

# **CNN-Fold: Protein Fold Recognition by Deep Convolutional Neural Networks**

Tyler Banks

Presented to:

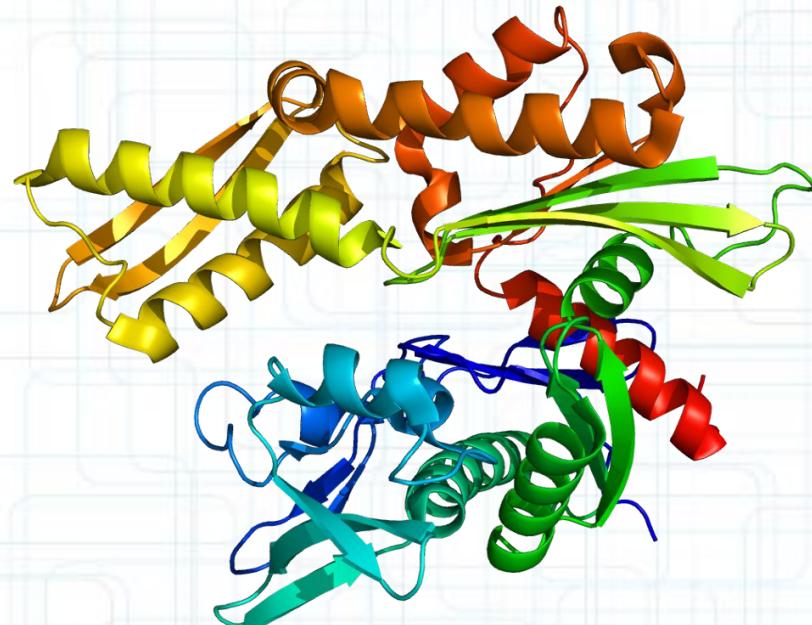
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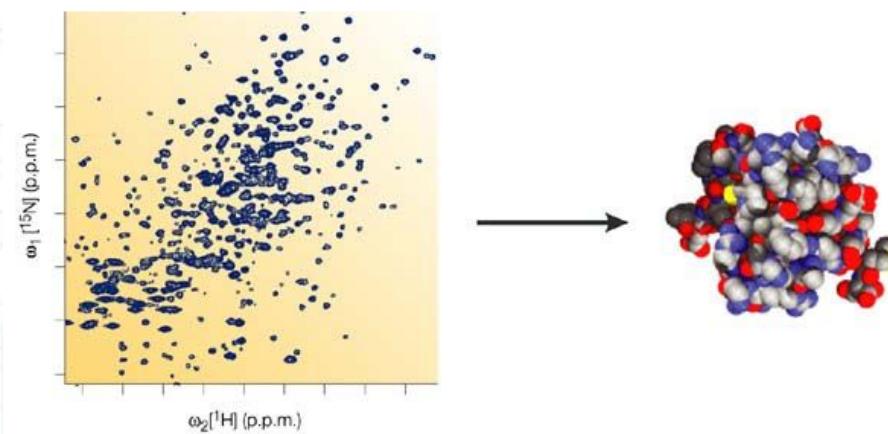
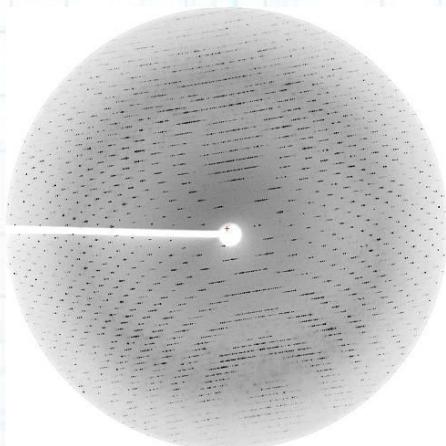
# Motivation

- Proteins structure determines function
  - Medicine
  - Biotechnology
- High discovery rate
- Known to unknown
  - 1:200



# X-Ray Crystallography and NMR Spectroscopy

- X-Ray Crystallography
  - High resolution microscopy
- NMR Spectroscopy
  - Quantum properties of the nucleus

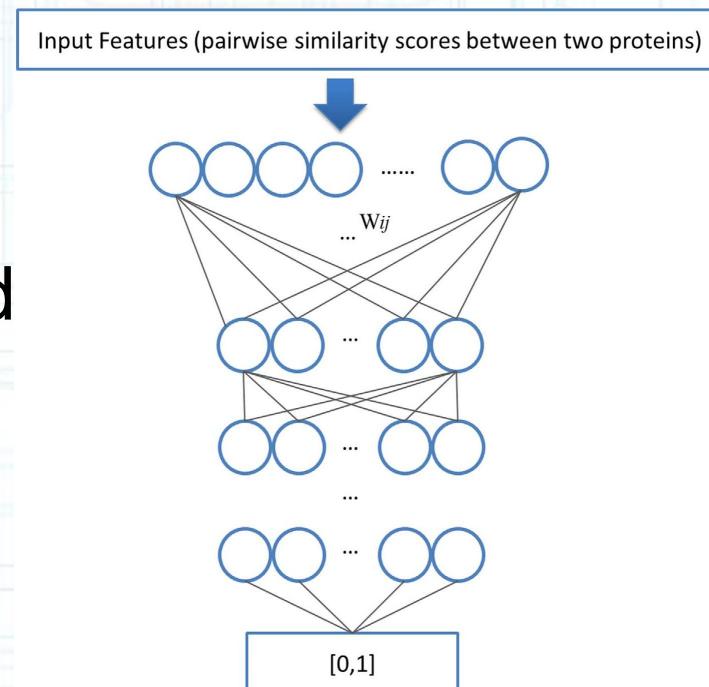


# Machine Learning Techniques

- Support Vector Machines (SVMs)
  - < 50% error rate
- Neural Networks
- Deep Learning
  - Deep Belief networks
  - 84.5% recognition rate

# DN-Fold

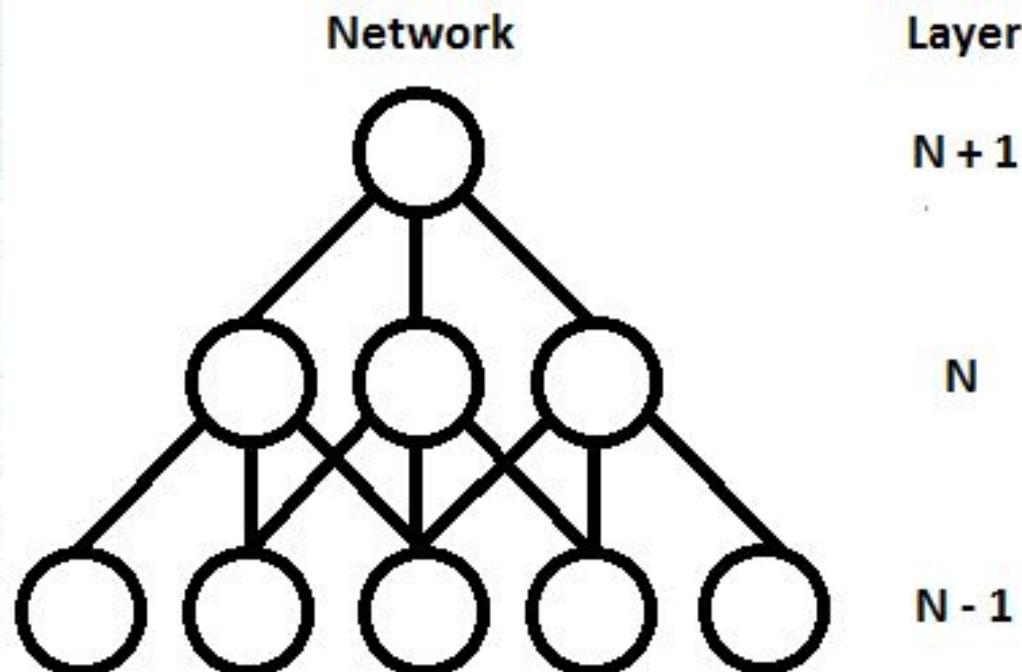
- Deep Belief Networks
  - Restricted Boltzman Machines
  - Generative Autoencoders
- Binary classification problem
  - 976 Proteins
  - $(n^2-n)$
  - Trained and Tested



# Convolutional Neural Networks

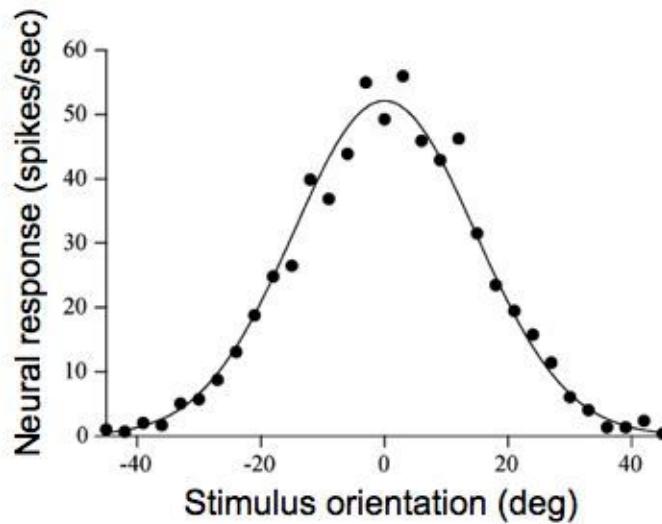
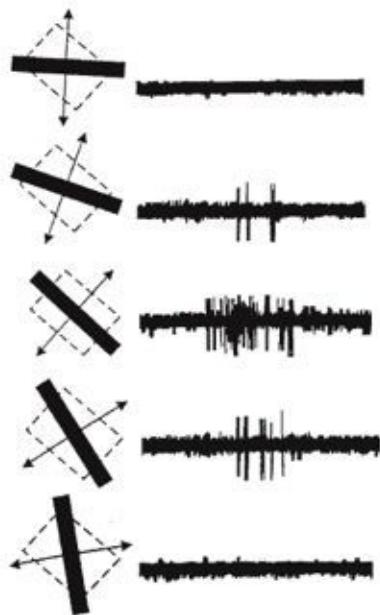
- Receptive Fields
- Surrounding area & Hidden properties
- Fully Connected Deep Neural Network

- Images
- Sounds



# Biological Inspiration

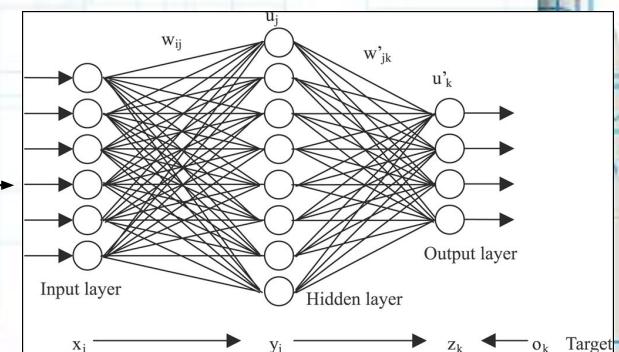
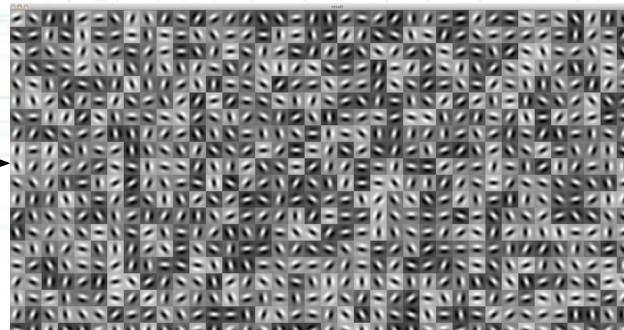
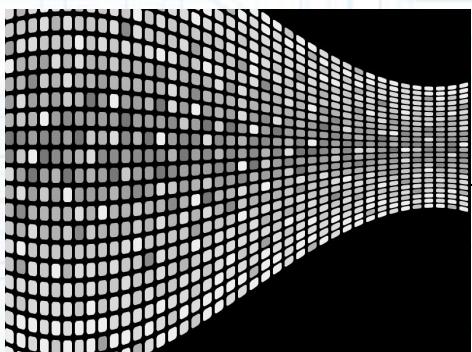
## V1 physiology: orientation selectivity



Hubel & Wiesel, 1968

# Convolutional Layers

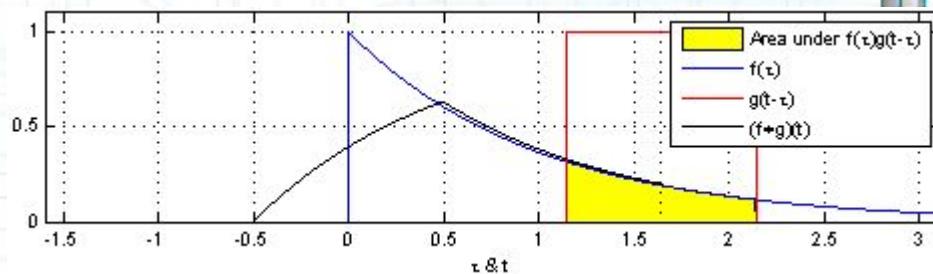
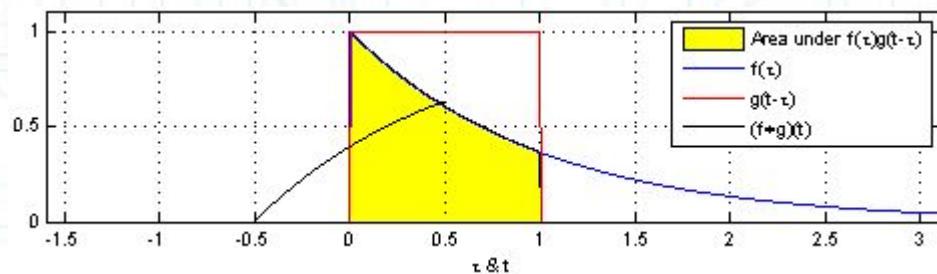
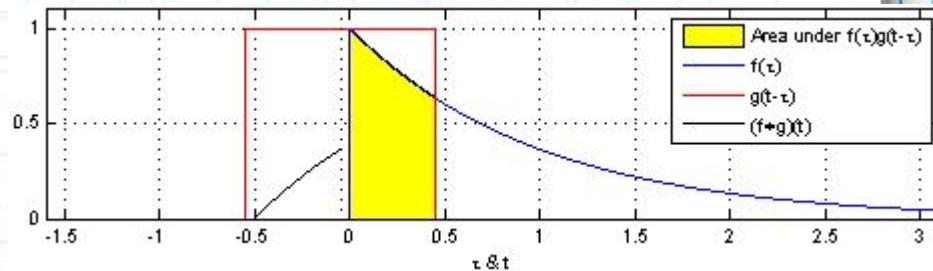
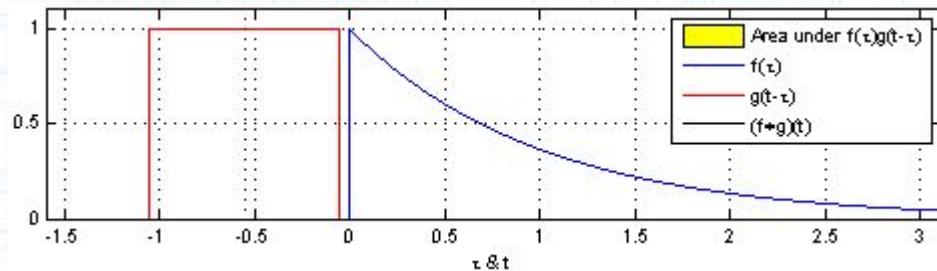
- V1 in the visual cortex
- Input → Eyes
- Filter → V1
- Output → Higher level cortical regions



# Mathematical Convolution

- Multiplying two function mathematically
- Produces an integral

$$\int_{-\infty}^{\infty} \delta(\tau) g(t - \tau) d\tau = g(t)$$



# Discrete Convolution

- Filters used have discrete stride lengths
- Snapshots taken

Data Input																
Step 1	0.1	0.2	0.5	0.62	0.12	0.52	0.23	0.12	0.99	0.04	0.72	0.41	0.55	0.24	0.11	0.12
Step 2	0.1	0.2	0.5	0.62	0.12	0.52	0.23	0.12	0.99	0.04	0.72	0.41	0.55	0.24	0.11	0.12
Step 3	0.1	0.2	0.5	0.62	0.12	0.52	0.23	0.12	0.99	0.04	0.72	0.41	0.55	0.24	0.11	0.12

- Activation Map

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2		

**Bias B0 (1x1x1)**

1

**(Input \* Filter) + Bias**

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	

**Bias B0 (1x1x1)**

1
---

**(Input \* Filter) + Bias**

# Producing an Activation Map

Input (with a pad of 1) (7x7x1)

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

Filter F0 (3x3x1)

-1	1	1
-1	-1	0
-1	0	1

Output (3x3x1)

2	-2	-5

Bias B0 (1x1x1)

1

(Input \* Filter) + Bias

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5		

**Bias B0 (1x1x1)**

1
---

**(Input \* Filter) + Bias**

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	<b>2</b>	<b>1</b>	<b>2</b>	2	0
0	0	<b>2</b>	<b>0</b>	<b>0</b>	1	0
0	1	<b>2</b>	<b>2</b>	<b>1</b>	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5	<b>-1</b>	

**Bias B0 (1x1x1)**

1

(Input \* Filter) + Bias

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5	-1	-1

**Bias B0 (1x1x1)**

1

**(Input \* Filter) + Bias**

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5	-1	-1
4		

**Bias B0 (1x1x1)**

1

**(Input \* Filter) + Bias**

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	<b>2</b>	<b>2</b>	<b>1</b>	1	0
0	0	<b>0</b>	<b>0</b>	<b>0</b>	1	0
0	0	<b>0</b>	<b>0</b>	<b>0</b>	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5	-1	-1
4	<b>2</b>	

**Bias B0 (1x1x1)**

1
---

**(Input \* Filter) + Bias**

# Producing an Activation Map

**Input (with a pad of 1) (7x7x1)**

0	0	0	0	0	0	0
0	1	1	2	2	2	0
0	0	2	1	2	2	0
0	0	2	0	0	1	0
0	1	2	2	1	1	0
0	0	0	0	0	1	0
0	0	0	0	0	0	0

**Filter F0 (3x3x1)**

-1	1	1
-1	-1	0
-1	0	1

**Output (3x3x1)**

2	-2	-5
5	-1	-1
4	2	0

**Bias B0 (1x1x1)**

1

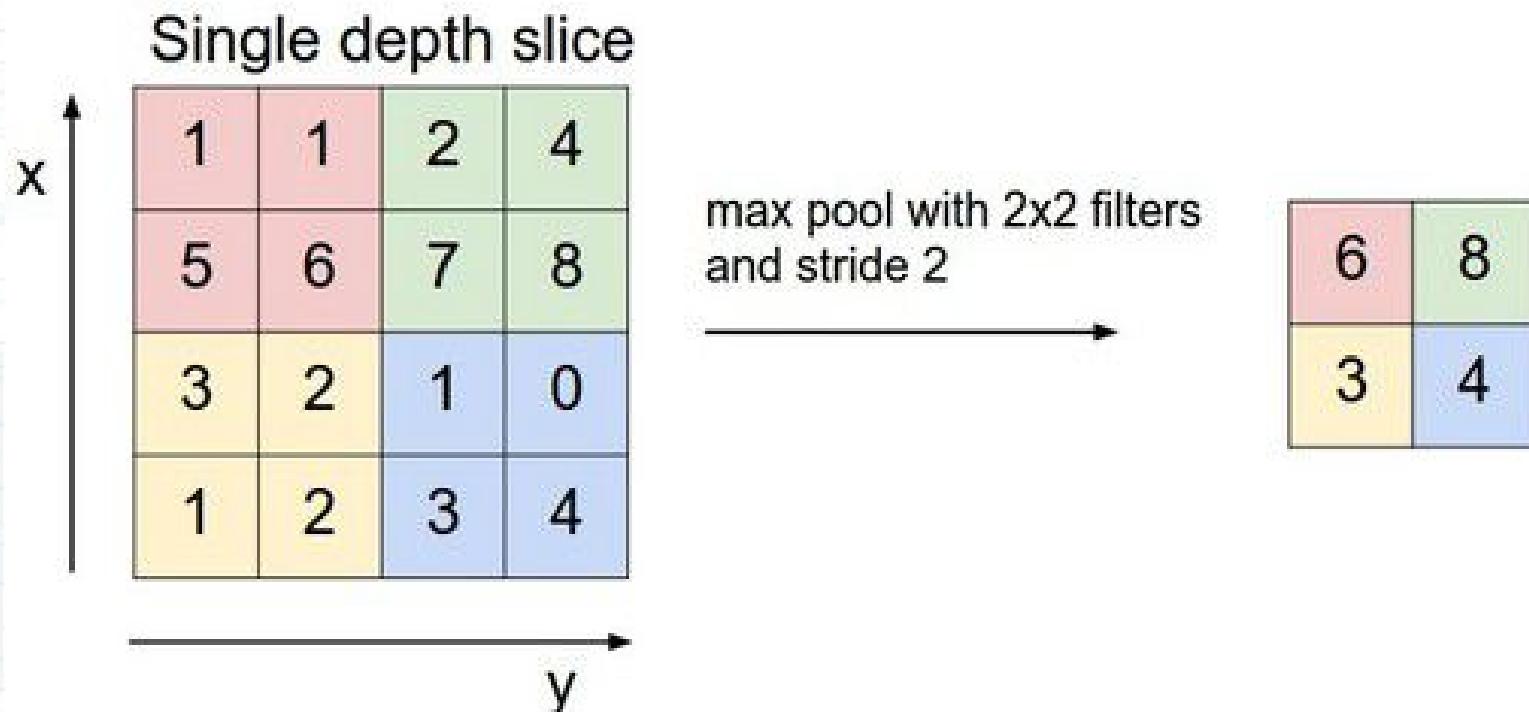
**(Input \* Filter) + Bias**

# Convolutional Network Parameters

- Parameters and their effects
  - Kernel Size
  - Stride length
  - Number of Filters
  - Depth of classifying network
- Optional layers and features
  - Downsampling
  - Dropout technique

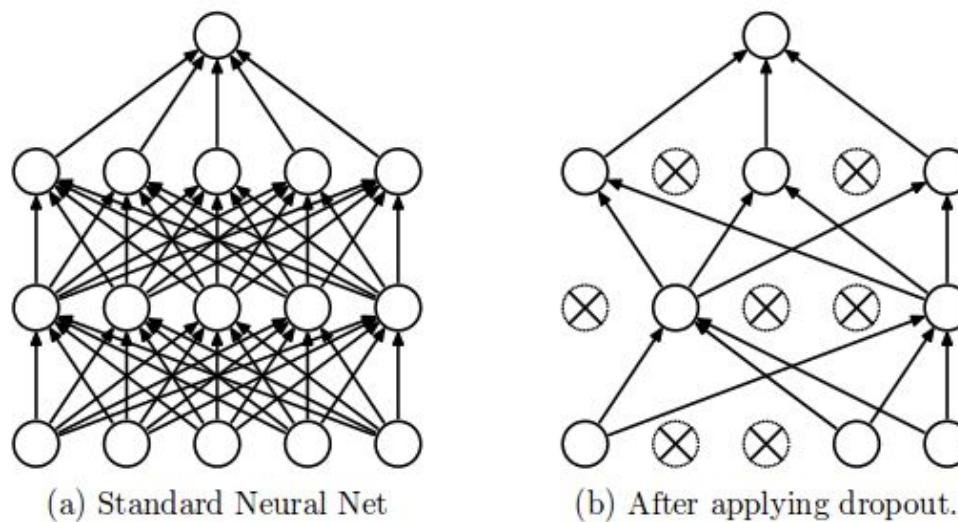
# The Downsampling Layer

- Decrease computational complexity



# Dropout Technique

- Temporarily disable neurons
- Prevents overfitting
- Only used in dense layers



# Dataset

- Derived from the SCOP database
- Proteins
  - Family (555)
  - Superfamily (434)
  - Fold (321)
- Sequence, profile, family alignment and structural information
- 84 data points

```
#1aca-d1aca 1abira-d1abira
-1 1:0.86 2:2.51 3:0.698011213320252 4:0.012914
0.87636911684164 9:0.868627475071523 10:0.45240
14:0.935274598806304 15:0.439352251735332 16:0
860465 20:0.169767441860465 21:-0.4307829160924
3255813953488 28:-0.198450938723838 29:0.136046
8604651162791 35:2.63905732961526 36:0.06976744
2686904762 42:0.526592567947394 43:0.1916390388
0.553370683027986 48:0.262824274759729 49:0.503
53:0.427663172777061 54:0.761204462628882 55:0
767253 59:0.71073499382554 60:0.470352794231018
```

# Models

- Generated 11 simi-random networks
- Varying kernels, strides, classifying networks

Model Number	Network Architecture
Model 1	C21K2S1-D100-D30-O1_30
Model 2	C21K4S1-D100-D30-O1_30
Model 3	C42K8S1-D100-D30-O1_30
Model 4	C42K2S2-D150-D35-O1_30
Model 5	C63K4S2-D150-D35-O1_30
Model 6	C63K8S2-D150-D35-O1_30
Model 7	C84K4S2-D150-D25-O1_30
Model 8	C84K8S2-D150-D25-O1_30
Model 9	C105K16S2-D150-D25-O1_30
Model 10	C105K2S2-D150-D25-O1_30
Model 11	C84K2S2-D100-D100-D30-O1_30

# Model Selection

- Initial testing on a randomized test set
- Chose the top 3 networks to train and test

```
Examples labeled as 0 classified by model as 0: 38582 times
Examples labeled as 0 classified by model as 1: 56221 times
Examples labeled as 1 classified by model as 0: 164 times
Examples labeled as 1 classified by model as 1: 583 times

=====Scores=====
Accuracy: 0.4099
Precision: 0.503
Recall: 0.5937
F1 Score: 0.5446
=====
```

# Results

- Overall CNNs did not outperform DN-Fold
- Provided comparable results to past methods
- Data format
- Label balance

Network	Family		Superfamily		Fold	
	Top 1	Top 5	Top 1	Top 5	Top 1	Top 5
C63K8S2-D150-D35-O1	25.4	51.7	3.7	66.4	4.1	46.3
C84K4S2-D150-D25-O1	33.2	56.8	8.1	67.9	15	58.5
C105K16S2-D150-D25-O1	24.1	37.8	5.6	42.4	10	36.2

Network	Family		Superfamily		Fold	
	Top 1	Top 5	Top 1	Top 5	Top 1	Top 5
PSI-Blast [18]	71.2	72.3	27.4	27.9	4	4.7
THREADER [19]	49.2	58.9	10.8	24.7	14.6	37.7
CNN-FOLD	33.2	56.8	8.1	67.9	15	58.5
DN-FOLD [1]	84.5	91.2	61.5	76.5	33.6	60.7

# Conclusions

- Tasked with CNNs applied to DN-Fold Dataset
- Lacked spacial properties
- More filters not always good

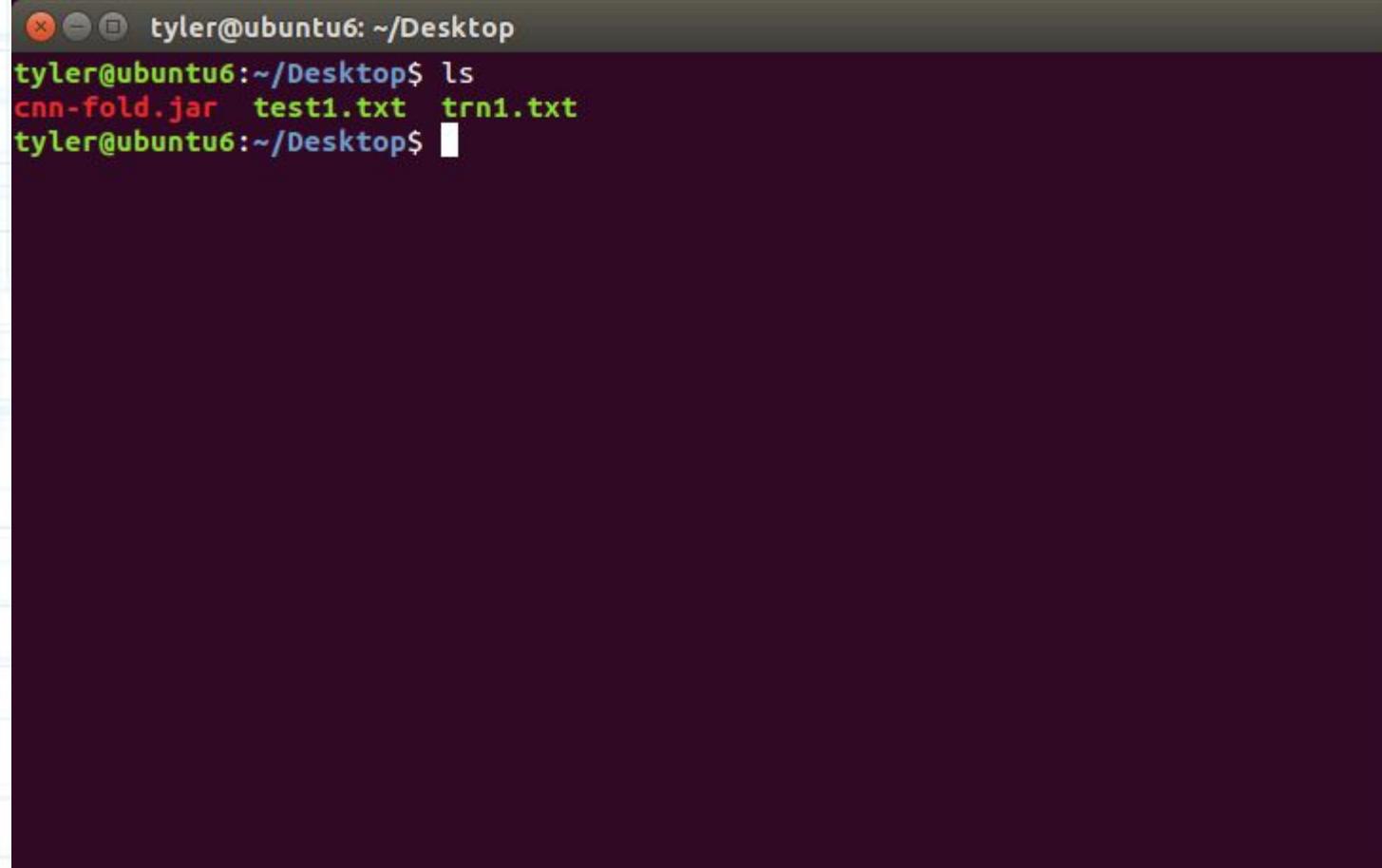
# CNN-Fold

- Program written to obtain results
- Specify CNN-Fold, DN-Fold, Command, or Json network architectures
- Train and Test modes
- Saves trained networks
- Download

```
.der builder = new NeuralNetConfiguration.Builder()
.seed(System.currentTimeMillis())
.iterations(iter)
.learningRate(learningRate)
.momentum(momentum)
.optimizationAlgo(OptimizationAlgorithm.STOCHASTIC_GRADIENT)
.list(numLayers);

for(int i = 0; i < numLayers; i++){
    if(layerConf[i][0] == CONVOLAYER) {
        convo = true;
        builder.layer(currentLayer++, new ConvolutionLayer.I
            .stride(1,layerConf[i][3])
            .nIn(1).nOut(layerConf[i][1]))//input is 1 b
```

# CNN-Fold Demo



```
tyler@ubuntu6: ~/Desktop
tyler@ubuntu6:~/Desktop$ ls
cnn-fold.jar  test1.txt  trn1.txt
tyler@ubuntu6:~/Desktop$
```

# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
tyler@ubuntu6:~/Desktop$ java -jar ./cnn-fold.jar -train -arch C84K4S2-D150-D25-
01 -param C84K4S2.bin -data ./trn1.txt -e 30 -i 30 -l .02 -m .08
```

# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
tyler@ubuntu6:~/Desktop$ java -jar ./cnn-fold.jar -train -arch C84K4S2-D150-D25-01 -param C84K4S2.bin -data ./trn1.txt -e 30 -i 30 -l .02 -m .08
May 02, 2016 10:28:24 PM com.github.fommil.jni.JniLoader liberalLoad
INFO: successfully loaded /tmp/jniloader4142235031727266110netlib-native_system-linux-x86_64.so
22:28:24.427 [main] DEBUG org.reflections.Reflections - going to scan these urls
:
jar:file:/home/tyler/Desktop/cnn-fold.jar!/
22:28:24.637 [main] DEBUG org.reflections.Reflections - could not scan file org/nd4j/linalg/cpu/javacpp/linux-x86_64/libjniLoop.so in url jar:file:/home/tyler/Desktop/cnn-fold.jar!/ with scanner SubTypesScanner
22:28:24.638 [main] DEBUG org.reflections.Reflections - could not scan file org/nd4j/linalg/cpu/javacpp/linux-x86_64/libjniLoop.so in url jar:file:/home/tyler/Desktop/cnn-fold.jar!/ with scanner TypeAnnotationsScanner
22:28:24.647 [main] INFO org.reflections.Reflections - Reflections took 217 ms to scan 1 urls, producing 114 keys and 357 values
Saving C84K4S2-D150-D25-01.json
22:28:25.099 [main] INFO edu.missouri.banks.Network - Train model...
22:28:25.277 [main] WARN o.d.optimize.solvers.BaseOptimizer - Objective function automatically set to minimize. Set stepFunction in neural net configuration to change default settings.
22:28:29.196 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 0 is 0.73593212890625
22:28:32.852 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 1
```

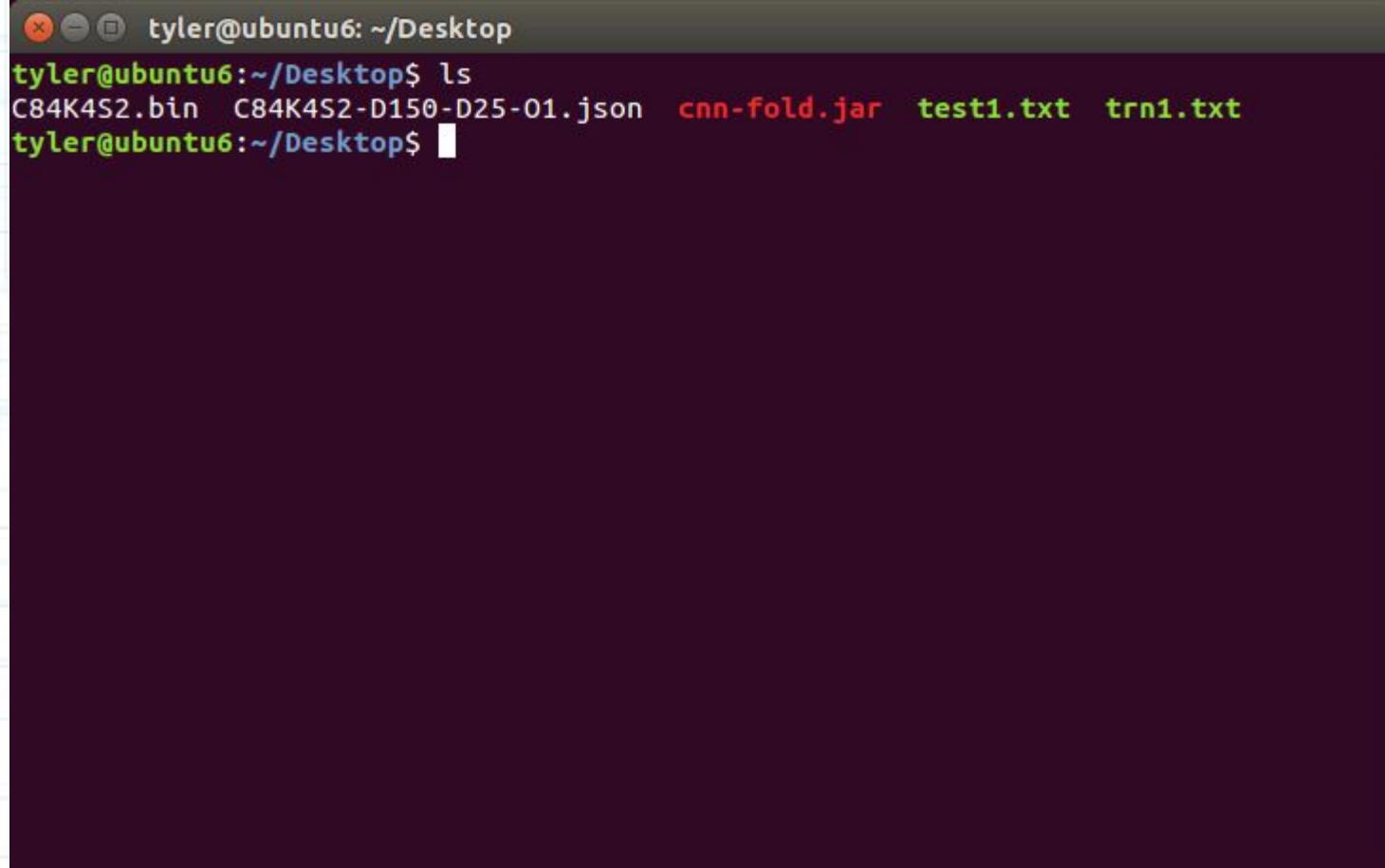
# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
22:29:41.127 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 22
is 0.045599266052246096
22:29:44.362 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 23
is 0.04348321151733398
22:29:47.615 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 24
is 0.04172942733764649
22:29:50.824 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 25
is 0.03977947235107422
22:29:54.035 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 26
is 0.037985916137695315
22:29:57.300 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 27
is 0.03653236389160156
22:30:00.500 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 28
is 0.03524551010131836
22:30:03.718 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 29
is 0.03398983001708984
22:30:03.744 [main] INFO edu.missouri.banks.Network - *** Completed epoch 0 ***
22:30:07.020 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 0
is 0.038803047180175784
22:30:10.274 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 1
is 0.03763349151611328
22:30:13.496 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 2
is 0.03655008697509766
```

# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
is 0.058407379150390626
23:16:59.594 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 21
is 0.058398063659667966
23:17:03.030 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 22
is 0.05838768768310547
23:17:06.405 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 23
is 0.05837990570068359
23:17:09.827 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 24
is 0.05836750030517578
23:17:13.126 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 25
is 0.05835894775390625
23:17:16.539 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 26
is 0.05835003662109375
23:17:19.975 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 27
is 0.058339103698730466
23:17:23.356 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 28
is 0.05832746124267578
23:17:26.684 [main] INFO o.d.o.l.ScoreIterationListener - Score at iteration 29
is 0.05831315994262695
23:17:26.712 [main] INFO edu.missouri.banks.Network - *** Completed epoch 29 ***
*
23:17:26.712 [main] INFO edu.missouri.banks.Network - *****Model tra
in finished*****
tyler@ubuntu6:~/Desktop$
```

# CNN-Fold Demo

A screenshot of a terminal window titled "tyler@ubuntu6: ~/Desktop". The window displays the output of the "ls" command, which lists several files: C84K4S2.bin, C84K4S2-D150-D25-01.json, cnn-fold.jar, test1.txt, and trn1.txt. The terminal has a dark background with light-colored text and standard window controls at the top.

```
tyler@ubuntu6:~/Desktop$ ls
C84K4S2.bin  C84K4S2-D150-D25-01.json  cnn-fold.jar  test1.txt  trn1.txt
tyler@ubuntu6:~/Desktop$
```

# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
tyler@ubuntu6:~/Desktop$ java -jar ./cnn-fold.jar -evaluate -config ./C84K4S2-D1
50-D25-01.json -param C84K4S2.bin -data ./test1.txt
```

# CNN-Fold Demo

```
tyler@ubuntu6:~/Desktop$ java -jar ./cnn-fold.jar -evaluate -config ./C84K4S2-D1  
50-D25-01.json -param C84K4S2.bin -data ./test1.txt  
May 02, 2016 11:20:18 PM com.github.fommil.jni.JniLoader liberalLoad  
INFO: successfully loaded /tmp/jniloader2472205380767391906netlib-native_system-  
linux-x86_64.so  
23:20:18.701 [main] DEBUG org.reflections.Reflections - going to scan these urls  
:  
jar:file:/home/tyler/Desktop/cnn-fold.jar!/br  
23:20:18.886 [main] DEBUG org.reflections.Reflections - could not scan file org/  
nd4j/linalg/cpu/javacpp/linux-x86_64/libjniLoop.so in url jar:file:/home/tyler/D  
esktop/cnn-fold.jar!/ with scanner SubTypesScanner  
23:20:18.887 [main] DEBUG org.reflections.Reflections - could not scan file org/  
nd4j/linalg/cpu/javacpp/linux-x86_64/libjniLoop.so in url jar:file:/home/tyler/D  
esktop/cnn-fold.jar!/ with scanner TypeAnnotationsScanner  
23:20:18.894 [main] INFO org.reflections.Reflections - Reflections took 191 ms  
to scan 1 urls, producing 114 keys and 357 values  
23:20:19.810 [main] INFO edu.missouri.banks.Network - Evaluate model....
```

# CNN-Fold Demo

```
Examples labeled as 0 classified by model as 0: 38582 times  
Examples labeled as 0 classified by model as 1: 56221 times  
Examples labeled as 1 classified by model as 0: 164 times  
Examples labeled as 1 classified by model as 1: 583 times
```

```
=====Scores=====
```

```
Accuracy: 0.4099  
Precision: 0.503  
Recall: 0.5937  
F1 Score: 0.5446
```

# CNN-Fold Demo

```
tyler@ubuntu6: ~/Desktop
tyler@ubuntu6:~/Desktop$ ls
C84K4S2.bin          cnn-fold.jar        test1.txt
C84K4S2-D150-D25-01.json eval-arch-test1.txt  trn1.txt
tyler@ubuntu6:~/Desktop$ head -n 10 eval-arch-test1.txt
[[0.46,0.54]
 [0.48,0.52]
 [0.65,0.35]
 [0.43,0.57]
 [0.46,0.54]
 [0.46,0.54]
 [0.47,0.53]
 [0.45,0.55]
 [0.42,0.58]
 [0.44,0.56]
tyler@ubuntu6:~/Desktop$
```

# CNN-Fold Demo

C84 Results

Family: The protein list size is: 555

Top1:184 Top5:315

Top1\_acc:0.331531531531532 Top5\_acc:0.567567567567568

Fold: The protein list size is: 321

Top1:26 Top5:218

Top1\_acc:0.0809968847352025 Top5\_acc:0.679127725856698

SuperFamily: The protein list size is: 434

Top1:65 Top5:254

Top1\_acc:0.149769585253456 Top5\_acc:0.585253456221198

# Future Work

- Additional methods of conveying proteins
  - Mimic image or sound
  - Data normalization
- CNN-Fold
  - Generalized input
  - Additional command line parameters

**Thank you!**

# Image Reference

- Protein - [https://commons.wikimedia.org/wiki/File:Protein\\_HSPA8\\_PDB\\_1atr.png](https://commons.wikimedia.org/wiki/File:Protein_HSPA8_PDB_1atr.png)
- X Ray - [http://chemwiki.ucdavis.edu/Core/Analytical\\_Chemistry/Instrumental\\_Analysis/Diffraction/X-ray\\_Crystallography](http://chemwiki.ucdavis.edu/Core/Analytical_Chemistry/Instrumental_Analysis/Diffraction/X-ray_Crystallography)
- NMR - [http://www.nature.com/horizon/proteinfolding/background/figs/technology\\_f3.html](http://www.nature.com/horizon/proteinfolding/background/figs/technology_f3.html)
- DNN - Jo, Taeho, et al. "Improving Protein Fold Recognition by Deep Learning Networks." *Scientific reports* 5 (2015).
- Filter - <http://eric-yuan.me/fake-cnn/>
- Random data- <http://www.thesearchagents.com/2013/06/google-wants-to-collect-our-data-to-show-us-what-we-want-why-do-our-governments-want-it/>
- Convolution gif - [https://upload.wikimedia.org/wikipedia/commons/b/b9/Convolution\\_of\\_spiky\\_function\\_with\\_box2.gif](https://upload.wikimedia.org/wikipedia/commons/b/b9/Convolution_of_spiky_function_with_box2.gif)
- Downsampling – deeplearning4j.org
- Dropout - Srivastava et. al.