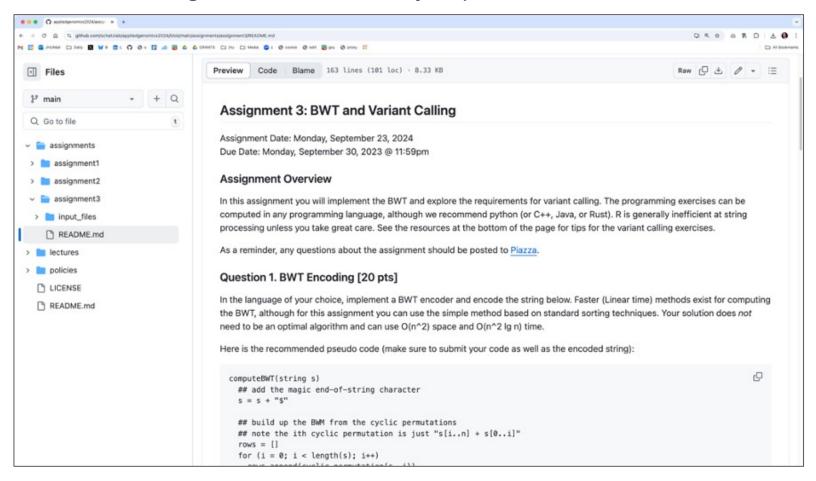
# ImageNet Classification with Deep Convolutional Neural Networks

Michael Schatz Oct 2, 2024

JHU EN.601.449/EN.601.649: Applied Comparative Genomics

#### Assignment 3 Due Monday September 30



Recap

#### "AlexNet"

#### ImageNet Classification with Deep Convolutional Neural Networks

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#### Abstract

We trained a large, deep convolutional neural network to classify the 1.2 million high-resolution images in the ImageNet LSVRC-2010 contest into the 1000 different classes. On the test data, we achieved top-1 and top-5 error rates of 37.5% and 17.0% which is considerably better than the previous state-of-the-art. The neural network, which has 60 million parameters and 650,000 neurons, consists of five convolutional layers, some of which are followed by max-pooling layers, and three fully-connected layers with a final 1000-way softmax. To make training faster, we used non-saturating neurons and a very efficient GPU implementation of the convolution operation. To reduce overfitting in the fully-connected layers we employed a recently-developed regularization method called "dropout" that proved to be very effective. We also entered a variant of this model in the ILSVRC-2012 competition and achieved a winning top-5 test error rate of 15.3%, compared to 26.2% achieved by the second-best entry.



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#### What is this?



#### GTX 580 GPU Network Architecture (annotated) only has 3GB! Network Architecture. normalizatio My iPhone's GPU has 6GB max-pooling GP41 11×11 5+5 448 3x2x 256 192 192 (48) (128) (192) (192) 2043 (123) Input image > Prediction! 224 + 274+ 1000 3432 545 3×3× 2041 11211 192 ×48 ¥3 142 (128) Saftmax (192) (192) (128) GPU2 (48) July- connected layers convolutional layers # neurons: 253,440 136,624 64,896 64,896 63,264 4096 1000

https://blog.acolyer.org/2016/04/20/imagenet-classification-with-deep-convolutional-neural-networks/

#### Kernels

Operation	Kernel ω	Image result g(x,y)
Identity	$\left[\begin{array}{ccc} 0 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{array}\right]$	6
Ridge or edge detection	$\begin{bmatrix} 0 & -1 & 0 \\ -1 & 4 & -1 \\ 0 & -1 & 0 \end{bmatrix}$	
Sharpen	$\begin{bmatrix} -1 & 8 & -1 \\ -1 & -1 & -1 \end{bmatrix}$ $\begin{bmatrix} 0 & -1 & 0 \\ -1 & 5 & -1 \\ 0 & -1 & 0 \end{bmatrix}$	
Bex blur (normalized)	$\frac{1}{9} \left[ \begin{array}{ccc} 1 & 1 & 1 \\ 1 & 1 & 1 \\ 1 & 1 & 1 \end{array} \right]$	e
Gaussian blur 3 x 3 (approximation)	$\frac{1}{16} \left[ \begin{array}{ccc} 1 & 2 & 1 \\ 2 & 4 & 2 \\ 1 & 2 & 1 \end{array} \right]$	C
Gaussian blur 5 x 5 (approximation)	$\begin{array}{c} 1 \\ \frac{1}{256} \begin{bmatrix} 1 & 4 & 6 & 4 & 1 \\ 4 & 16 & 24 & 16 & 4 \\ 6 & 24 & 36 & 24 & 6 \\ 4 & 16 & 24 & 16 & 4 \\ 1 & 4 & 6 & 4 & 1 \end{bmatrix} \end{array}$	9
Unsharp masking 5 x 5 Based on Gaussian blur with amount as 1 and threshold as 0 (with no image mask)	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	4



= input!A1 \* filter!\$A\$1 + input!A2 \* filter!\$A\$2 + input!A3 \* filter!\$A\$3 + input!A4 \* filter!\$A\$4 + input!A5 \* filter!\$A\$5 + input!B1 \* filter!\$B\$1 + input!B2 \* filter!\$B\$2 + input!B3 \* filter!\$B\$3 + input!B4 \* filter!\$B\$4 + input!B5 \* filter!\$B\$5 +

input!C1 \* filter!\$C\$1 + input!C2 \* filter!\$C\$2 + input!C3 \* filter!\$C\$3 + input!C4 \* filter!\$C\$4 + input!C5 \* filter!\$C\$5 + input!D1 \* filter!\$D\$1 + input!D2 \* filter!\$D\$2 + input!D3 \* filter!\$D\$3 + input!D4 \* filter!\$D\$4 + input!D5 \* filter!\$D\$5 + input!D5 \* filter!\$

input!E1 \* filter!\$E\$1 + input!E2 \* filter!\$E\$2 + input!E3 \* filter!\$E\$3 + input!E4 \* filter!\$E\$4 + input!E5 \* filter!\$E\$5

#### **Learned Convolutional Kernels**

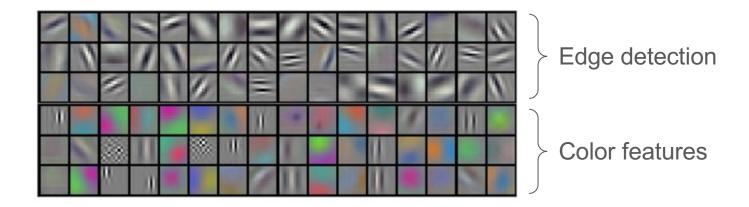
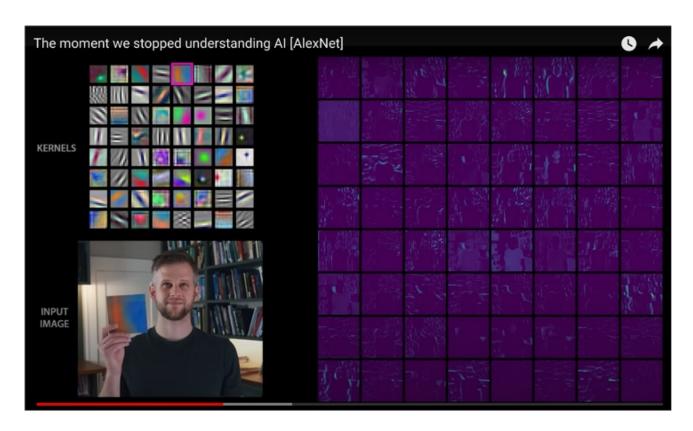
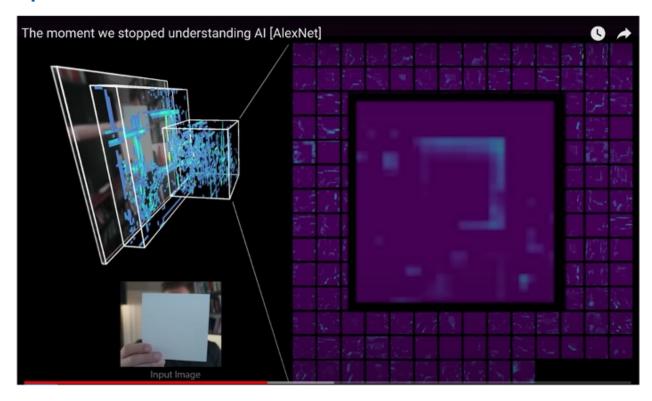


Figure 3: 96 convolutional kernels of size  $11 \times 11 \times 3$  learned by the first convolutional layer on the  $224 \times 224 \times 3$  input images. The top 48 kernels were learned on GPU 1 while the bottom 48 kernels were learned on GPU 2. See Section 6.1 for details.

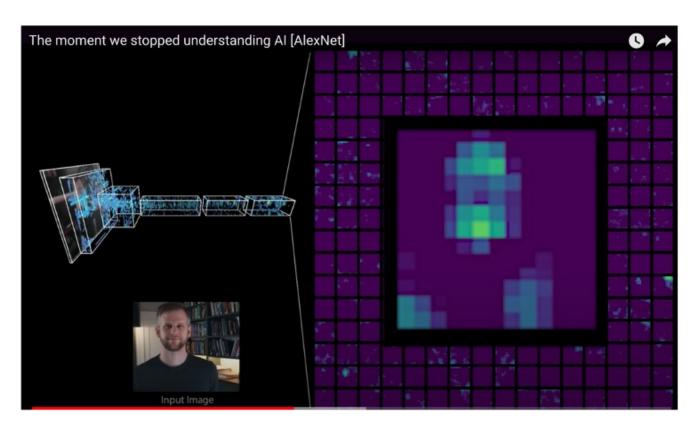
## From 1 image to 96 images: Activation Maps



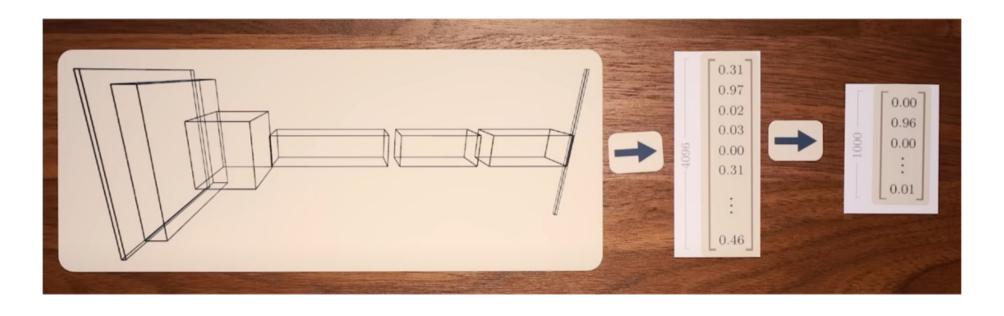
## Testing images shows the deeper layers are learning more complicated features



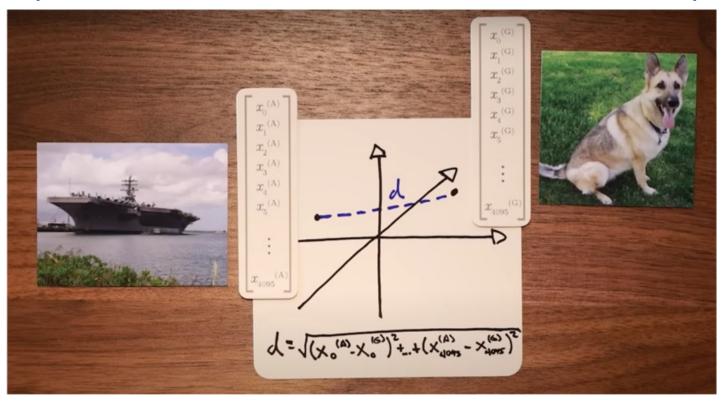
### Deeper layers recognize very complex features: faces!



## Final two layers are special: Last layer [1x1000]: Probability of the image in class i Second last [1x4096]: Projection of image into 4096-dim space

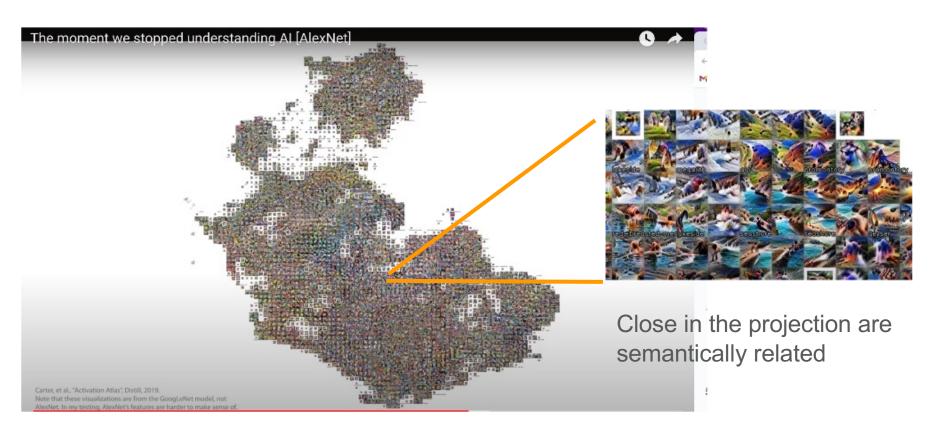


## Each image creates a unique vector Compute the distance between vectors in 4096-dim space

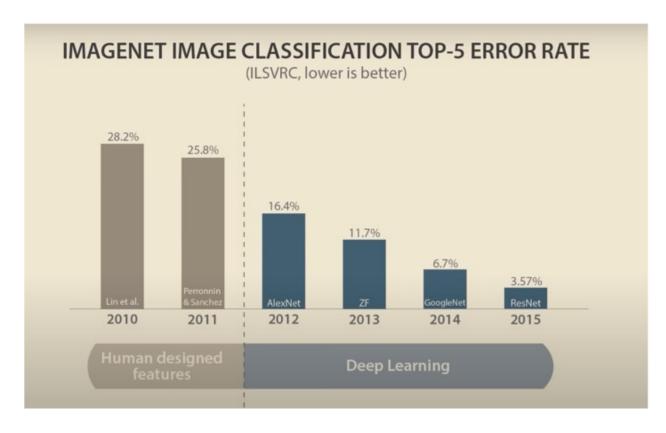


https://www.youtube.com/watch?v=UZDiGooFs54

#### Activation Atlas: tSNE plot of images embeddings



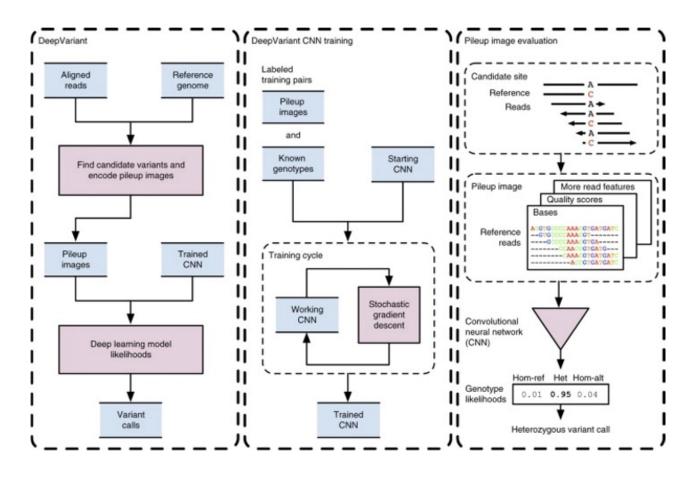
#### **Deep Learning Success!**



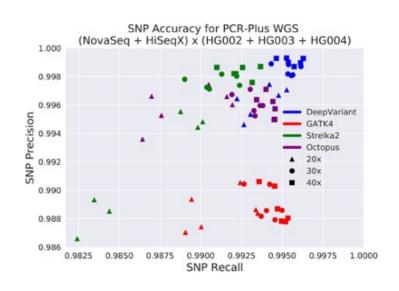
"Our results show that a large, deep convolutional neural network is capable of achieving record-breaking results on a highly challenging dataset using purely supervised learning. It is notable that our network's performance degrades if a single convolutional layer is removed. For example, removing any of the middle layers results in a loss of about 2% for the top-1 performance of the network. So the depth really is important for achieving our results."

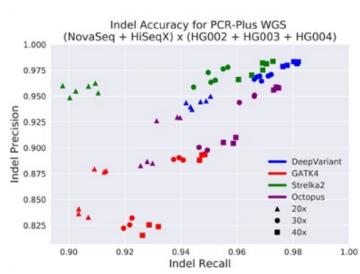
#### Outline

- 1. The Problem
- 2. pre-AlexNet approaches
- 3. How do animals & humans approach this?
- 4. Artificial Neural Networks
- 5. Convolutional Neural Networks
- 6. AlexNet
- 7. Impact



A universal SNP and small-indel variant caller using deep neural networks Poplin et al (2018) Nature Biotechnology. https://doi.org/10.1038/nbt.4235









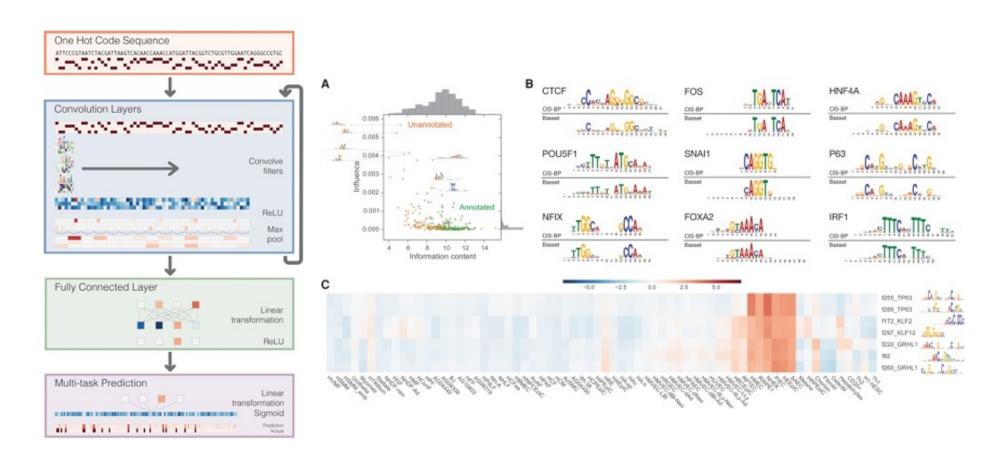




## An Extensive Sequence Dataset of Gold-Standard Samples for Benchmarking and Development

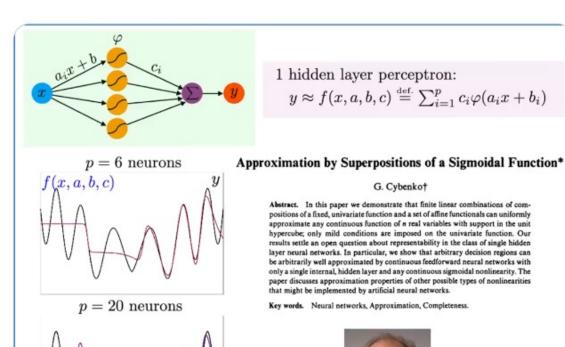
Baid et al (2020) bioRxiv

https://www.biorxiv.org/content/10.1101/2020.12.11.422022v1.full



Basset: learning the regulatory code of the accessible genome with deep convolutional neural networks Kelly et al (2016) Genome Research. doi: 10.1101/gr.200535.115

#### ANNs are "Universal Approximators"



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224\*224 = 50k pixels

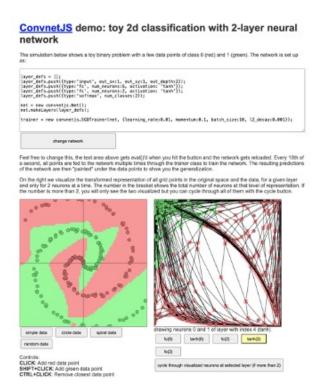
50k pixels \* 3 bytes (RGB) 838B possible inputs

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https://cs.stanford.edu/people/karpathy/convnetjs/demo/classify2d.html

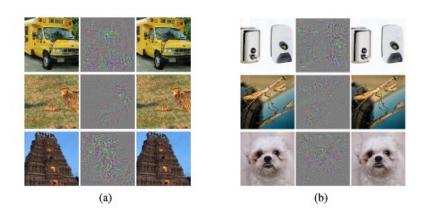


Figure 5: Adversarial examples generated for AlexNet [9].(Left) is a correctly predicted sample, (center) difference between correct image, and image predicted incorrectly magnified by 10x (values shifted by 128 and clamped), (right) adversarial example. All images in the right column are predicted to be an "ostrich, Struthio camelus". Average distortion based on 64 examples is 0.006508. Plase refer to http://goo.gl/huaGPb for full resolution images. The examples are strictly randomly chosen. There is not any postselection involved.

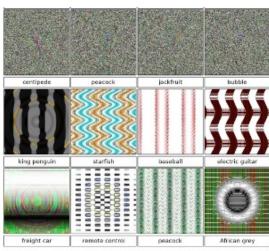


Figure 1. Evolved images that are unrecognizable to humans, but that state-of-the-art DNNs trained on ImageNet believe with ≥ 99.6% certainty to be a familiar object. This result highlights differences between how DNNs and humans recognize objects. Images are either directly (top) or indirectly (bottom) encoded.

#### Intriguing properties of neural networks

Szegedy et al (2013) arXiv:1312.6199

Deep Neural Networks are Easily Fooled: High Confidence Predictions for Unrecognizable Images

Nguyen et al (2015) arXiv:1412.1897

#### Reflections

- CNNs are incredibly powerful for image recognition (and similar tasks)
  - Heavily inspired by how the human brain processes visual information from the real world
  - Start with the most basic feature kernels and then iterate into higher level concepts leveraging their spatial proximity
  - Ultimately projects images into abstract semantic embeddings
- Certain problems are a great fit for CNNs:
  - ImageNet: Lots of test data, nicely annotated into 1000 categories
  - DeepVariant/Clairvoyante: Lots of test data; One network architecture made it the worlds best variant caller for all types of long read data: PacBio, ONT, 10x
- Deeper network architectures generally perform better
  - Largely constrained by GPU RAM & Cores

#### "Images" in Bioinformatics

- DNA sequence: one-hot encode
  - Are there other encoding schemes? (modDotPlot)
  - Predict features (genes, repeat classes); Predict species (kraken)

#### Coverage tracks:

- Predict: SNVs (DeepVariant), CNVs, SVs (Cue)
- RNAseq, ChIPseq (overcome error, allele specific)

#### Functional Genomics

- Inputs
  - Expression from multiple tissues (GTEx) from one patient
  - Multiomics: Variants + expression + methylation
- Predict phenotype: Gender (easy), cell types (easy), age (pretty easy)
  - Phenopredict: <a href="https://academic.oup.com/nar/article/46/9/e54/4920847">https://academic.oup.com/nar/article/46/9/e54/4920847</a>
- Disease outcome: Needs the right training data, right tissues
- Multiple alignments, position weight matrix
  - Predict conserved regions, predict binding sites, etc



https://www.youtube.com/watch?v=UZDiGooFs54