CSI5180. Machine Learning for Bioinformatics Applications

Deep learning: fundamentals

,

Marcel Turcotte

Preamble

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Preamble

Deep learning: fundamentals

In this lecture, we study artificial neural networks (ANN) and specifically the multilayer architectures known as deep learning. This is the first of three lectures on this topic. Herein, we focus on the building blocks, namely the units, their connectivity, and the training algorithms.

General objective :

▶ Discuss the similarities and differences between other machine learning algorithms and deep learning.

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Learning objectives

- **Explain** in your own words the threshold logic unit
- Discuss the role of the activation function
- Describe the multilayer perceptron

Reading:

- James Zou, Mikael Huss, Abubakar Abid, Pejman Mohammadi, Ali Torkamani, and Amalio Telenti, A primer on deep learning in genomics, *Nat Genet* **51**:1, 1218, 2019.
- Webb, S. Deep Learning for Biology. *Nature* **554**, 555557 2018.

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Reading

- ▶ Vanessa Isabell Jurtz, Alexander Rosenberg Johansen, Morten Nielsen, Jose Juan Almagro Armenteros, Henrik Nielsen, Casper Kaae Sønderby, Ole Winther, and Søren Kaae Sønderby, An introduction to deep learning on biological sequence data: examples and solutions, *Bioinformatics* 33:22, 36853690, 2017.
- Mufti Mahmud, Mohammed Shamim Kaiser, Amir Hussain, and Stefano Vassanelli, Applications of deep learning and reinforcement learning to biological data, *IEEE Transactions on Neural Networks and Learning Systems* **29**, 20632079, 2018.

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Reading

- Seonwoo Min, Byunghan Lee, and Sungroh Yoon, Deep learning in bioinformatics, *Brief Bioinform* **18**:5, 851869, 2017.
- Gökcen Eraslan, Ziga Avsec, Julien Gagneur, and Fabian J Theis, Deep learning: new computational modelling techniques for genomics, *Nat Rev Genet* **20**:7, 389403, 2019.
- ▶ Binhua Tang, Zixiang Pan, Kang Yin, and Asif Khateeb, Recent advances of deep learning in bioinformatics and computational biology, Frontiers in Genetics 10, 214, 2019.

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Plan

- 1. Preamble
- 2. Application
- 3. Introduction
- 4. Implementations
- **5.** Prologue

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3Blue1Brown on deep learning (videos)

But what is a Neural Network?

https://youtu.be/aircAruvnKk

19 minutes

Gradient descent, how neural networks learn

https://youtu.be/IHZwWFHWa-w

21 minutes

What is backpropagation really doing?

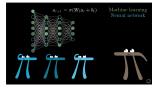
https://youtu.be/Ilg3gGewQ5U

14 minutes

Backpropagation calculus

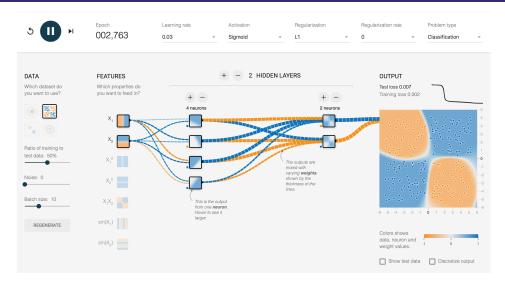
https://youtu.be/tIeHLnjs5U8

10 minutes



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playground.tensorflow.org

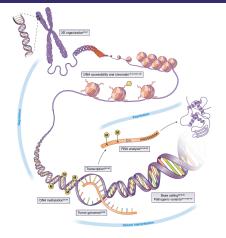


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Application

Application 10/46

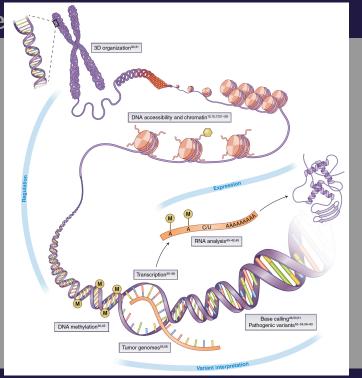
What are the applications?



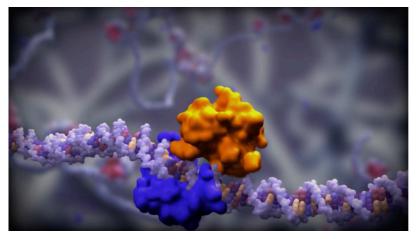
Source [Zou et al., 2019] Figure 2

Application 11/46

What are



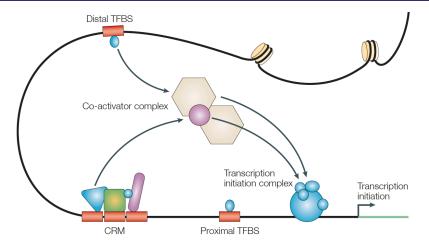
Transcription factors



Source: https://youtu.be/MkUgkDLp2iE

Application 12/46

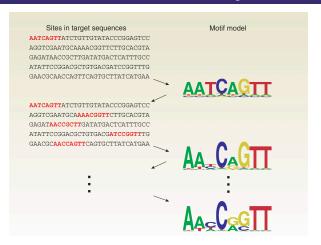
Essential Cellular Biology: Gene Regulation



Wasserman, W. & Sandelin, A. Applied bioinformatics for the identification of regulatory elements. *Nat Rev Genet* **5**, 276287 (2004).

Application 13/46

DNA sequence motif discovery



Patrik Dhaeseleer, How does DNA sequence motif discovery work?, *Nat Biotechnol* **24**:8, 95961, 2006.

Application 14/46

Deep Learning in Genomics Primer

- James Zou, Mikael Huss, Abubakar Abid, Pejman Mohammadi, Ali Torkamani, and Amalio Telenti, A primer on deep learning in genomics, *Nat Genet* **51**:1, 1218, 2019.
 - Google Colab Notebook

Application 15/46

Introduction

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Artificial neural networks

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Introduction 17/46

Artificial neural networks

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- Likewise, artificial neural networks (ANN) were inspired by the human brain.

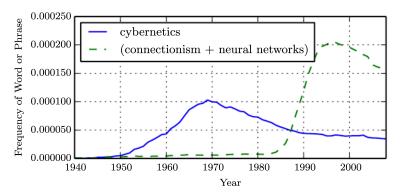
Introduction 17/46

Artificial neural networks

- "Although planes were inspired by birds, they don't have to flap their wings." [Géron, 2019]
- Likewise, artificial neural networks (ANN) were inspired by the human brain.
 - However, ANN do not (necessarily) mimic the brain or explain its functioning.

Introduction 17/46

1940-1960, 1980-mid 1990, 2010-



Source: [Goodfellow et al., 2016] Figure 1.7

 McCulloch, W. S. & Pitts, W. A logical calculus of the ideas immanent in nervous activity. The Bulletin of Mathematical Biophysics 5, 115133 (1943).

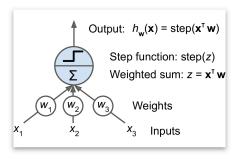
Introduction 18/46

What has changed?

- Vast amounts of data are now available
- Powerful computers (including fast GPU)
- Improved **training algorithms**

Introduction 19/46

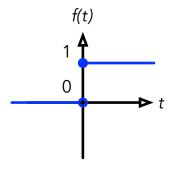
Neuron - threshold logic unit

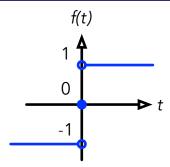


Source: [Géron, 2019] Figure 10.4

Common **step functions** include the **heavyside** function (0 if the input is negative and 1 otherwise) or the **sign** function (-1 if the input is negative, 0 if the input is zero, 1 otherwise).

Simple step functions - heavyside and sign





$$heavyside(t) =$$

- ightharpoonup 1, if t > 0
- 0, if t < 0

$$sign(t) =$$

- 1, if t > 0
- 0, if t = 0
- -1, if t = 0

Does this sound familiar?

• One **threshold logic unit** (TLU) is similar to a **logistic regression** and both are solving the same kinds of problems.

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- Just like the Linear Regression, the Logistic Regression computes a weighted sum of the input features:

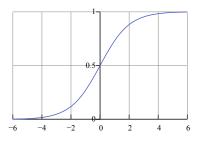
$$\theta_0 + \theta_1 x_i^{(1)} + \theta_2 x_i^{(2)} + \ldots + \theta_D x_i^{(D)}$$

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▶ The image of this function is $-\infty$ to ∞ !

In mathematics, a **standard logistic function** maps a real value (\mathbb{R}) to the interval (0,1):



Source: Wikipedia

$$\sigma(t) = \frac{1}{1 + e^{-t}}$$

The **Logistic Regression** model, in its vectorized form is:

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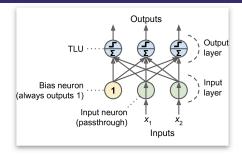
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- $y_i = 0$, if $h_{\theta}(x_i) < 0.5$
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- The values of θ are learnt using **gradient descent**.

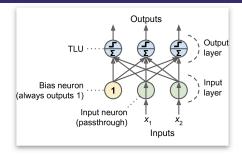
Perceptron



Source: [Géron, 2019] Figure 10.5

A **Perceptron** consists of a single layer of threshold logic units.

Perceptron



Source: [Géron, 2019] Figure 10.5

- A **Perceptron** consists of a single layer of threshold logic units.
- It computes the following function:

$$h_{W,b}(X) = \phi(WX + b)$$

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- **b: bias vector** (same size as the number of neurons in the output layer).
- Activation function: maps its input domain to a restricted set of values (heavyside and sign are commonly used with threshold logic unit perceptrons).

sklearn.linear_model.Perceptron

```
from sklearn.linear_model import Perceptron
# ...
model = Perceptron()
model.fit(X, y)

y_pred = model.predict(X_new)
```

Standard logistic (sigmoid) function,

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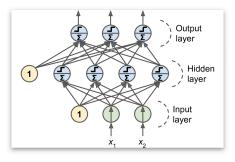
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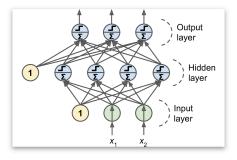
Rectified Linear Unit function (ReLU),

$$ReLU(z) = max(0, z)$$



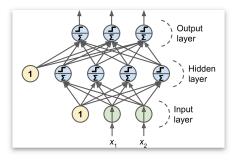
Source: [Géron, 2019] Figure 10.7

One input layer



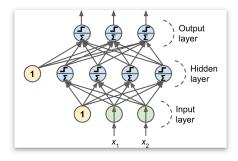
Source: [Géron, 2019] Figure 10.7

- One input layer
- One or more hidden layers



Source: [Géron, 2019] Figure 10.7

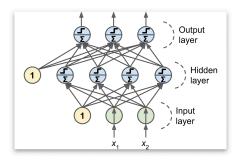
- One input layer
- One or more hidden layers
- One **output layer**



Source: [Géron, 2019] Figure 10.7

- One input layer
- One or more hidden layers
- One output layer
- With the exception of the output layer, every layer has a bias unit

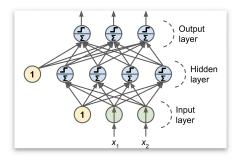
Feed-forward network (FFN)



Source: [Géron, 2019] Figure 10.7

Since information flows from the input to the output, this architecture is called a feed-forward network (FFN)

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- Since information flows from the input to the output, this architecture is called a feed-forward network (FFN)
- Generally, FFN with more than three (3) layers are called **deep learning networks**.

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- With these activation functions the network is calculating a non-linear function.

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Repeat the gradient descent until the network converges to a solution.

Backpropagation (2)

Going through all the examples for every iteration would be too slow.

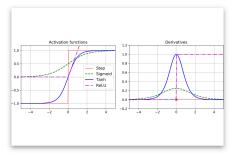
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 - Rather, examples are grouped together into **mini-batches** (32).
 - Going through all the examples is called an **epoch**.

Backpropagation (3)



Source: [Géron, 2019] Figure 10.8

Regularization

Early stopping

Regularization

- **Early stopping**
- L1 and L2 norm

Regularization

- **Early** stopping
- L1 and L2 norm
- Dropouts

Implementations

Implementations 37/4

Frameworks Widely popular implementations include,

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Implementations 38/46

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Reading:

- Ladislav Rampasek and Anna Goldenberg, TensorFlow: Biologys gateway to deep learning?, *Cell Syst* **2**:, no. 1, 124, (2016).
- ➤ Kathleen M Chen, Evan M Cofer, Jian Zhou, and Olga G Troyanskaya, Selene: a PyTorch-based deep learning library for sequence data, *Nat Methods* **16**:4, 315318, (2019).
- Avsec, Z. et al. Kipoi: accelerating the community exchange and reuse of predictive models for genomics. bioRxiv 375345 (2018). doi:10.1101/375345

Implementations 38/4

Suggested exercises

- 1. Review the Google Colab Notebook by [Zou et al., 2019]
- 2. Experiment with https://playground.tensorflow.org

Implementations 39/4

Prologue

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- In a **Perceptron**, the units of the output layer are connected to **all** the input units, forming a **dense layer**.
- The **multilayer perceptron** has intermediate layers called **hidden layers**.
- Activation functions are playing a key role, allowing the network to compute a non-linear function of its input. This is important to model complex phenomenon.

Next module

Deep learning (continued)

References

- Burkov, A. (2019).

 The Hundred-Page Machine Learning Book.

 Andriv Burkov.
- Chen, K. M., Cofer, E. M., Zhou, J., and Troyanskaya, O. G. (2019). Selene: a PyTorch-based deep learning library for sequence data. *Nat Methods*, 16(4):315–318.
- Chollet, F. (2017).

 Deep learning with Python.

 Manning Publications.
- D'haeseleer, P. (2006). How does DNA sequence motif discovery work? Nat Biotechnol, 24(8):959–61.
- Eraslan, G., Avsec, Ž., Gagneur, J., and Theis, F. J. (2019). Deep learning: new computational modelling techniques for genomics. *Nat Rev Genet*, 20(7):389–403.

References

Hands-on Machine Learning with Scikit-Learn, Keras, and TensorFlow. O'Reilly Media, 2nd edition.

Goodfellow, I., Bengio, Y., and Courville, A. (2016). Deep Learning.

MIT Press.

Jurtz, V. I., Johansen, A. R., Nielsen, M., Almagro Armenteros, J. J., Nielsen, H., Sønderby, C. K., Winther, O., and Sønderby, S. K. (2017).

An introduction to deep learning on biological sequence data: examples and solutions.

Bioinformatics, 33(22):3685-3690.

- Mahmud, M., Kaiser, M. S., Hussain, A., and Vassanelli, S. (2018). Applications of deep learning and reinforcement learning to biological data. *IEEE Transactions on Neural Networks and Learning Systems*, 29:2063–2079.
- Min, S., Lee, B., and Yoon, S. (2017). Deep learning in bioinformatics. *Brief Bioinform*, 18(5):851–869.

References

- Rampasek, L. and Goldenberg, A. (2016). TensorFlow: Biology's gateway to deep learning? *Cell Syst*, 2(1):12–4.
- Tang, B., Pan, Z., Yin, K., and Khateeb, A. (2019).

 Recent advances of deep learning in bioinformatics and computational biology.

 Frontiers in Genetics, 10:214.
- Webb, S. (2018).

 Deep learning for biology.

 Nature, 554(7693):555–557.
- Zou, J., Huss, M., Abid, A., Mohammadi, P., Torkamani, A., and Telenti, A. (2019).

A primer on deep learning in genomics.

Nat Genet, 51(1):12-18.



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