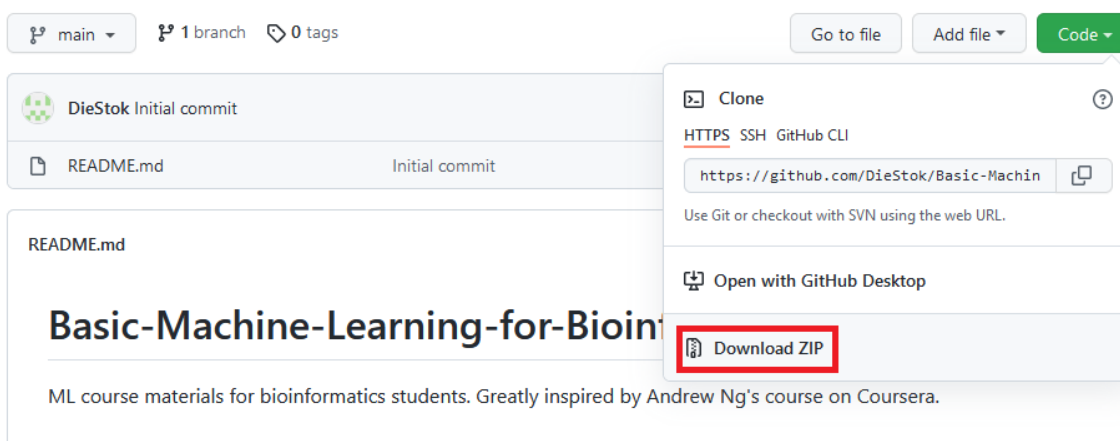


Figure 3. Manually downloading the files from the GitHub page.

Doing the practicals

Normal method

The practicals all take place in Jupyter Notebooks, which are similar to R notebooks or R markdown files you might know. Read more on Jupyter Notebooks [here](#). All you need to do to access the practicals is:

1. Open an Anaconda Prompt (search for Anaconda Prompt in your start menu)
2. (**optional**) activate the environment for the course (if using)
3. Navigate to where you cloned the GitHub repository. For me, that's "J:\OnderwijsmedewerkerUU2021-2022\Basic-Machine-Learning-for-Bioinformatics". To change drive in Windows, simply type the drive letter with a colon (*J:* in my example). Then use *cd OnderwijsmedewerkerUU2021-2022\Basic-Machine-Learning-for-Bioinformatics*.
4. Finally, type *jupyter notebook* in the anaconda prompt. A web browser should open up, where you can navigate to the practical you need. Take care not to close the prompt in which you are running jupyter notebook.

If you didn't manage to install Anaconda

First come to us and let us troubleshoot the issues together. Perhaps there are quick fixes. If we cannot get things working on your local machine, do the following:

1. Make sure you have a Google account
2. Go to <https://colab.research.google.com/>
3. Open a new notebook (File>new notebook)
4. In that new notebook, write in your code cell:

```
from google.colab import drive
drive.mount('/content/drive')
```

and then run it (ctrl+enter with the code cell selected). This will ask you to authenticate that this notebook can access your drive. Allow this.
5. Insert a new code cell below (shortcut keys: esc+b). In it, write:

```
import os
os.chdir("/content/drive/MyDrive/")
```

run this code cell too.
6. Insert a new code cell below. There, clone the course Github. You do that with the following command: `!git clone https://github.com/DieStok/Basic-Machine-Learning-for-Bioinformatics.git`

7. You can't open the notebooks you download there directly from Colab, unfortunately. You need to go to your google drive in the browser, locate the appropriate notebook there, and then right click and open with Colab. See Figure 4.

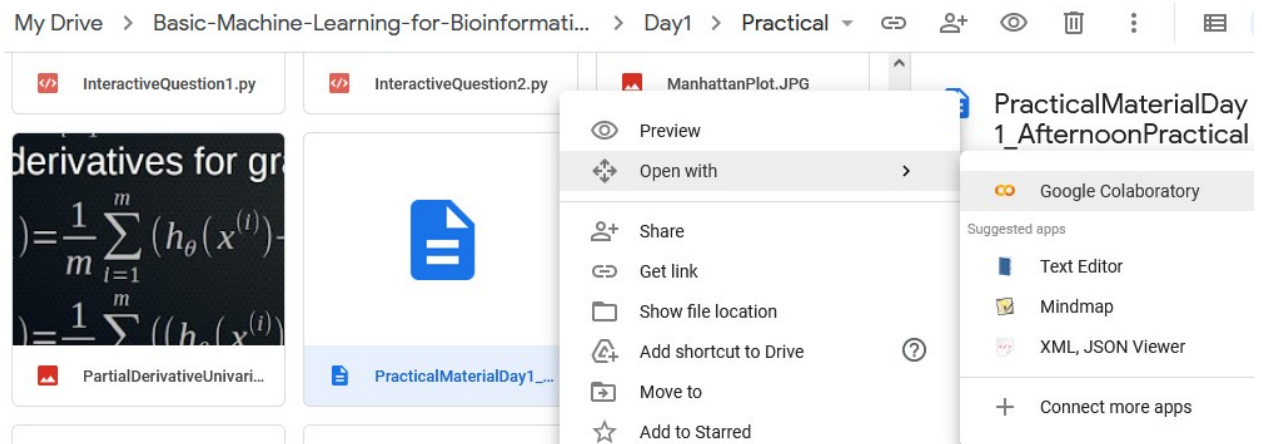


Figure 4. Opening a Jupyter Notebook from Google Drive. Navigate to your Drive in the browser; right-click, and select open with Colab.

8. **Before you can run the notebook you need to do four things:**

-If it says somewhere in the notebook `%matplotlib notebook`, change this to `%matplotlib inline`. Otherwise plotting won't show anything.

-Again mount your google drive:

```
from google.colab import drive
drive.mount('/content/drive')
```

-insert the same os code as above, but now navigating to the correct practical directory, say for practical one:

```
import os
os.chdir("/content/drive/MyDrive/Basic-Machine-Learning-for-
Bioinformatics/Day1/Practical")
```

-Insert calls to install any dependencies that are not yet there. These use the package manager pip. So for when we are using pandas-plink:

```
!pip install pandas-plink
```

You can also use:

```
!pip list
```

 to see what's installed. You put these in a code cell and execute. You will know soon enough when you are missing a dependency, because you will see errors.

Day 1 materials

Description

Today we focus on the most basic supervised ML algorithm: linear regression. We talk about the equations for linear regression, its cost function, using gradient descent to minimise the cost function, going from univariate linear regression (one feature, x , one predicted quantity based on it, y) to multivariate linear regression (predicting, say, disease severity based on levels of 10 biomarkers in the blood). We also discuss the bias-variance trade-off and using cross-validation for measuring ML performance. Finally, you are introduced to linear algebra for programming ML algorithms efficiently.

Recommended videos for review:

[Video 1](#); [Video 2](#); [Video 3](#); [Video 4](#) (~55 minutes)

Extra resources

Videos

- Linear Regression short StatQuest: <https://www.youtube.com/watch?v=PaFPbb66DxQ>
- More in-depth StatQuest Linear Regression: https://www.youtube.com/watch?v=nk2CQITm_eo
- Cross-validation: <https://www.youtube.com/watch?v=fSytzGwwBVw>
- Hyperparameters: <https://www.youtube.com/watch?v=VTE2KlfoO3Q>
- Bias and variance (Video 2 above): <https://www.youtube.com/watch?v=EuBBz3bI-aA>
- Linear algebra: https://www.youtube.com/watch?v=fNk_zzaMoSs&list=PLZHQObOWTQDPD3MizzM2xVFitgF8hE_ab (3Blue1Brown playlist)

Reading

- Cross-validation: <https://machinelearningmastery.com/k-fold-cross-validation/>
- Hyperparameters: <https://machinelearningmastery.com/difference-between-a-parameter-and-a-hyperparameter/>
- In-depth treatment of bias-variance trade-off: <http://scott.fortmann-roe.com/docs/BiasVariance.html>
- Linear algebra: <https://cbmm.mit.edu/sites/default/files/documents/algebra.pdf> (slide deck with a lot of depth) ; https://minireference.com/static/tutorials/linear_algebra_in_4_pages.pdf
- Math symbols primer: <https://amitnesh.com/2019/08/math-for-programmers/>
- L2-norm: <https://kawahara.ca/what-does-the-l2-or-euclidean-norm-mean/>

- Jupyter (Notebook) tutorial:
https://www.tutorialspoint.com/jupyter/jupyter_notebook_introduction.htm

Other

- Bias-variance trade-off and overfitting discussed with interactive examples:
https://machinelearningcompass.com/model_optimization/bias_and_variance/
- Jupyter Guide to Linear Algebra: https://bvanderlei.github.io/jupyter-guide-to-linear-algebra/Matrix_Algebra.html
- Linear algebra course Khan Academy: <https://www.khanacademy.org/math/linear-algebra>

Day 2 materials

Description

Today we focus on a simple ML algorithm for classification: predicting discrete classes from data (as opposed to continuous variables, which we covered yesterday). This is logistic regression, which you've probably heard of before. It will turn out that the implementation is extremely similar to that of linear regression, just with a different cost function. After this, we will look into regularisation to prevent overfitting, and get started on the basics of neural networks.

Recommended videos for review:

[Video 1](#); [Video 2](#); [Video 3](#); [Video 4](#) (~60 minutes)

Extra resources

Videos

- Regularisation: <https://www.youtube.com/watch?v=KvtGD37Rm5I> ; <https://www.youtube.com/watch?v=Q81RR3yKn30>
- Neural networks: <https://www.youtube.com/watch?v=HGwBXDKFk9I&list=PLblh5JKOoLUIxGDQs4LFFD--41Vzf-ME1> (StatQuest playlist)
- (AUC) ROC: <https://www.youtube.com/watch?v=4jRBRDbJemM>
- Nested cross-validation: <https://www.youtube.com/watch?v=LpOxxBeggM0>; <https://www.youtube.com/watch?v=DuDtXtKNpZs>

Reading

- Logistic regression: <https://towardsdatascience.com/logistic-regression-detailed-overview-46c4da4303bc>
- Regularisation: <https://explained.ai/regularization/index.html> (highly recommended for a *deep* understanding)
- Neural networks: <http://neuralnetworksanddeeplearning.com/>
- (AUC) ROC and (AUC) PRC: <https://glassboxmedicine.com/2019/02/23/measuring-performance-auc-auROC/> ; <https://glassboxmedicine.com/2019/03/02/measuring-performance-auprc/> ; <https://machinelearningmastery.com/roc-curves-and-precision-recall-curves-for-classification-in-python/>

Other

Day 3 materials

Description

Today we focus on neural networks in a bit more depth. Specifically, we take a look at backpropagation and how it allows us to optimise the weights and biases of a neural network. You'll be implementing backpropagation yourself and training a simple network. However, simple (or dense) neural networks are not what have made the huge strides in image recognition (and some other areas) in the last ~10 years possible. For that, we need to look to convolutional neural networks. We'll discuss what convolution is, and the core ideas that make these networks tick. You'll see how you can train a simple convolutional neural net quickly and easily in Keras.

Recommended videos for review:

[Video 1](#); [Video 2](#); [Video 3](#) (~50 minutes)

Extra resources

Videos

- Whole Neural Network playlist StatQuest: <https://www.youtube.com/watch?v=CqOfi41LfDw&list=PLblh5JKOoLUIxGDQs4LFFD--41Vzf-ME1>
- Backpropagation: <https://www.youtube.com/watch?v=tIeHLnjs5U8> ; https://www.youtube.com/watch?v=8d6jf7s6_Qs&t=0s ; <https://www.youtube.com/watch?v=GlcnxUIrtek&list=PLiaHhY2iBX9hdHaRr6b7XevZtgZRa1PoU&index=4>
- Numerical gradient checking: <https://www.youtube.com/watch?v=pHMzNW8Agq4&list=PLiaHhY2iBX9hdHaRr6b7XevZtgZRa1PoU&index=5>
- Batch normalisation: <https://youtu.be/DtEq44FTPM4?t=127>
- Dropout: <https://www.youtube.com/watch?v=ARq74QuavAo> ; <https://www.youtube.com/watch?v=vAVOY8frLIQ> (quite in-depth)

Reading

- Backpropagation: <http://neuralnetworksanddeeplearning.com/chap2.html> ; <https://mattmazur.com/2015/03/17/a-step-by-step-backpropagation-example/>
- Neural networks in general: <http://neuralnetworksanddeeplearning.com>
- Great 4-part series of Medium articles on neural networks and the (linear algebra) math behind optimising them: [1](#), [2](#), [3](#), [4](#).
- SENet architecture (complicated!): <https://github.com/hujie-frank/SENet>
- Understanding convolutions intuitively: <https://towardsdatascience.com/intuitively-understanding-convolutions-for-deep-learning-1f6f42faee1>
- Batch normalisation: <https://towardsdatascience.com/batch-normalisation-explained-5f4bd9de5feb>

- Dropout: <https://towardsdatascience.com/simplified-math-behind-dropout-in-deep-learning-6d50f3f47275>
- Learning logical functions with neurons illustrated (until halfway through subheading 4): [Solving XOR with a single Perceptron | by Lucas Araújo | Medium](#)

Other

- Interactive TensorFlow Neural Network: <http://playground.tensorflow.org/>
- **Highly recommended** backpropagation slide deck: http://cs231n.stanford.edu/slides/2019/cs231n_2019_lecture04.pdf (especially slide 42-55).

Day 4 materials

Description

Today we change gears completely and switch from supervised to unsupervised learning. In unsupervised learning, there are no labels, no known training data on which to train our algorithms. Instead, we want to find some structure in the data, without knowing what we are looking for: clustering. This means we must contend with the fact that there is no ‘correct’ clustering, and that depending on your exact criterion for clustering, there may be many possible clusters to make. We’ll look at prototype clustering (K-means as an example) and hierarchical clustering. We’ll then take a slightly in-depth look at how we use clustering in phylogeny and what extra problems sequences have in that respect.

Recommended videos for review:

[Video 1](#); [Video 2](#); [Video 3](#); [Video 4](#) (~45 minutes)

Extra resources

Videos

- Clustering algorithms overview: https://www.youtube.com/watch?v=Se28XHI2_xE
- K-means maths: <https://www.youtube.com/watch?v=Z2VXrwMYHFk> ;
<https://www.youtube.com/watch?v=0MQEt10e4NM>
- Lecture series hierarchical clustering: https://www.youtube.com/watch?v=GVz6Y8r5AkY&list=PLBv09BD7ez_7qIbBhyQDr-LAKWUeycZtx

Reading

- Clustering algorithms overview: <https://towardsdatascience.com/the-5-clustering-algorithms-data-scientists-need-to-know-a36d136ef68>
- K-means clustering: <https://towardsdatascience.com/k-means-clustering-algorithm-applications-evaluation-methods-and-drawbacks-aa03e644b48a>
- Hierarchical clustering:
https://www.tutorialspoint.com/machine_learning_with_python/machine_learning_with_python_clustering_algorithms_hierarchical.htm
- Multiple sequence alignment technique overview: <http://www.cs.cmu.edu/~durand/03-711/2009/Lectures/MSA09-2-notes.pdf>

Other

Day 5 materials

Description

Today we continue on with unsupervised learning, now focussing on dimensionality reduction and its flagship/most well-known method: Principal Component Analysis. We first take a look at the curse of dimensionality: how more dimensions cause problems by exponentially increasing the need for training data, inducing overfitting, and making it impossible to use distance metrics. Luckily, we have intelligent ways to reduce the dimensionality of data. Using StatQuest's excellent basis, we'll dive a bit deeper into performing PCA ourselves, and the mathematics behind it. Finally, with knowledge of PCA and multivariate linear regression in hand, we know enough to perform our very own GWAS, which we will do in the afternoon practical.

Recommended videos for review:

[Video 1](#) ; [Video 2](#) (28 minutes)

Extra resources

Videos

- Counter-intuitive phenomena in high dimensions: <https://www.youtube.com/watch?v=dr2sIoD7eeU>
- Worked example of PCA calculations: <https://www.youtube.com/watch?v=S51bTyIwxFs>
- PCA lecture series: https://www.youtube.com/watch?v=IbE0tbjy6JQ&list=PLBv09BD7ez_5_yapAg86Od6JeypkS4YM
- PCA in GWAS: <https://youtu.be/qMkuYrlh7jw?t=370>
- t-SNE and UMAP: <https://www.youtube.com/watch?v=NEaUSP4YerM> ; <https://www.youtube.com/watch?v=6BP181wGGP8>
- Eigenvalue and eigenvector calculations: <https://www.youtube.com/watch?v=TQvxWaQnrqI>
- Why eigenvalues correspond to variance on each component:

Reading

- Curse of dimensionality: <https://www.visiondummy.com/2014/04/curse-dimensionality-affect-classification/>
- PCA in GWAS: <https://pubmed.ncbi.nlm.nih.gov/16862161/> ; <https://stats.stackexchange.com/a/8780>
- PCA stackoverflow gems: <https://stats.stackexchange.com/questions/217995/what-is-an-intuitive-explanation-for-how-pca-turns-from-a-geometric-problem-wit?noredirect=1&lq=1> (mathy) ; <https://stats.stackexchange.com/questions/117695/why-is-the-eigenvector-in-pca-taken-to-be-unit-norm> (unit length vector) ; <https://stats.stackexchange.com/questions/2691/making-sense-of-principal-component->

[analysis-eigenvectors-eigenvalues/140579#140579](#) (simple explanation from the level you'd tell your grandmother to more of an expert)

- In-depth PCA tutorial with examples:
http://www.cs.otago.ac.nz/cosc453/student_tutorials/principal_components.pdf
(unfortunately some images are missing but other than that it is a great read!)

Other

- Interactive exploration of the UMAP technique: <https://pair-code.github.io/understanding-umap/>

Day 6 materials

Description

Today we switch gears again. You've now been duly introduced to the basics of ML: supervised and unsupervised learning, (nested) cross-validation, bias and variance, and implementation of various classifiers yourself. There are great open-source libraries that vastly speed up Machine Learning workflows and do all the heavy lifting for you. We didn't start with those because it is important to know what's really going on, but when applying ML in future projects you'd be foolish to do everything by yourself. Therefore, today we switch resolutely to Scikit-learn and Keras, and also discuss some other best practices. In the afternoon, you will be introduced to the project and start work on that. At 16:00 there is a UBC seminar on ML in brain that is sure to be interesting!

Recommended videos for review:

[Video 1](#) ; [Video 2 \(you can probably watch on 1.5 speed\)](#); [Video 3](#) (~5 hours 10 minutes: You don't need to watch all of this! The latter two videos are full courses on scikit-learn and Keras which you can browse as you see fit.)

Extra resources

Videos

- Scikit-learn pipelines tutorial (with code): <https://www.youtube.com/watch?v=XvnkUg1yVmk> ; <https://www.youtube.com/watch?v=9vz0n1cyMbc>
- Freecodecamp 2+-hour Keras introduction: <https://www.youtube.com/watch?v=qFJeN9V1ZsI>
- What a tensor is: <https://www.youtube.com/watch?v=f5liqUk0ZTw>
- Optimizers (gradient descent, stochastic gradient descent, mini-batch gradient descent, AdaGrad, Adam): <https://www.youtube.com/watch?v=mdKjMPmcWjY>
- Pandas: <https://www.youtube.com/watch?v=vmEHCJofslg>

Reading

- Hyperparameter optimisation strategies (GridSearchCV and/or RandomSearchCV): <https://machinelearningmastery.com/hyperparameter-optimization-with-random-search-and-grid-search/> ; https://scikit-learn.org/stable/modules/grid_search.html
- One-hot encoding <https://machinelearningmastery.com/why-one-hot-encode-data-in-machine-learning/>
- Nested cross-validation: <https://weina.me/nested-cross-validation/> ; https://scikit-learn.org/stable/auto_examples/model_selection/plot_nested_cross_validation_iris.html#sphx-glr-auto-examples-model-selection-plot-nested-cross-validation-iris-py ; <https://stats.stackexchange.com/questions/65128/nested-cross-validation-for-model-selection>

- Global average pooling: <https://adventuresinmachinelearning.com/global-average-pooling-convolutional-neural-networks/> ; <https://paperswithcode.com/method/global-average-pooling>
- Keras training loop from scratch: https://keras.io/guides/writing_a_training_loop_from_scratch/
- RMSprop (explanation and Python implementation): <https://machinelearningmastery.com/gradient-descent-with-rmsprop-from-scratch/>
- Adam (standard optimisation method deep learning): <https://machinelearningmastery.com/adam-optimization-algorithm-for-deep-learning/>
- Mini-batch gradient descent: <https://towardsdatascience.com/batch-mini-batch-stochastic-gradient-descent-7a62ecba642a>
- Tuning models with KerasTuner: https://keras.io/guides/keras_tuner/getting_started/
- Gradient descent total overview: <https://runder.io/optimizing-gradient-descent/>
- Detailed hyperparameter optimisation guide: <https://nanonets.com/blog/hyperparameter-optimization/>

Other

- Adam gradient optimizer slide deck: https://moodle2.cs.huji.ac.il/nu15/pluginfile.php/316969/mod_resource/content/1/adam_pres.pdf
- Different optimizers visualised (very cool, though does not translate 1-on-1 to the real high-dimensional case): <https://github.com/Jaewan-Yun/optimizer-visualization>

Day 7 + 8 materials

Description

These days will be spent on the group project. **Details on the group project will be given at the beginning of week 2.** On Wednesday(day 8), we will have 10-minute group presentations from 15:00-17:00. They should showcase your approach, snags you ran into and how you did or did not solve them, and what you learned. Grading of the project will be mostly on the basis of how well you implemented the basics: did you correctly do the guided exercises, did you correctly cross-validate, did you correctly normalise and correct missing values, etc. A small component will be your creativity and use of advanced methods (if any), and another small component how clear your presentation was. Be sure to look at least as professional and pleased with yourself when presenting as our relative below (Figure 5).



Figure 5: Taken from: <http://media.gettyimages.com/photos/male-chimpanzee-in-business-clothes-picture-id184941527>

Day 9 materials

Description

Today we have a question hour in the morning (probably from 11:00-12:00). Questions should be sent in by e-mail before 10:30. **If no questions are sent in, I will communicate this to you and there will be no question hour.** Other than that, there's the pen-and-paper exam from 14:00-17:00. After that, you are free! You could spend your time wisely, or you could read [Saturday Morning Breakfast Cereal](#). I recommend the latter.

Extra Resources

Basic

- Simple overview of steps of Machine Learning: <https://www.youtube.com/watch?v=nKW8Ndu7Mjw> + <https://towardsdatascience.com/the-7-steps-of-machine-learning-2877d7e5548e>
- Numpy video tutorial: <https://www.youtube.com/watch?v=GB9ByFAIAH4>
- Pandas video tutorial: <https://www.youtube.com/watch?v=vmEHCJofslg>

Libraries

- eli5 (from explain like I'm five) for making ML models interpretable: <https://eli5.readthedocs.io/en/latest/overview.html#>
- imbalanced-learn for imbalanced datasets: <https://imbalanced-learn.org/stable/>

Words of gratitude

I am indebted to Dr. Jeroen de Ridder for expert support in preparing the lectures and practical material. The teachings in the course follow [Andrew Ng's Coursera course](#) in much of the material, which I highly recommend for further learning and consolidation of the material presented here. The PCA section is based mostly on [the lecture series by Victor Lavrenko](#). I highly recommend both this lecture series and the other content on this channel for further viewing. [StatQuest](#) is invaluable for intuitive and simple explanations of the ideas behind various ML methods, stripped of their mathy mystery. I'd like to thank the countless volunteers who make and maintain open-source libraries like Matplotlib, Keras, and Scikit-learn and prevent the world from falling apart ([relevant XKCD](#)). Finally, I'd like to thank Prof. Dr. Berend Snel and Dr. Bas van Breukelen for giving me the opportunity to make this course (and/or cobble it together based on many different sources!).

Stock photos of a woman threatening a goldfish with a gun

What course reader can finish without them? There's a whole slew of these. Amazing. That's life for ya. No ML algorithm can hope to explain why this is the case. Note: I am against harming goldfish, but have reason to believe this particular specimen was photoshopped in.

