

CS 66: Machine Learning

Prof. Sara Mathieson

Spring 2019



Outline for April 12

- Proposal and Project highlights
- Finish Convolutional Neural Networks
 - Weights on CONV layers
 - Parameter analysis
 - Strides and pooling
- Other NN architectures
- Next week: unsupervised learning
 - Lab 7 due MONDAY
 - Final project details posted
 - Office hours TODAY 1-3pm

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Project Proposal

- 1) Dataset
- 2) Methods/Algorithms
- 3) Scientific Question
- 4) Evaluation and Interpretation of Results
- 5) References

Datasets

- [Kaggle](#): Wide variety of datasets (may need to create an account).
- [UCI Machine Learning Repository](#): Also contains a wide variety of datasets (options on the left allow you to search by task, attribute type, etc which can be very useful).
- [ImageNET](#): Large database of images (larger than CIFAR-10, which is also an option).
- **Climate data**: The faculty at Swarthmore have been encouraged to incorporate climate analysis into our curriculum. If you are interested, I would encourage you to choose a climate-oriented project. There are a number of government and climate research sites with data. Here are a few:
 - [Climate.gov](#)
 - [GlobalChange.gov](#)
 - [NOAA](#)
 - [EPA](#)
- [Wikipedia Data List](#): Up-to-date list of datasets organized by category (may or may not be freely available):
 - Image data
 - Text data
 - Sound data
 - Signal data
 - Physical data
 - Biological data
 - Anomaly data
 - Question answering data
 - Multivariate data
- [1000 genomes \(human DNA data\)](#): If you're interested in exploring a biological project, let me know. This specific dataset contains DNA data from humans around the world, but there are many other datasets from other species.
- [The 50 Best Free Datasets for Machine Learning](#): There are a number of these type of lists floating around - this one looks decent but make sure that you can actually download the data.

Project Lab Notebook

- As you as you receive your git repo, start creating a “lab notebook” in your README
- This should say who was working, what date, how long, and briefly what you did

Sara: 03-07-18 (2hrs)

- now averaging the Markov chain, fixed all the results
- combined ancestral 1000 genomes still running (need to start similar for SGDP)
- started new runs with filtering to only have selected alleles in the “selected pop” and only have ancestral alleles in the “reference panel”

Project Deliverables

- Main deliverable: presentation

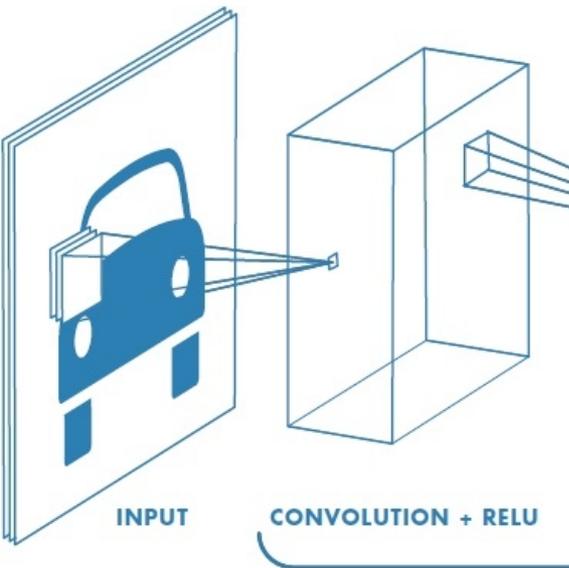
- Group of 1: 5 min
- Group of 2: 9 min
- Group of 3: 12 min

- On git:
 - Lab Notebook
 - Project Code
 - Presentation Slides

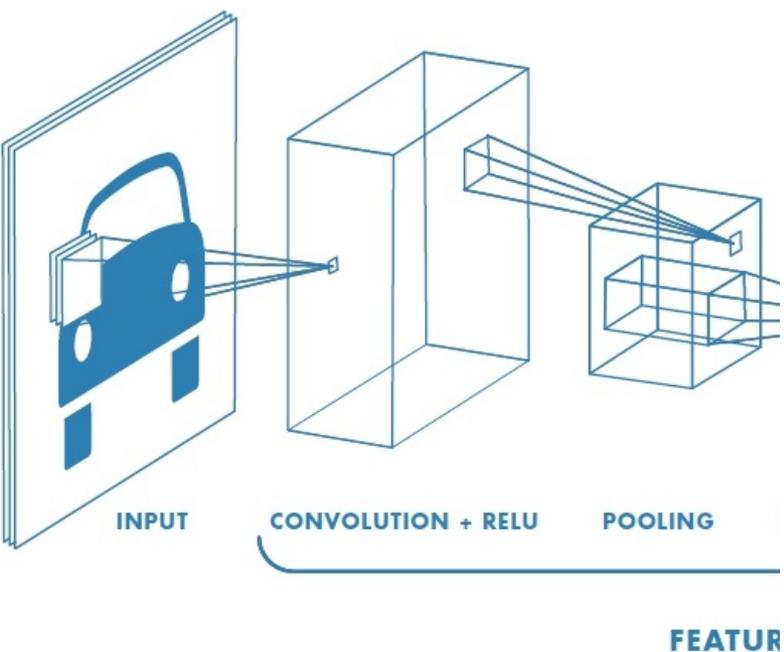
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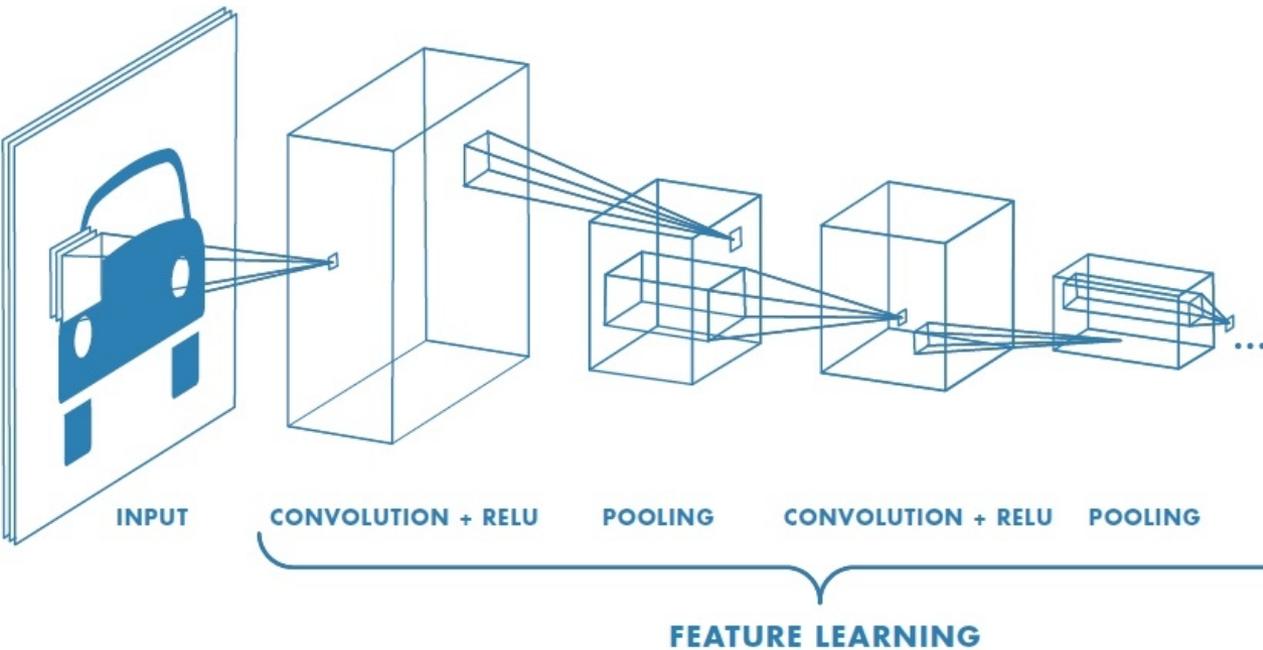
Visualization of an entire network



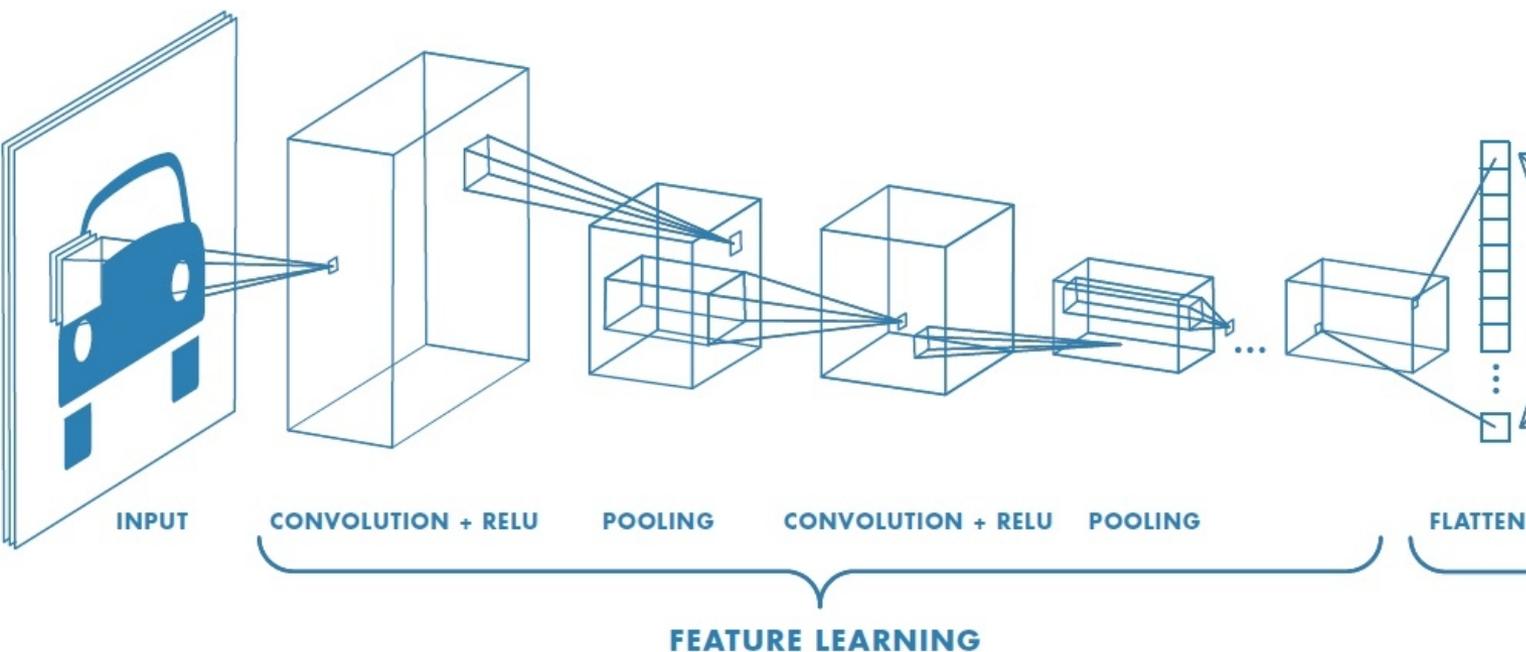
Visualization of an entire network



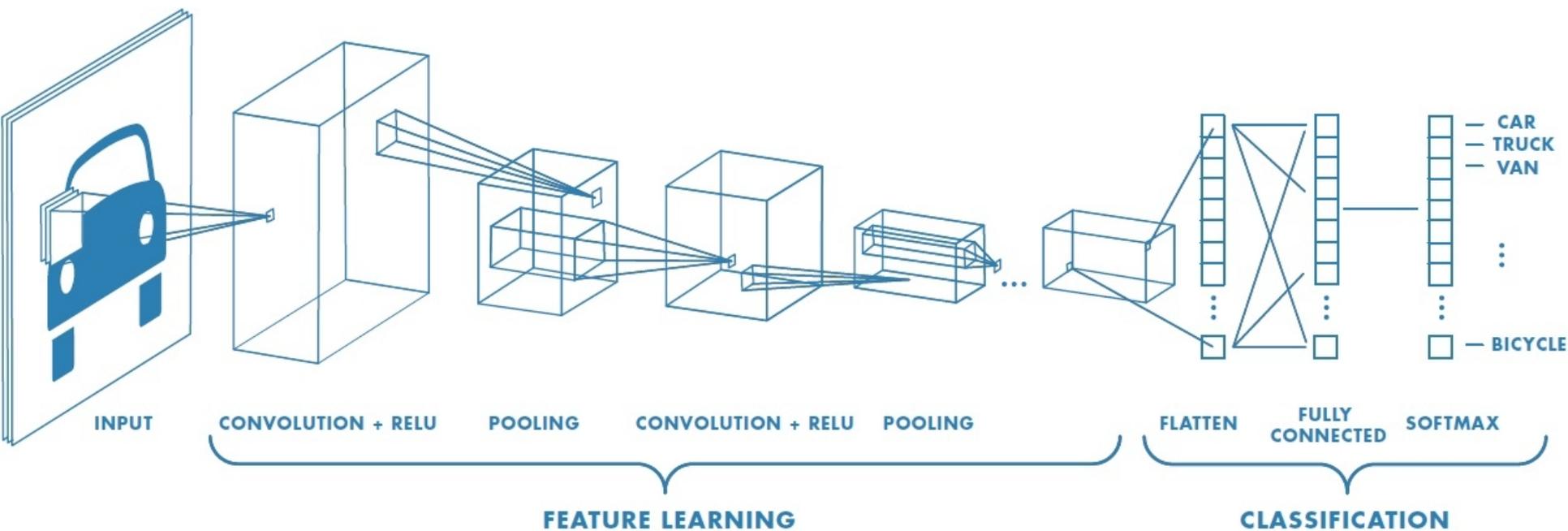
Visualization of an entire network



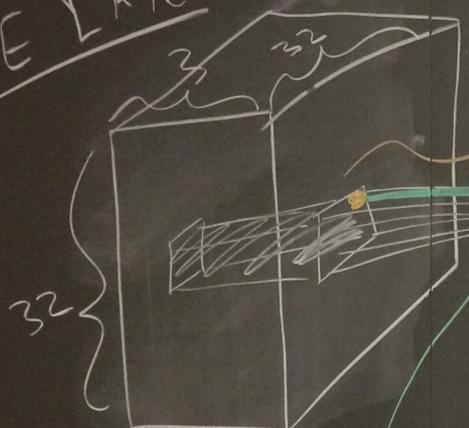
Visualization of an entire network



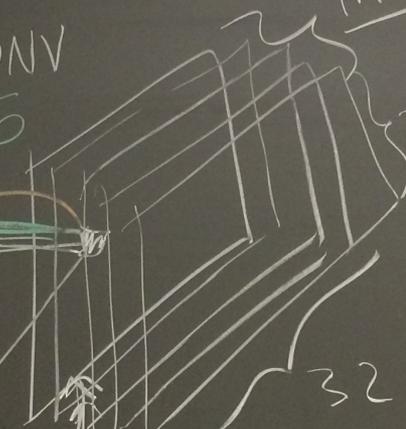
Visualization of an entire network



ONE LAYER

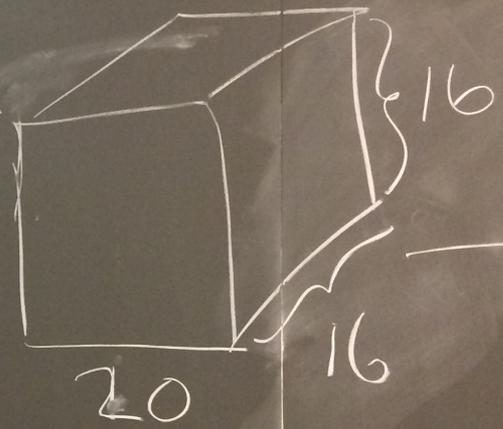


CONV
3x5



independent filters

RELU
POOL



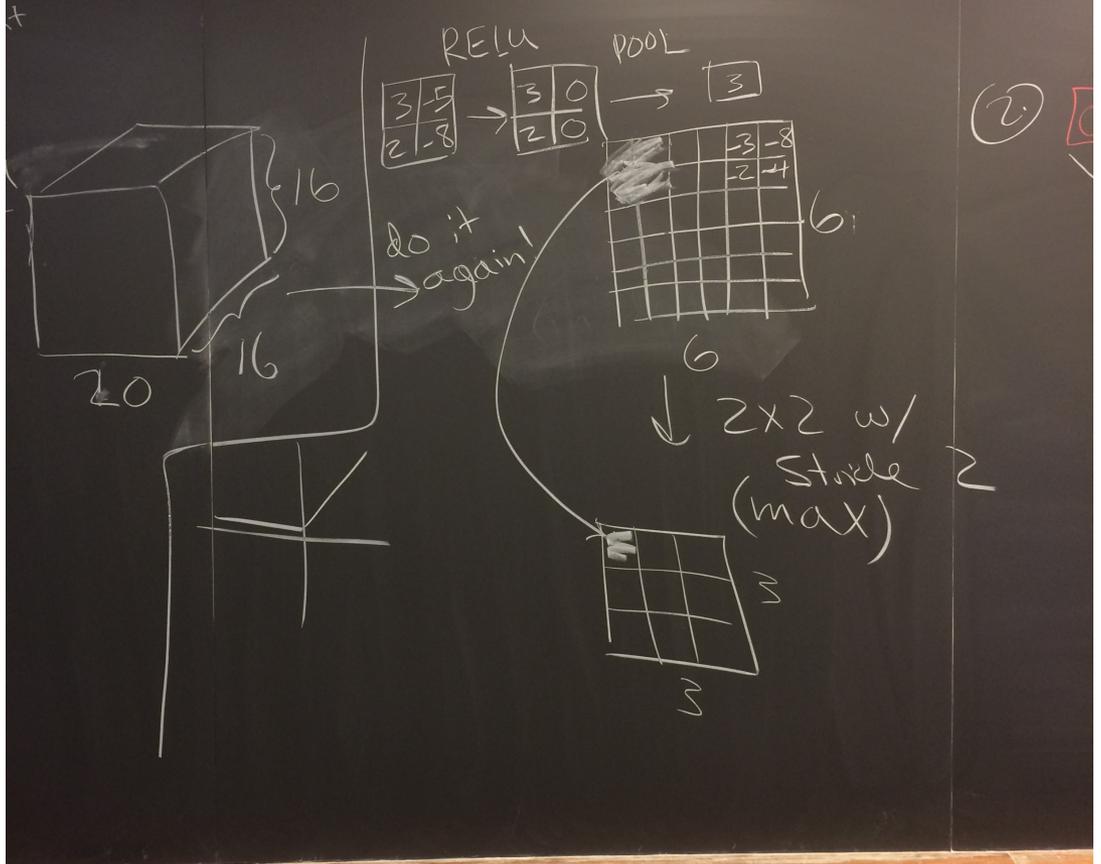
+ bias

random to start

output of
one filter

filters
(20)





Handout 16

(a) params:

CONV, FC

no params:

RELU, POOL, FLATTEN

(b) $5 \times$
weights for one filter

(c) $3 \times$

Note that some of these operations require hyper-parameters!

- CONV: filter size, number of filters, stride, padding
- POOL: filter size, stride
- FC: if using a hidden layer, number of units in this layer

(b) $5 \times 5 \times 3 \times 20 + 20$

weights for one filter → $5 \times 5 \times 3$ (one 3D filter)

↑ # filters new depth! → 20 (from input)

$+ 20$

1520

(c) $3 \times 3 \times 20 \times 10 + 10$

1810

(d) $8 \cdot 8 \cdot 10 \times 10 + 10$ (f)

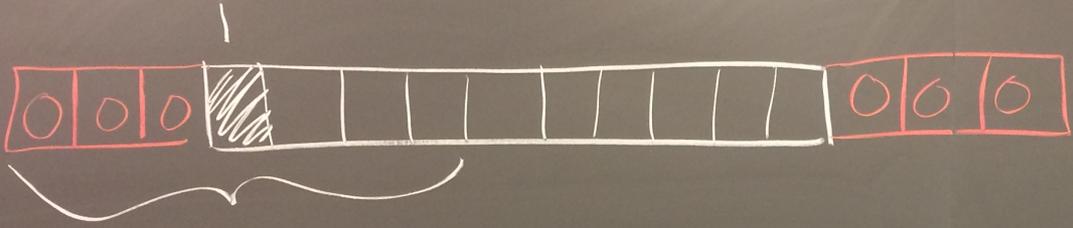
6410

(e) **9740**

* much better!



②



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