CS 696 Intro to Big Data: Tools and Methods
Spring Semester, 2019
Doc 19 Clustering & Deep Learning
Apr 16, 2019

Software 2.0

https://medium.com/@karpathy/software-2-0-a64152b37c35

Andrej Karpathy

Director of AI at Tesla

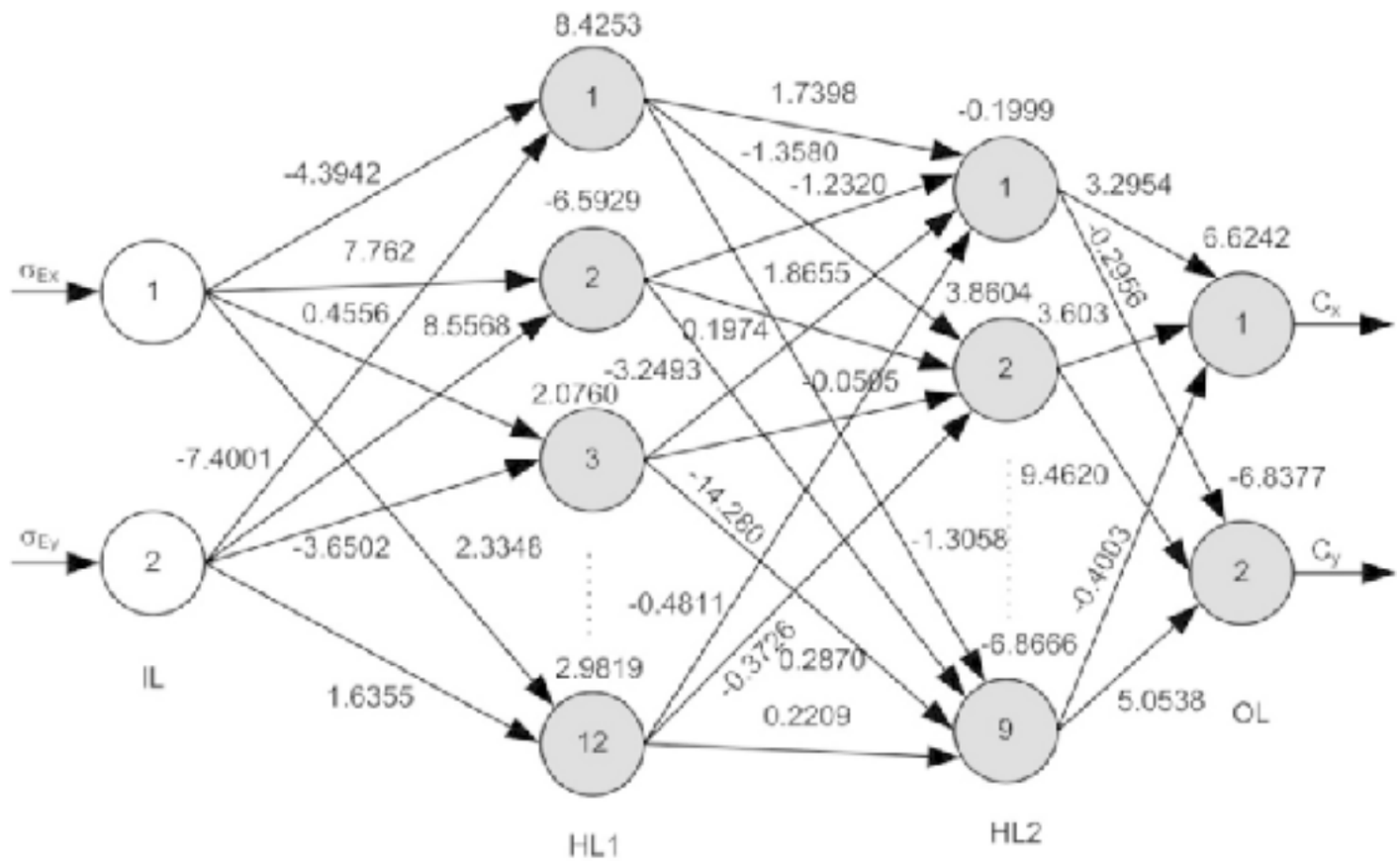
Nov 11, 2017

Software 1.0

Writing code in Java, Python, Scala, Kotlin, C, ...

Software 2.0

Training neural nets



"Software 2.0 is not going to replace 1.0 (indeed, a large amount of 1.0 infrastructure is needed for training and inference to “compile” 2.0 code), but it is going to take over increasingly large portions of what Software 1.0 is responsible for today."

Visual Recognition

Speech Recognition

Speech Synthesis

Machine Translation

Robotics

Games

Benefits of Software 2.0

Computationally homogeneous
  matrix multiplication and thresholding

Simple to bake into silicon

Constant running time

Constant memory use

It is highly portable

It is very agile
  Too slow use fewer layers
  More data -> better software

It is better than you

Limitations of Software 2.0

Don't know why models does what it does

Can do odd things

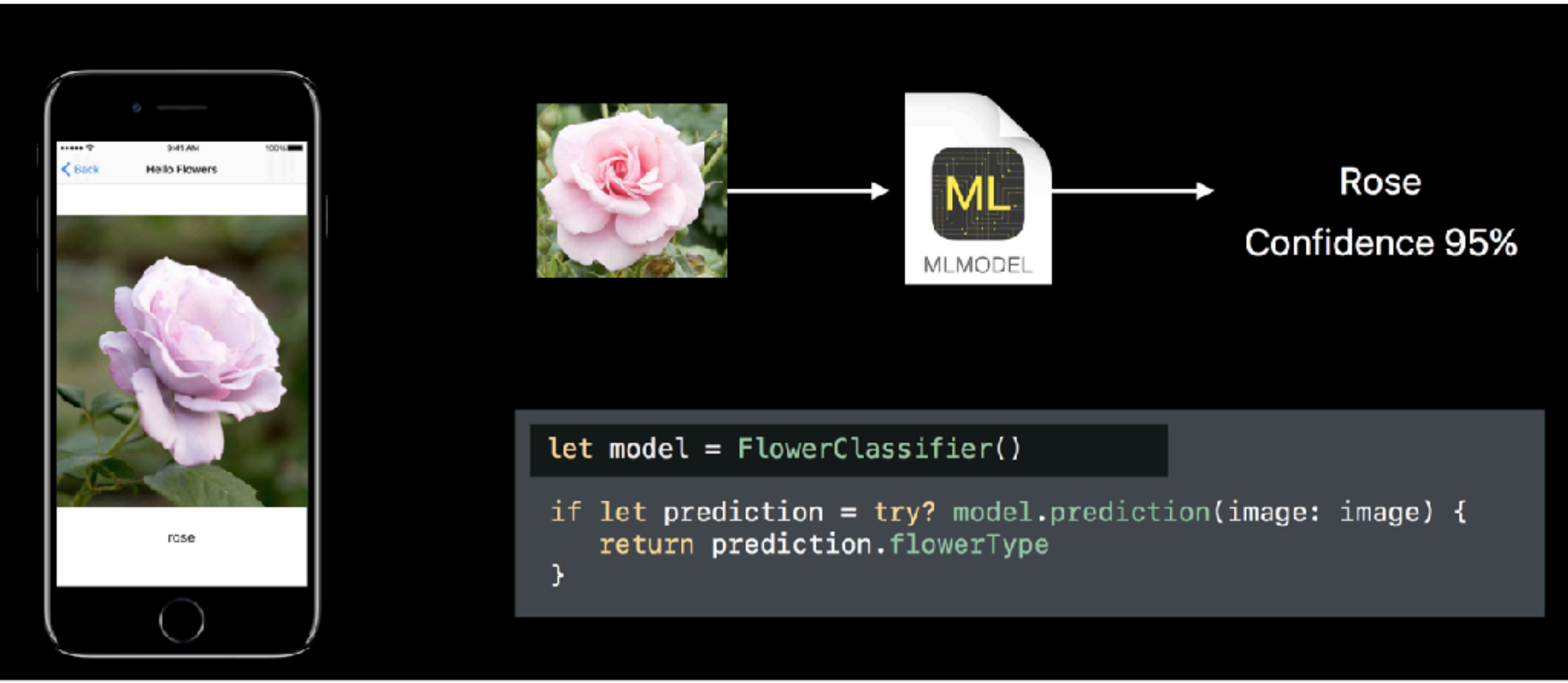
Microsoft's Tay bot

- Became a Nazi

APPLE CAN TRY TO FIX THE AUTOCORRECT BUG, BUT I'VE ALREADY INCORPORATED IT INTO MY HANDWRITING.

Apple MLKit

Apple MLKit


Back
Hello Flowers
rose
ML
MLMODEL
Rose
Confidence 95%
let model = FlowerClassifier()
if let prediction = try? model.prediction(image: image) {
return prediction.flowerType
}

Apple Watch detects Hypertension & Sleep

Cardiogram & University of California San Fransisco

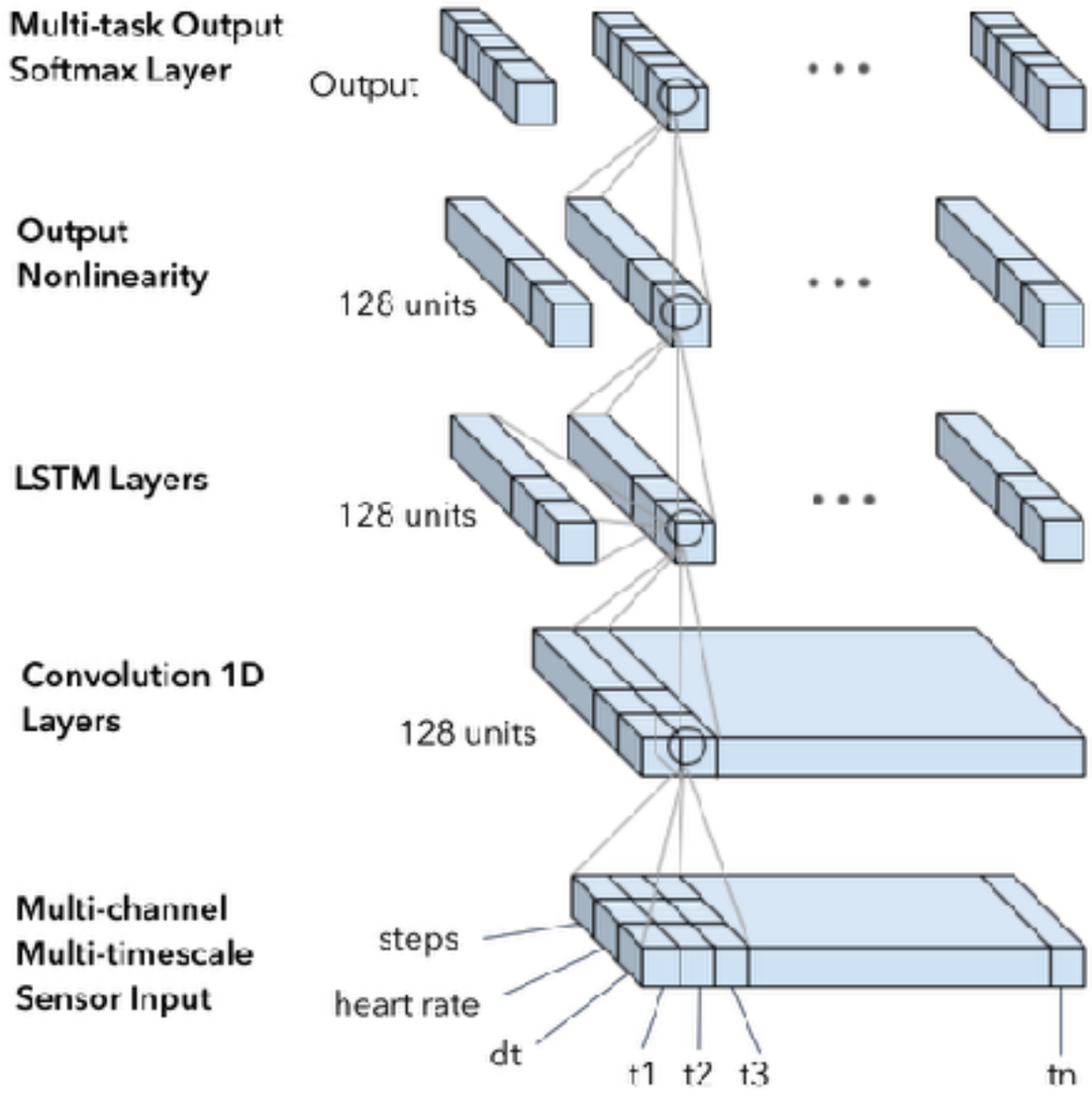
Trained deep learning algorithm DeepHeart on existing data

Had 6,158 people in test
- 70% training set
- 30% test set
- People used watch for 1 to 53 weeks

Sleep apnea 90 accuracy

Hypertension 82% accuracy

Detecting irregular heart rhythms 97%



iOS & Android

iOS ML Kit

- Scikit Learn models

Android

- TensorFlow Lite model

Spark ML

Classification

- Logistic Regression
- Decision tree classier
- Random forest classifier
- Gradient-boosted tree classifier
- Multilayer perceptron classifier
- Linear Support Vector Machine
- One-vs-Rest classifier
- Naive Bayes

Clustering

- K-means
- Latent Dirichlet allocation
- Gaussian Mixture Model

How to Export Spark Models

Spark.mllib supports PMML for some models

`spark.mllib` model	PMML model
KMeansModel	ClusteringModel
LinearRegressionModel	RegressionModel
RidgeRegressionModel	RegressionModel
LassoModel	RegressionModel
SVMModel	RegressionModel
Binary LogisticRegressionModel	RegressionModel

Predictive Model Markup Language

XML-based predictive model interchange format

ML libraries and Model interoperability

https://www.andrey-melentyev.com/model-interoperability.html

Spark k-Means

pyspark.ml.clustering.KMeans

Need to set number of clusters
Can set a seed

iris.txt

```
sepal_length,sepal_width,petal_length,petal_
5.1,3.5,1.4,0.2,setosa
4.9,3.0,1.4,0.2,setosa
4.7,3.2,1.3,0.2,setosa
4.6,3.1,1.5,0.2,setosa
5.0,3.6,1.4,0.2,setosa
5.4,3.9,1.7,0.4,setosa
4.6,3.4,1.4,0.3,setosa
5.0,3.4,1.5,0.2,setosa
```

```
import pyspark.sql as sql
spark = sql.SparkSession.builder \
        .master("local[4]") \
        .appName("Sample") \
        .getOrCreate()

iris = spark.read.format("csv"). \
       option("header",True).\
       option("inferschema",True).\
       load("iris.txt")

from pyspark.ml.feature import StringIndexer
from pyspark.ml.feature import VectorAssembler
from pyspark.ml import Pipeline

iris_indexer = StringIndexer(inputCol="species", outputCol="label").fit(iris)
iris_assembler = VectorAssembler(inputCols=["sepal_length","sepal_width", "petal_length",
"petal_width"], outputCol="features")

pipeline = Pipeline(stages=[iris_indexer, iris_assembler])
iris_formated = pipeline.fit(iris).transform(iris)
```

iris_formated.show()

```
+------------+-----------+------------+-----------+-------+-----+-----------------+
|sepal_length|sepal_width|petal_length|petal_width|species|label|         features|
+------------+-----------+------------+-----------+-------+-----+-----------------+
|         5.1|        3.5|         1.4|        0.2| setosa|  2.0|[5.1,3.5,1.4,0.2]|
|         4.9|        3.0|         1.4|        0.2| setosa|  2.0|[4.9,3.0,1.4,0.2]|
|         4.7|        3.2|         1.3|        0.2| setosa|  2.0|[4.7,3.2,1.3,0.2]|
|         4.6|        3.1|         1.5|        0.2| setosa|  2.0|[4.6,3.1,1.5,0.2]|
|         5.0|        3.6|         1.4|        0.2| setosa|  2.0|[5.0,3.6,1.4,0.2]|
|         5.4|        3.9|         1.7|        0.4| setosa|  2.0|[5.4,3.9,1.7,0.4]|
|         4.6|        3.4|         1.4|        0.3| setosa|  2.0|[4.6,3.4,1.4,0.3]|
|         5.0|        3.4|         1.5|        0.2| setosa|  2.0|[5.0,3.4,1.5,0.2]|
|         4.4|        2.9|         1.4|        0.2| setosa|  2.0|[4.4,2.9,1.4,0.2]|
|         4.9|        3.1|         1.5|        0.1| setosa|  2.0|[4.9,3.1,1.5,0.1]|
|         5.4|        3.7|         1.5|        0.2| setosa|  2.0|[5.4,3.7,1.5,0.2]|
|         4.8|        3.4|         1.6|        0.2| setosa|  2.0|[4.8,3.4,1.6,0.2]|
|         4.8|        3.0|         1.4|        0.1| setosa|  2.0|[4.8,3.0,1.4,0.1]|
```

```
from pyspark.ml.clustering import KMeans, KMeansModel
clusters = KMeans(k = 3)

iris_model = clusters.fit(iris_formated)

centers = iris_model.clusterCenters()
print("Cluster Centers: ")
for center in centers:
    print(center)
```

```
Cluster Centers:
[5.88360656 2.74098361 4.38852459 1.43442623]
[6.85384615 3.07692308 5.71538462 2.05384615]
[5.006 3.418 1.464 0.244]
```

```
predictions = iris_model.transform(iris_formated)

predictions.filter(predictions.label != predictions.prediction).show()
```

```
+------------+-----------+------------+-----------+----------+-----+-----------------+----------+
|sepal_length|sepal_width|petal_length|petal_width|   species|label|         features|prediction|
+------------+-----------+------------+-----------+----------+-----+-----------------+----------+
|         7.0|        3.2|         4.7|        1.4|versicolor|  0.0|[7.0,3.2,4.7,1.4]|         1|
|         6.9|        3.1|         4.9|        1.5|versicolor|  0.0|[6.9,3.1,4.9,1.5]|         1|
|         6.7|        3.0|         5.0|        1.7|versicolor|  0.0|[6.7,3.0,5.0,1.7]|         1|
|         5.8|        2.7|         5.1|        1.9| virginica|  1.0|[5.8,2.7,5.1,1.9]|         0|
|         4.9|        2.5|         4.5|        1.7| virginica|  1.0|[4.9,2.5,4.5,1.7]|         0|
|         5.7|        2.5|         5.0|        2.0| virginica|  1.0|[5.7,2.5,5.0,2.0]|         0|
|         5.8|        2.8|         5.1|        2.4| virginica|  1.0|[5.8,2.8,5.1,2.4]|         0|
|         6.0|        2.2|         5.0|        1.5| virginica|  1.0|[6.0,2.2,5.0,1.5]|         0|
|         5.6|        2.8|         4.9|        2.0| virginica|  1.0|[5.6,2.8,4.9,2.0]|         0|
|         6.3|        2.7|         4.9|        1.8| virginica|  1.0|[6.3,2.7,4.9,1.8]|         0|
|         6.2|        2.8|         4.8|        1.8| virginica|  1.0|[6.2,2.8,4.8,1.8]|         0|
|         6.1|        3.0|         4.9|        1.8| virginica|  1.0|[6.1,3.0,4.9,1.8]|         0|
|         6.3|        2.8|         5.1|        1.5| virginica|  1.0|[6.3,2.8,5.1,1.5]|         0|
|         6.0|        3.0|         4.8|        1.8| virginica|  1.0|[6.0,3.0,4.8,1.8]|         0|
|         5.8|        2.7|         5.1|        1.9| virginica|  1.0|[5.8,2.7,5.1,1.9]|         0|
|         6.3|        2.5|         5.0|        1.9| virginica|  1.0|[6.3,2.5,5.0,1.9]|         0|
|         5.9|        3.0|         5.1|        1.8| virginica|  1.0|[5.9,3.0,5.1,1.8]|         0|
+------------+-----------+------------+-----------+----------+-----+-----------------+----------+
```

17/151 incorrect

```
from pyspark.ml.evaluation import ClusteringEvaluator
evaluator = ClusteringEvaluator()
silhouette = evaluator.evaluate(predictions)
print("Silhouette with squared euclidean distance = " + str(silhouette))
```

```
Silhouette with squared euclidean distance = 0.7342113066202725
```

Silhouettes

Method for validating clusters of data

a(i) = average distance from i-th point to other points within the same cluster

if i is in wrong cluster a(i) will be high

b(i, k) = average distance from the i-th point to the points in the k-th cluster

b(i) = min b(i,k) over all all k except for k = i

if i-th point is in wrong cluster b(i) will be low

s(i) = b(i) - a(i)/ max(a(i), b(i))

-1 <= s(i) <= 1

Silhouettes

$$s(i) = b(i) - a(i) / \max(a(i), b(i))$$

$-1 <= s(i) <= 1$

s(i) close to 1 indicates i-th point well within cluster

PCA - Principle Component Analysis

Used to reduce the dimensionality of data

Changes the dimension of the data so

First dimension has the greatest variance
Second dimension has second greatest variance
...

Can then select first K dimensions to work with

Data is transformed into different corrdinate system

```
from pyspark.ml.feature import PCA
pca = PCA(k=3, inputCol="features", outputCol="pca_features")
model = pca.fit(iris_formated)
iris_transformed = model.transform(iris_formated)

model.explainedVariance
```

```
DenseVector([0.9246, 0.053, 0.0172])
```

Sum 0.9948

```
pca = PCA(k=2, inputCol="features", outputCol="pca_features")
model.explainedVariance
```

```
DenseVector([0.9246, 0.053])
```

Sum 0.9776

```
iris_transformed.select("features", "pca_features").show(n=20, truncate=60)
```

```
+-----------------+------------------------------------------------------------+
|         features|                                                pca_features|
+-----------------+------------------------------------------------------------+
|[5.1,3.5,1.4,0.2]| [-2.827135972679027,-5.641331045573321,0.6642769315107171]|
|[4.9,3.0,1.4,0.2]|[-2.795952482148843 7,-5.145166883252896,0.8462865195142029]|
|[4.7,3.2,1.3,0.2]|[-2.6215235581650584,-5.177378121203909,0.6180558535097703]|
|[4.6,3.1,1.5,0.2]| [-2.7649059004742402,-5.003599415056946,0.605093119223434]|
|[5.0,3.6,1.4,0.2]|[-2.782750115951660 3,-5.648648294377395,0.546535394734156 9]|
|[5.4,3.9,1.7,0.4]|[-3.231445736773378,-6.062506444034077,0.4684394754923788 5]|
|[4.6,3.4,1.4,0.3]|[-2.690452415602345,-5.232619219784267,0.3785140093180462 4]|
|[5.0,3.4,1.5,0.2]|[-2.884861104459156 3,-5.485129079769225,0.6585666047730699]|
|[4.4,2.9,1.4,0.2]|[-2.623384532447340 6,-4.743925704477345,0.6154296883942059]|
|[4.9,3.1,1.5,0.1]|[-2.837498411063853 7,-5.208032027056187,0.8342983942443872]|
```

Neural Networks

All you really need to know for the moment is that the universe is a lot more complicated than you might think, even if you start from a position of thinking it's pretty damn complicated in the first place.

--- Douglas Adams, Hitchhikers Guide to the Universe

Example

Apples	Oranges	Total Cost
2	3	5
9	4	16
4	8	10.5

Find w(a) and w(o)

let w(a) = cost of apple

n(a) = number of apples

w(o) = cost of orange

n(o) = number of oranges

t = transaction fee

Total Cost = w(a)*n(a) + w(o)*n(o) + t

Apples	Oranges	Total Cost	Guess
2	3	5	2.5
9	4	16	6.5
4	8	10.5	6

w(a) - guess 0.5
w(o) - guess 0.5
t - guess 0

Too low

Apples	Oranges	Total Cost	Guess
2	3	5	6
9	4	16	14
4	8	10.5	13

w(a) - guess 1
w(o) - guess 1
t - guess 1

Too high in two cases

Apples	Oranges	Total Cost	Guess
2	3	5	4.5
9	4	16	10.5
4	8	10.5	9.75

w(a) - guess 0.75
w(o) - guess 0.75
t - guess 0.75

Too low

Need

Measure of how far off guess is from data

Systematic way to change weights

Loss Function

Measure of how the data differs from estimate

Linear case

$$\mathfrak{L}(\mathbf{W}, \boldsymbol{b}) = \frac{1}{N} \sum_{i=1}^{N} (\hat{Y}_i - Y_i)^2$$

Y_i = data value

Y_i_hat = computed value

Activation Function

Function that we are trying to fit

In example linear function with two independent variables

$$f(x1,x2) = a*x1 + b*x2 + c$$

$$= w1*x1 + w2*x2 + b$$

w1, w1 are the weights

b is the bias

Bias

Prejudice in favor of one thing

Positive values being for
Negative values being against

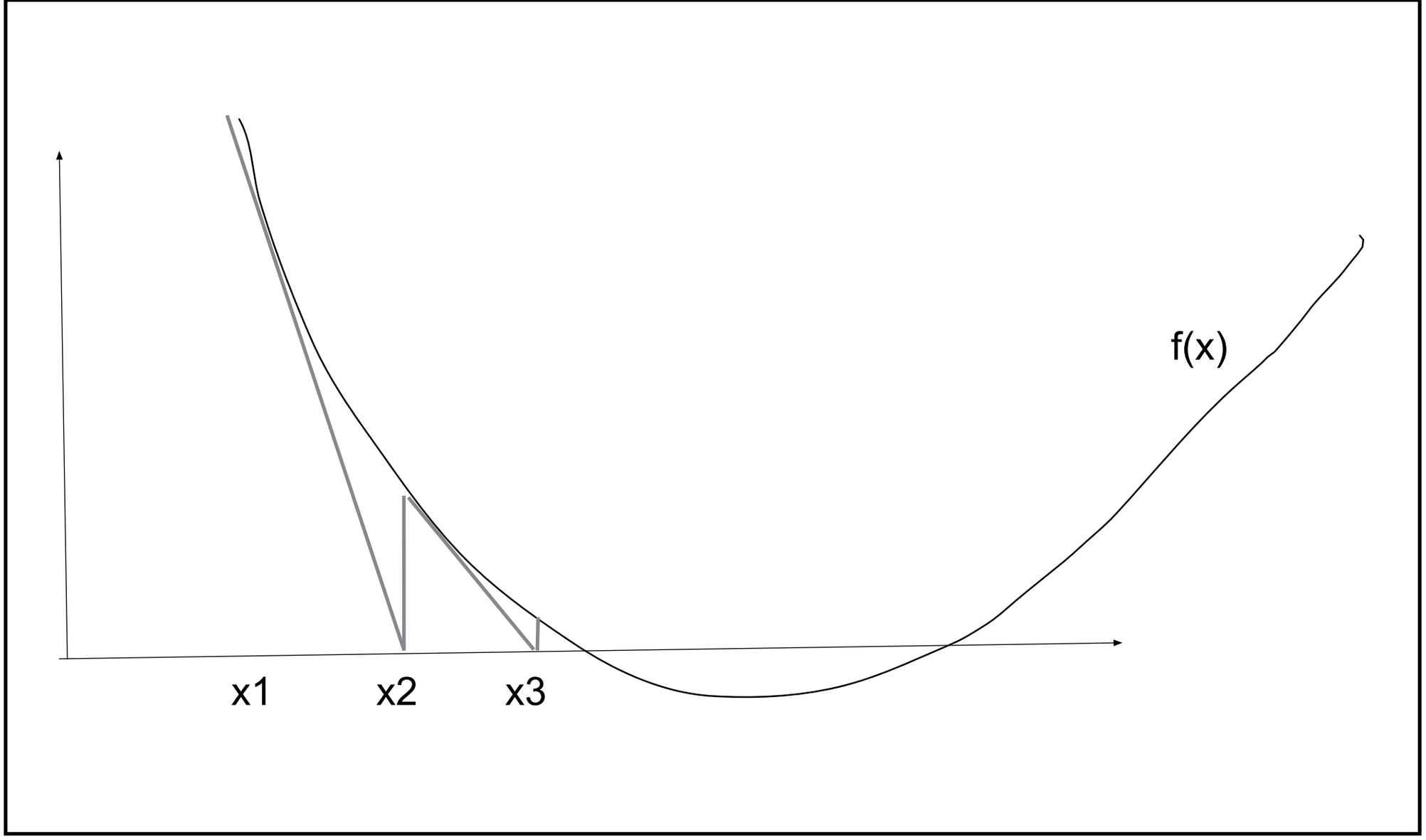
Consider x = 0 neutral input

Then if f is neutral function f(0) == 0

f(x1, x2) = w1*x1 + w2*x2 + b

f(0, 0) = b

So f has a bias



Pick x1

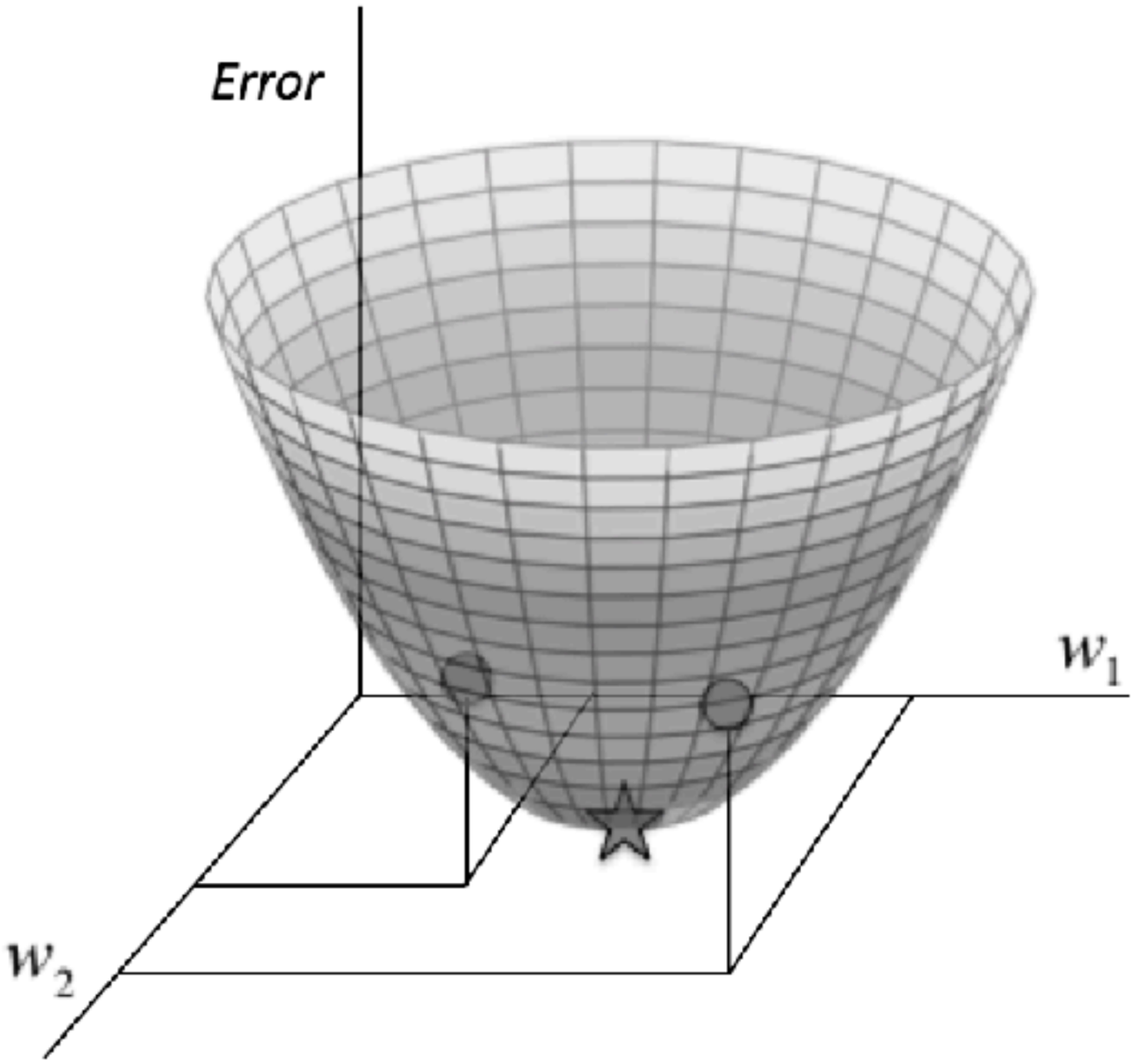
Find the slope at f(x1) ie take derivative

Use slope to estimate where f(x) is zero = x2

Repeat process until f(xn) is really close to 0

Gradient Descent

gradient is the derivative of multi-dimensional function



Systematic way to change weights

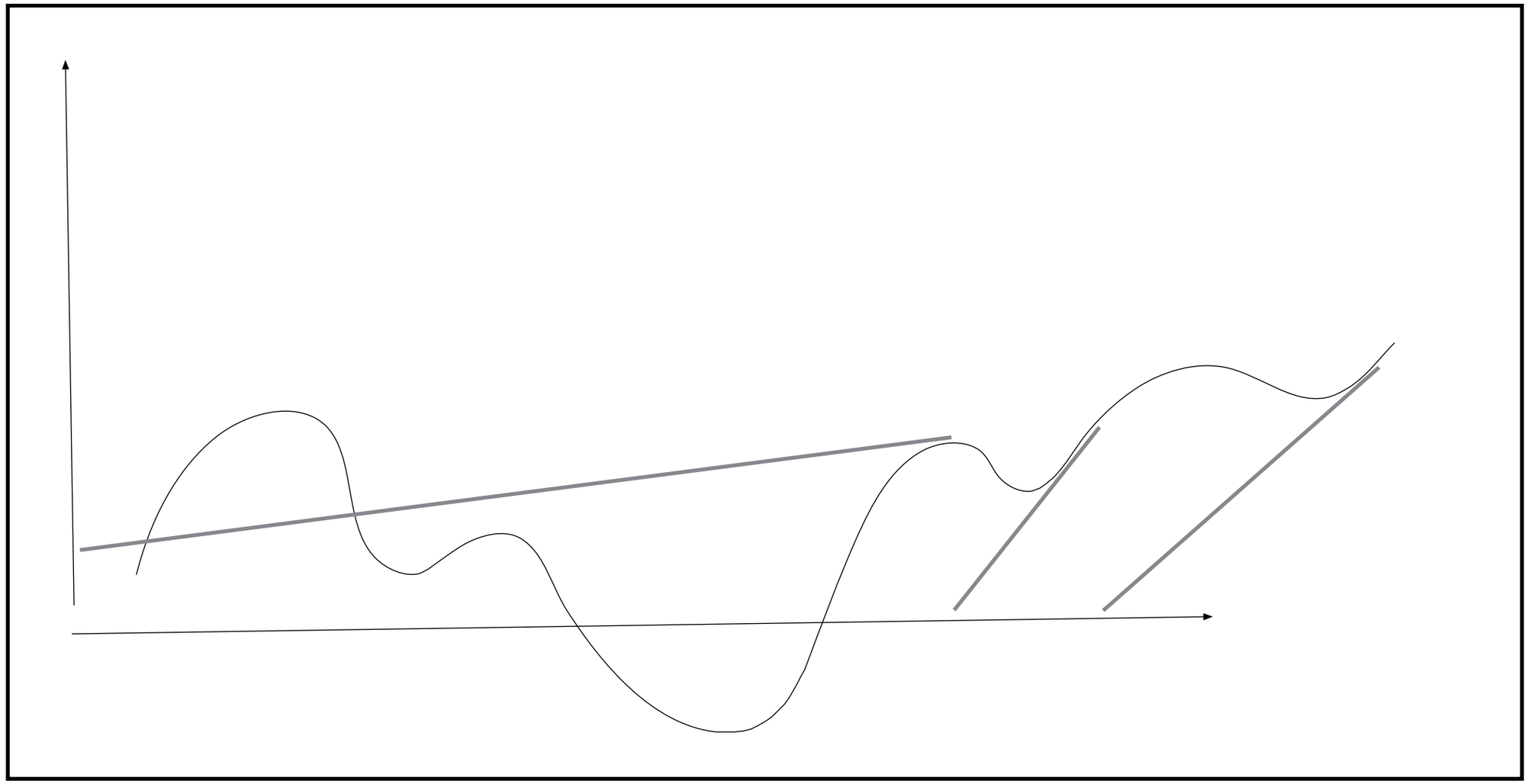
f(x1, x2) = w1*x1 + w2*x2 + b

Take derivative of activation function get gradient

Use the slope in the x1 dimension to adjust w1

Use the slope in the x2 dimension to adjust w2

How far to go?



Learning Rate

To avoid overshooting multiply the gradient by a factor - say 0.1

This is called the learning rate

Take derivative of activation function get gradient

Use the slope in the x1 dimension * learning rate to adjust w1

Use the slope in the x2 dimension * learning rate to adjust w2

Terms

Loss Function

Activation Function

Learning Rate

Weights

Bias

Basic Algorithm

f(x1, x2) = w1*x1 + w2*x2 + b

Select initial values for w1, w2, b

1. Compute loss function on data to find the error

2. Update w1, w2, b

Take derivative of activation function get gradient

w1 = w1 + the slope in the x1 dimension * learning rate * Error

w2 = w2 + the slope in the x2 dimension * learning rate * Error

b = b + gradient * learning rate * Error

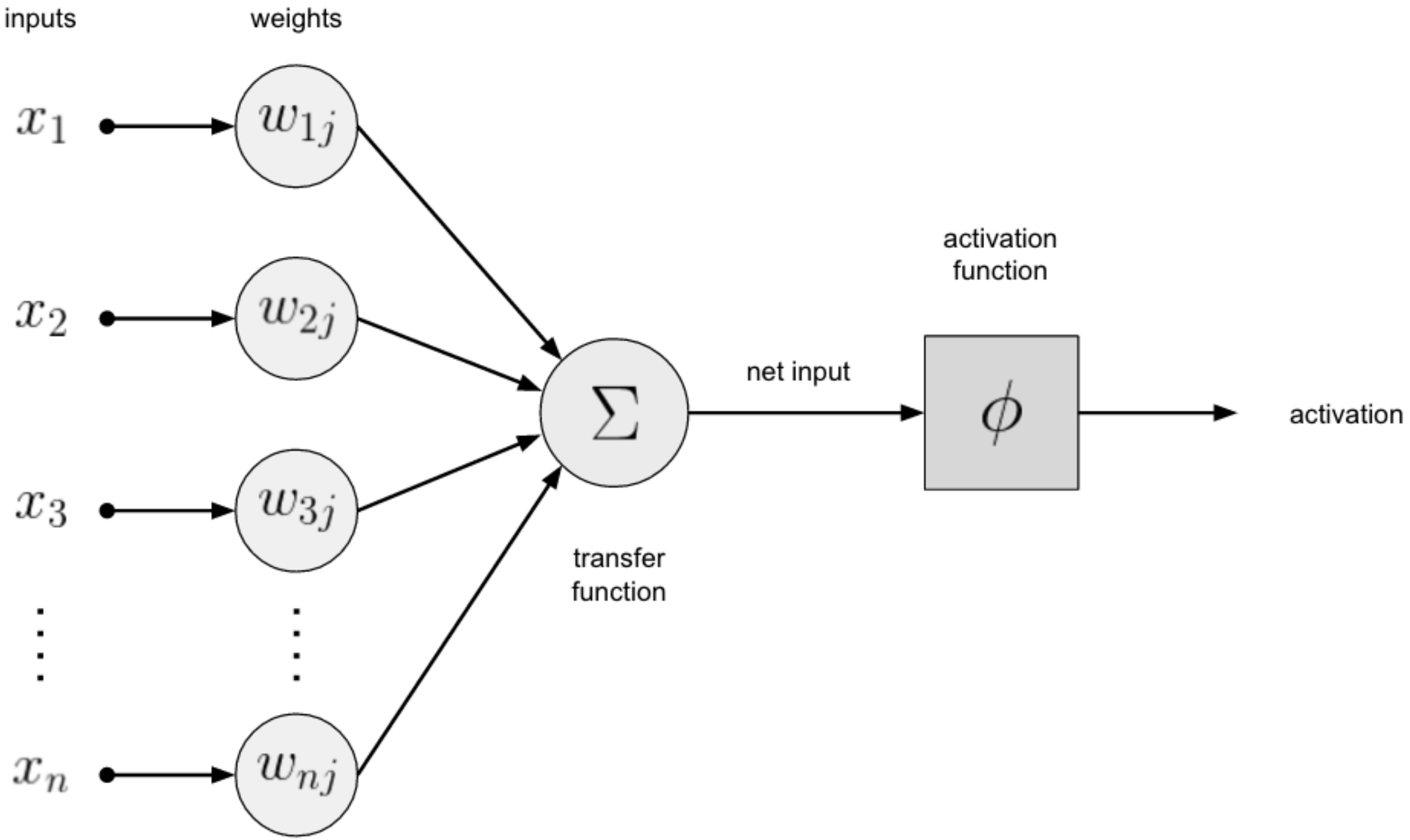
Repeat 1 & 2 until error is acceptable

Learning Rate

If too small then take too long for result to converge

If too large then algorithm will jump around too much and not converge

Basic Structure of Neuron



Linear Knet Example

```
using Knet

activation(w,x) = w[1]*x .+ w[2]

loss(w,x,y) = sumabs2(y - activation(w,x)) / size(y,2)

lossgradient = grad(loss)                    # grad computed gradient
```

```
function train(w, data; learning_rate=.1)
    for (x,y) in data
        dw = lossgradient(w, x, y)
        for i in 1:length(w)
            w[i] -= learning_rate * dw[i]
        end
    end
end
```

```
x = rand(10)
y = 2 .* x .+ 3          #exact model so we know
x = x'
y = y'
w = [2.5,3.5]

for i in 1:20
  train(w,[(x,y)], learning_rate = 0.1)
  println(loss(w,x,y))
end
```

Loss value

First 0.34

Last 0.001

w:

2.09855

2.94161

Varying Learning Rate

Learning rate 0.01

Loss value

First 0.43
Last 0.26

w:

2.45
3.29

Learning rate 0.1

Loss value

First 0.34
Last 0.001

w:

2.10
2.94

Learning rate 1.0

Loss value

First 0.83
Last 21299.5

w:

53.0
129

Varying Starting Point

Learning rate 0.01
w = [0.0, 0.0]

Loss value	w:
First 9.1	1.76
Last 0.005	3.12

Learning rate 0.01
w = [-10.0, -10.0]

Loss value	w:
First 208	-1.01
Last 0.76	4.56

Learning rate 0.01
w = [10.0, -10.0]

Loss value	w:
First 52	10.9
Last 6.76	-1.81

Neural Networks Parameters

Input weights

Learning Rate

Linear Neurons - Perceptrons

Linear neurons even when combined have limited use

Need more types of neurons

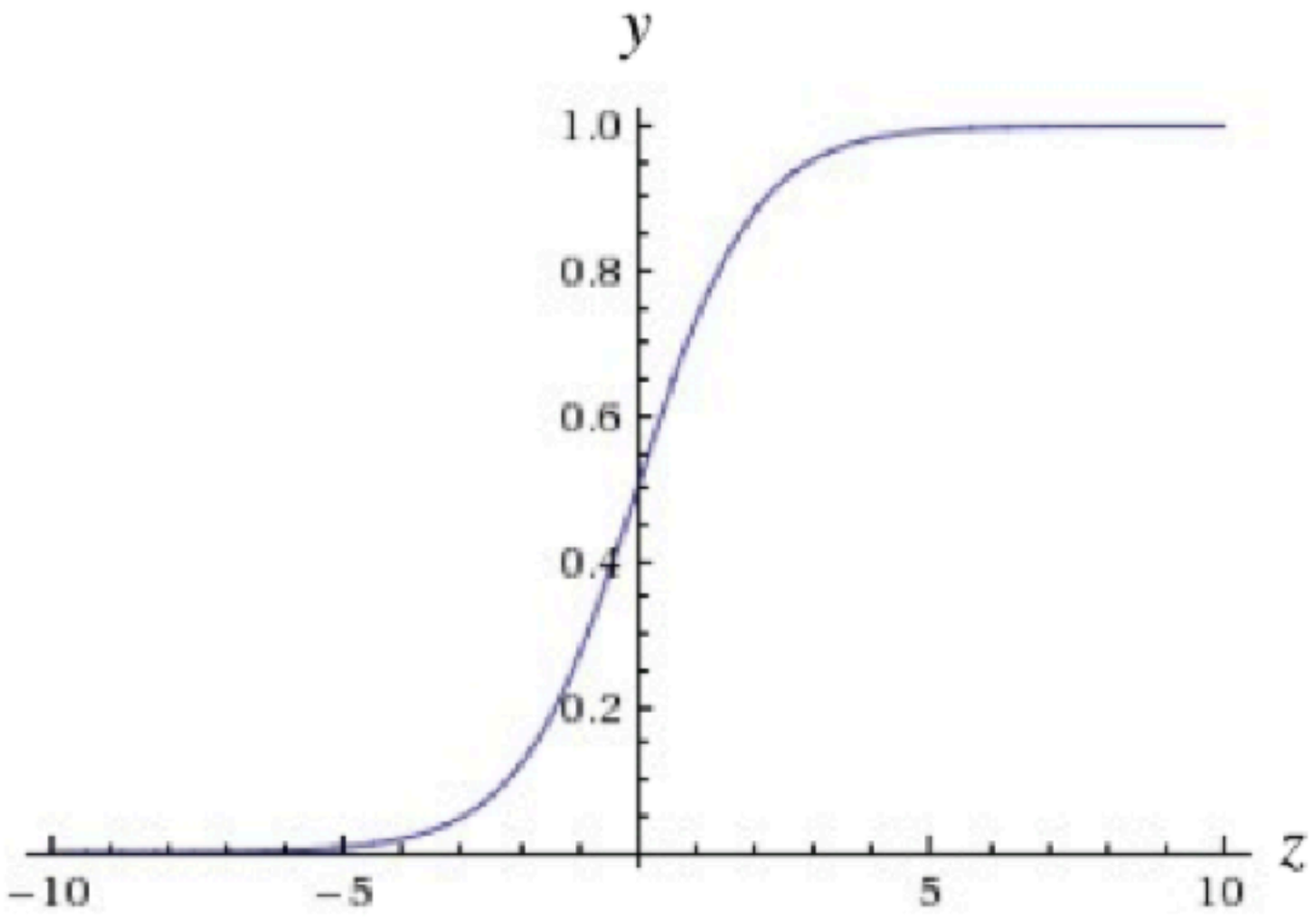
Each type needs gradient function & loss function

Layers of neurons

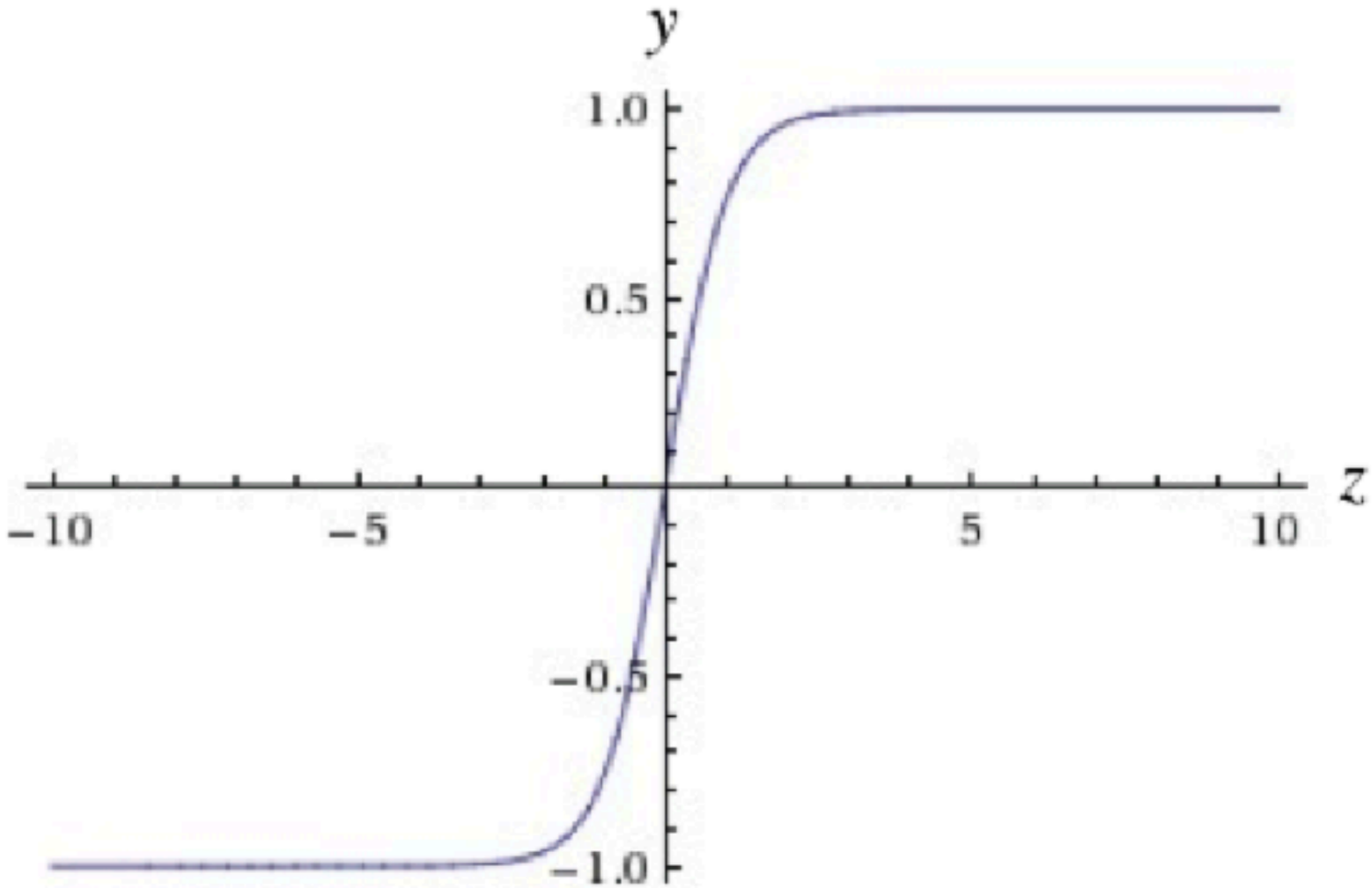
Types of Neurons/Activation Functions

Sigmoid

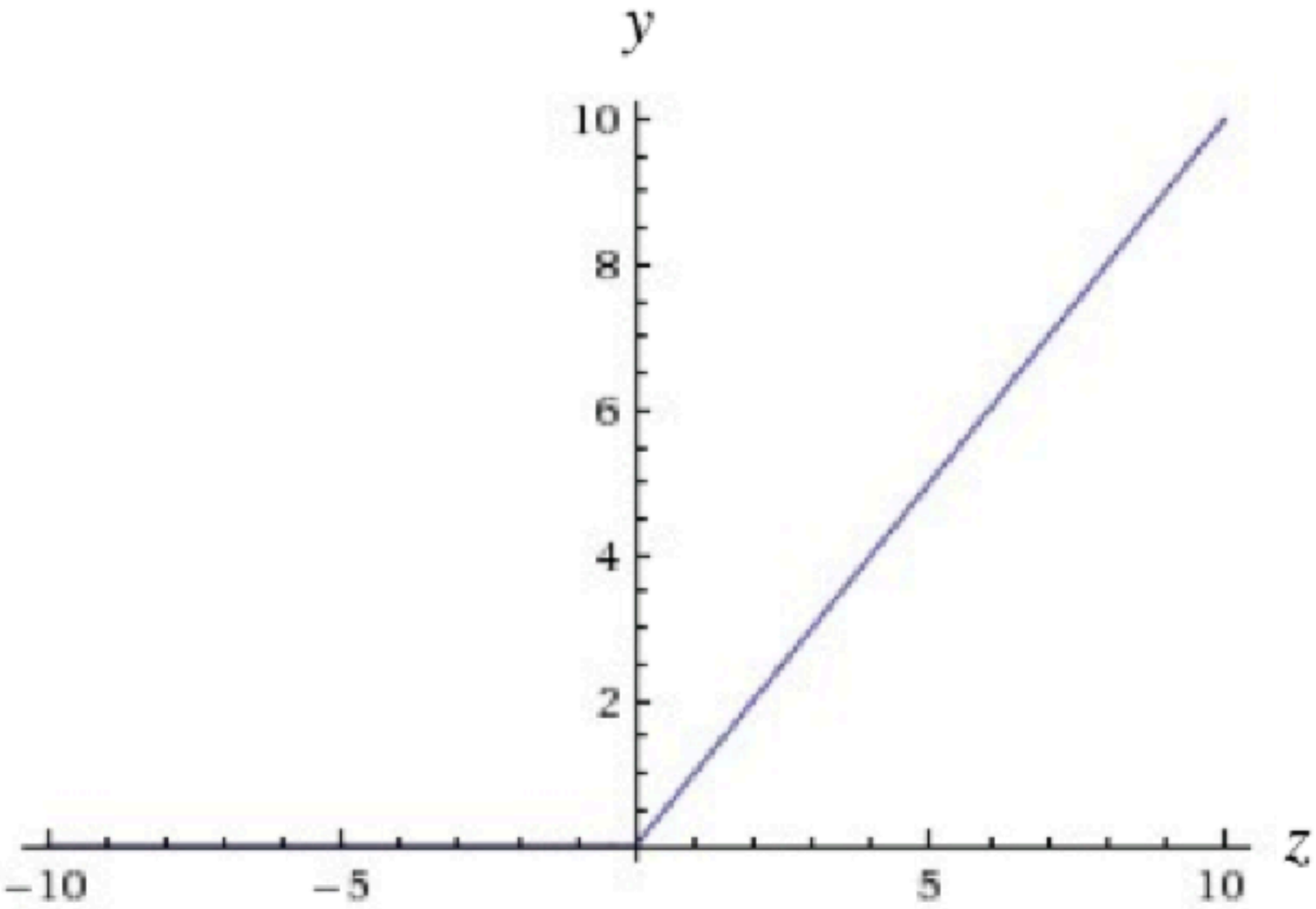
$$f(z) = \frac{1}{1 + e^{-z}}$$



Tanh



Restricted Linear Unit (ReLU)



Softmax

softmax_norm(x) = 1 ./(1 + exp(-(x - mean(x))/std(x)))

Recall from clustering

Often used as output neuron

Loss functions

$$\mathfrak{L}(\mathbf{W}, \boldsymbol{b}) = \frac{1}{N}\sum_{i=1}^{N}\sum_{j=1}^{M}(\log \hat{y_{ij}} - \log y_{ij})^2$$

mean square log error
MSLE

$$\mathfrak{L}(\mathbf{W}, \boldsymbol{b}) = \frac{1}{N}\sum_{i=1}^{N} max(0, 1 - y_{ij} \times \hat{y_{ij}})$$

Hinge loss
Binary classification

$$\mathfrak{L}(\mathbf{W}, \boldsymbol{b}) = -\sum_{i=1}^{N}\sum_{j=1}^{M} y_{ij} \times \log \hat{y}_{ij}$$

Logisitic loss

Neural Networks Parameters

Input weights

Learning Rate

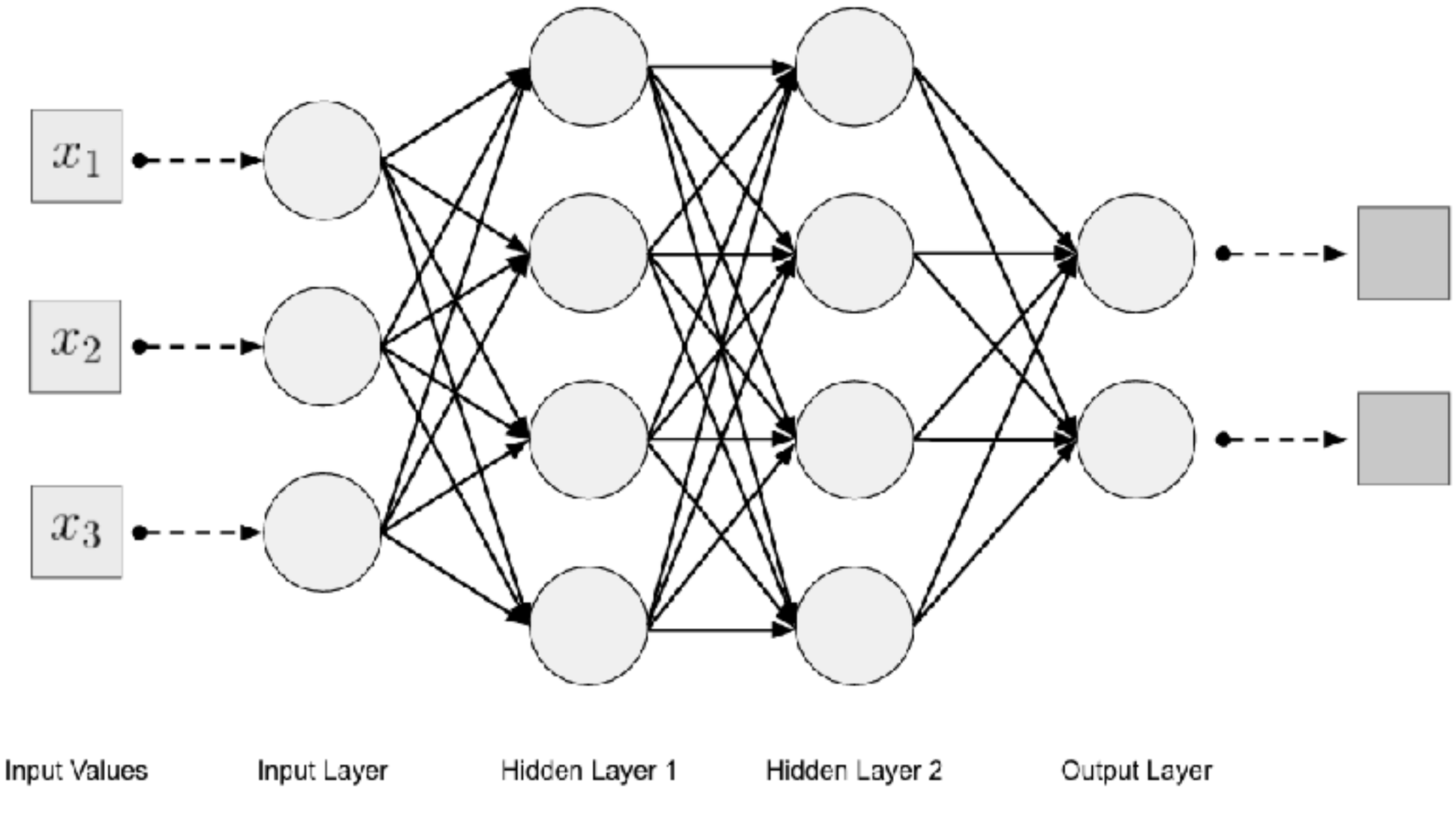
Loss function

Activation function

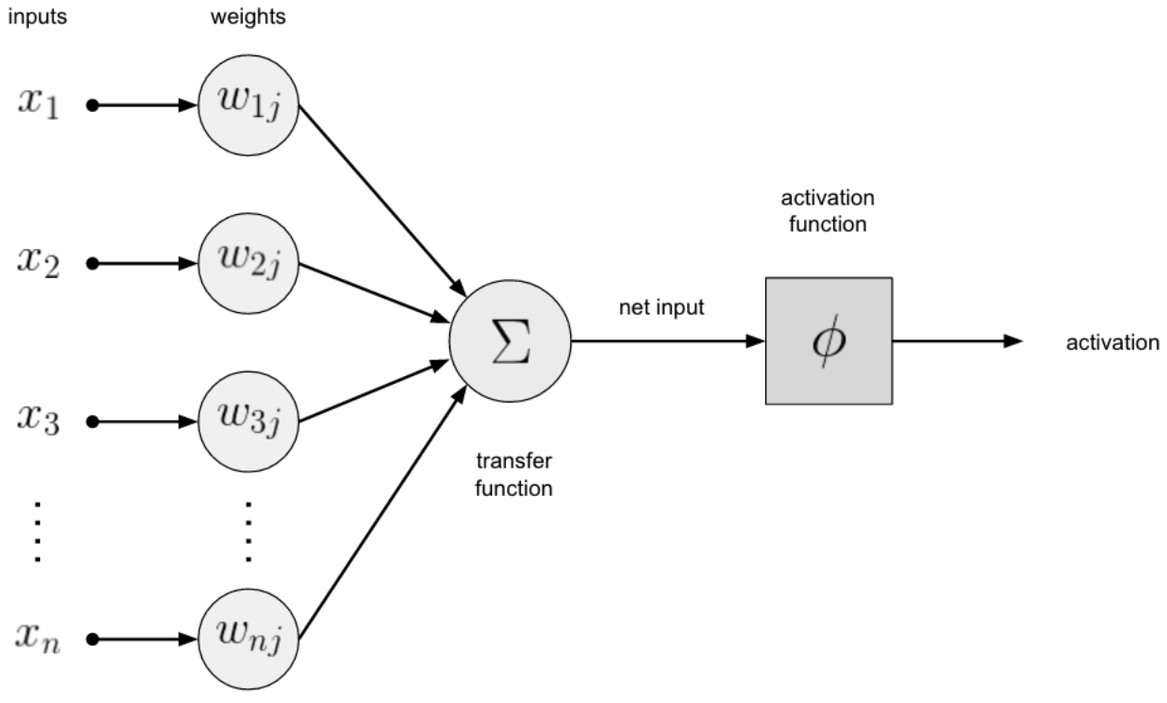
Layers

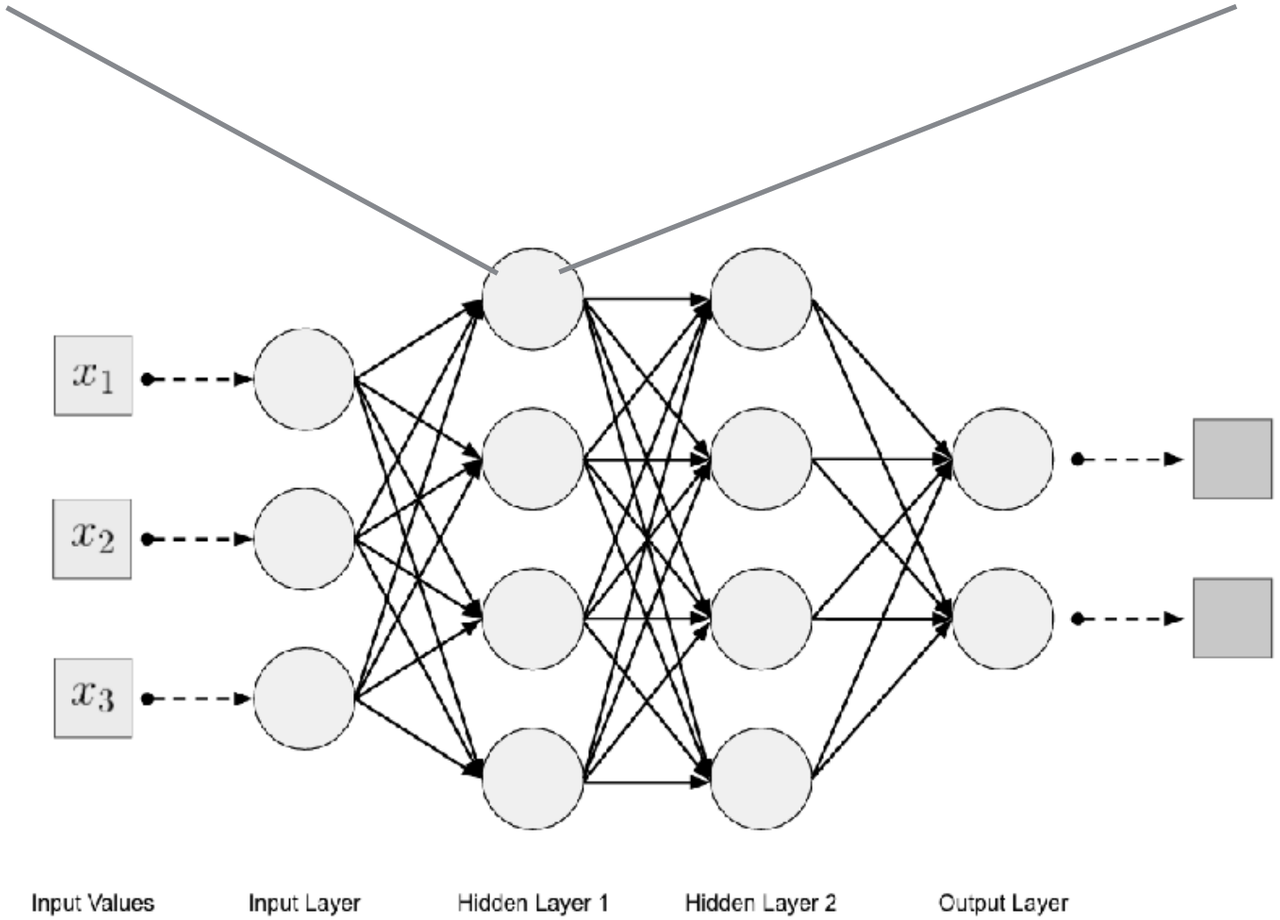
Even with different types of neurons single neurons are not very useful

Create layers of neurons



One neuron


inputs
weights
x_1
x_2
x_3
x_n
w_{1j}
w_{2j}
w_{3j}
w_{nj}
Σ
transfer function
net input
activation function
ϕ
activation

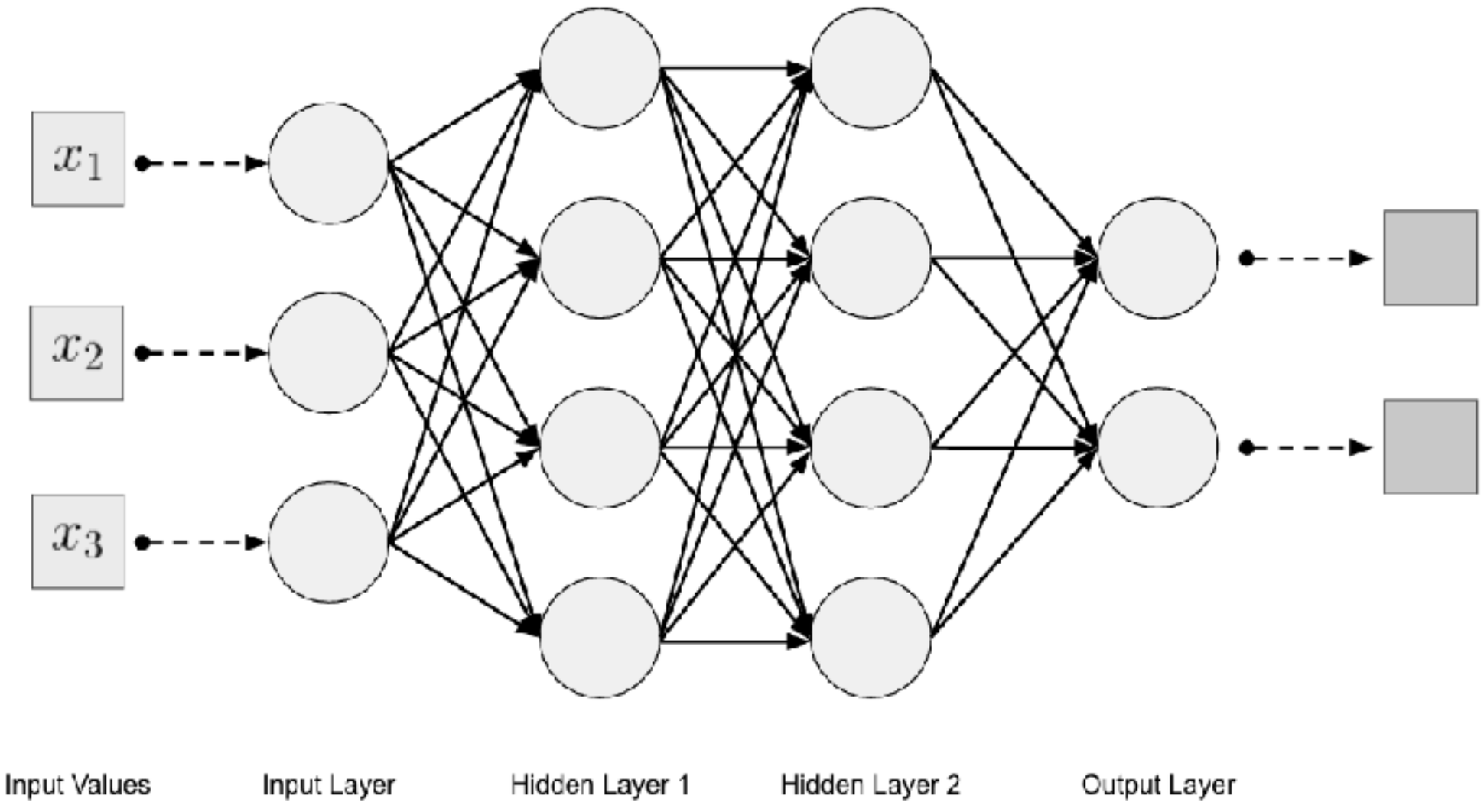

x_1
x_2
x_3
Input Values
Input Layer
Hidden Layer 1
Hidden Layer 2
Output Layer

Forward Propagation

Input data goes to input layer

Each neuron passes its output to the next layer

Below is a fully connected neural network



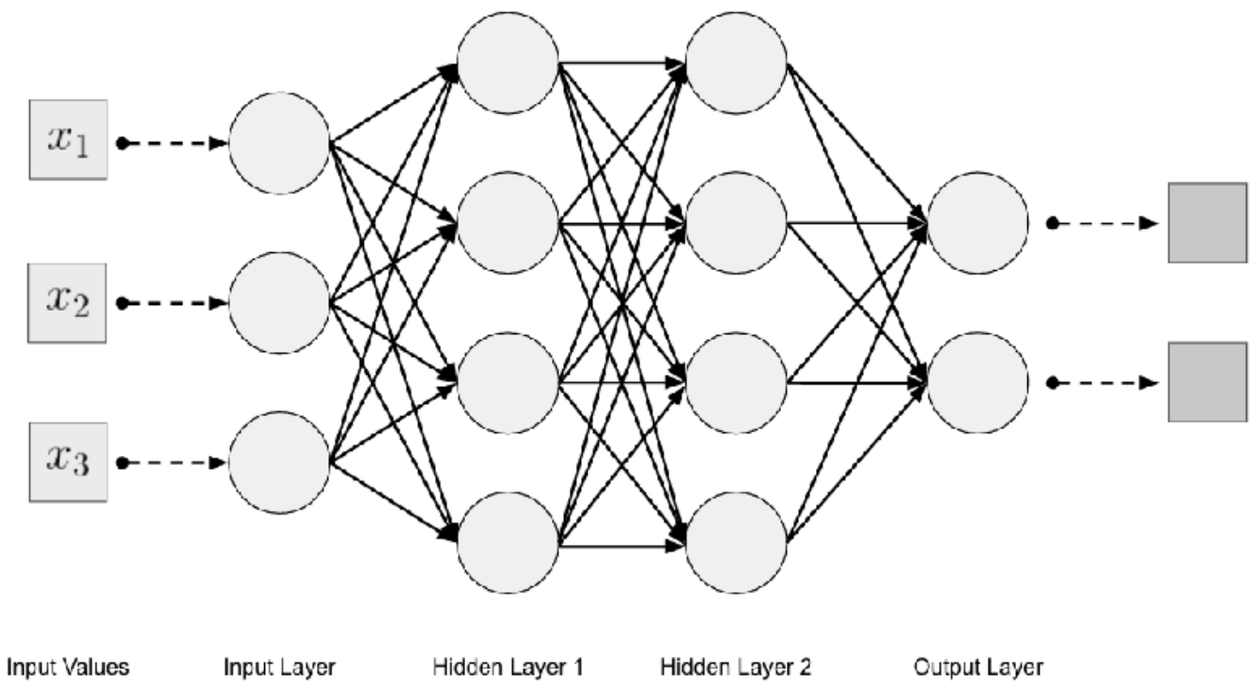
Back Propagation

How to adjust weights for each neuron?

Adjust the weights of the last layer as before

Using these weights we can compute what the inputs to last layer should be

We can now use those estimates to adjust the previous layers weights



Neural Networks Parameters

Input weights per neuron

Learning rate per neuron

Loss function per neuron

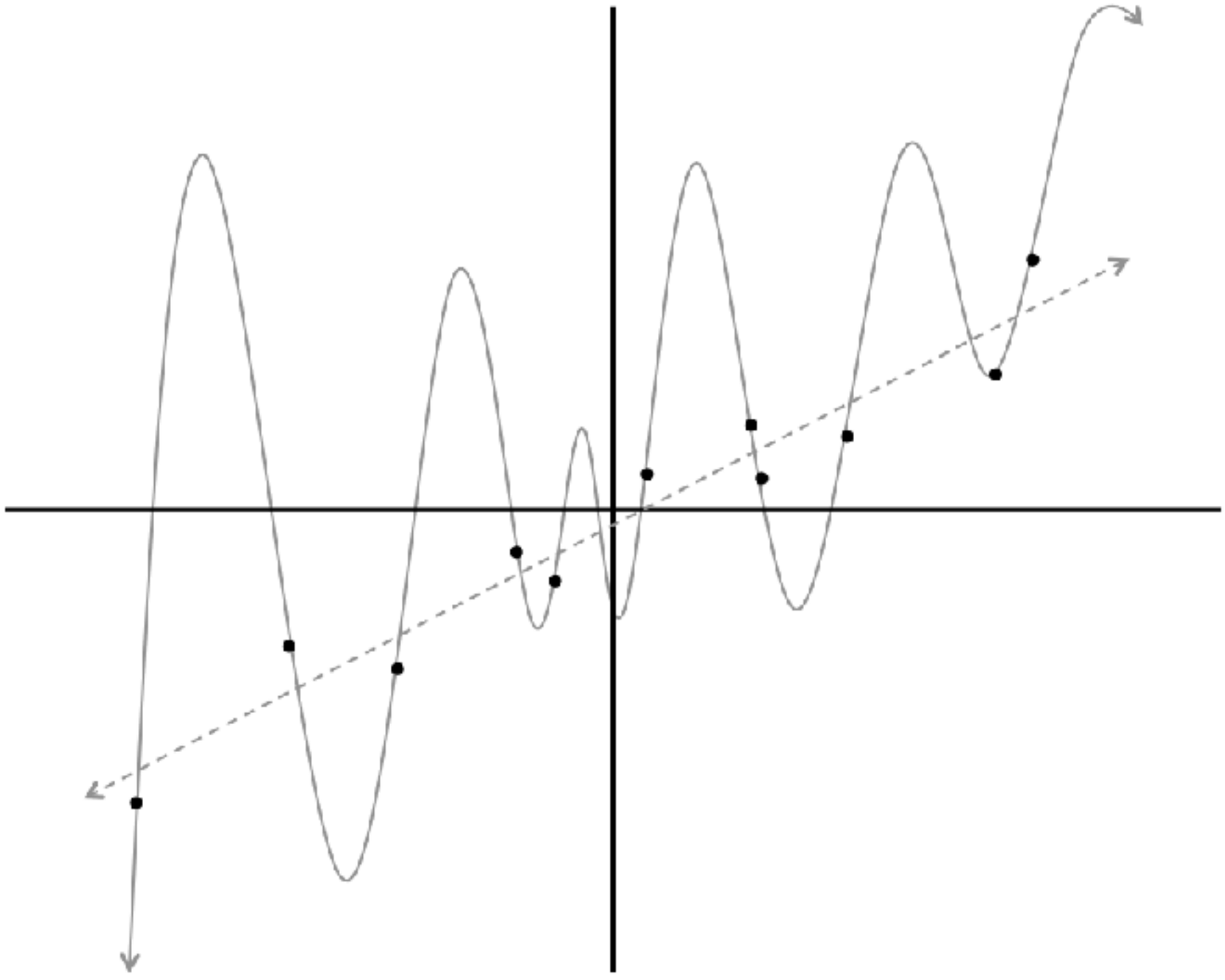
Activation function per neuron

Number of layers

Number of neurons per layer

How neurons are connected

Overfitting



Hyperparameters

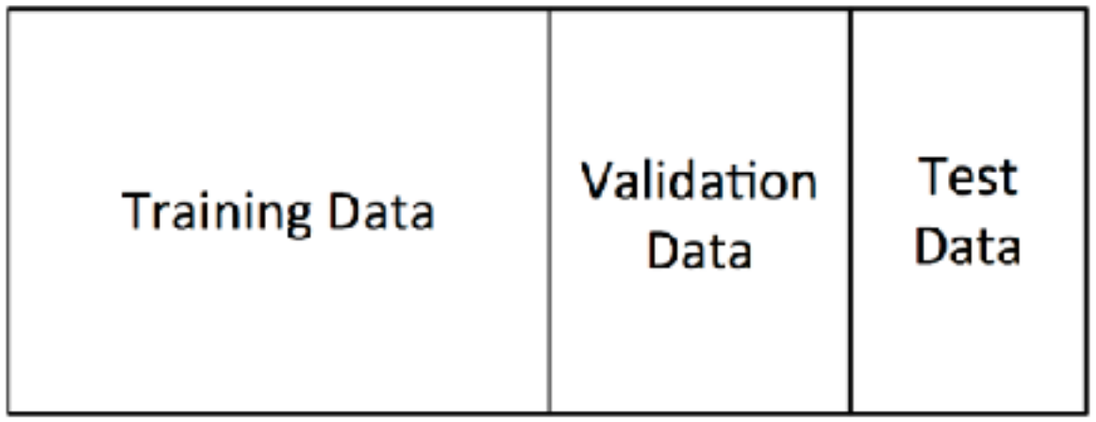
Things we can change to make neural networks train better

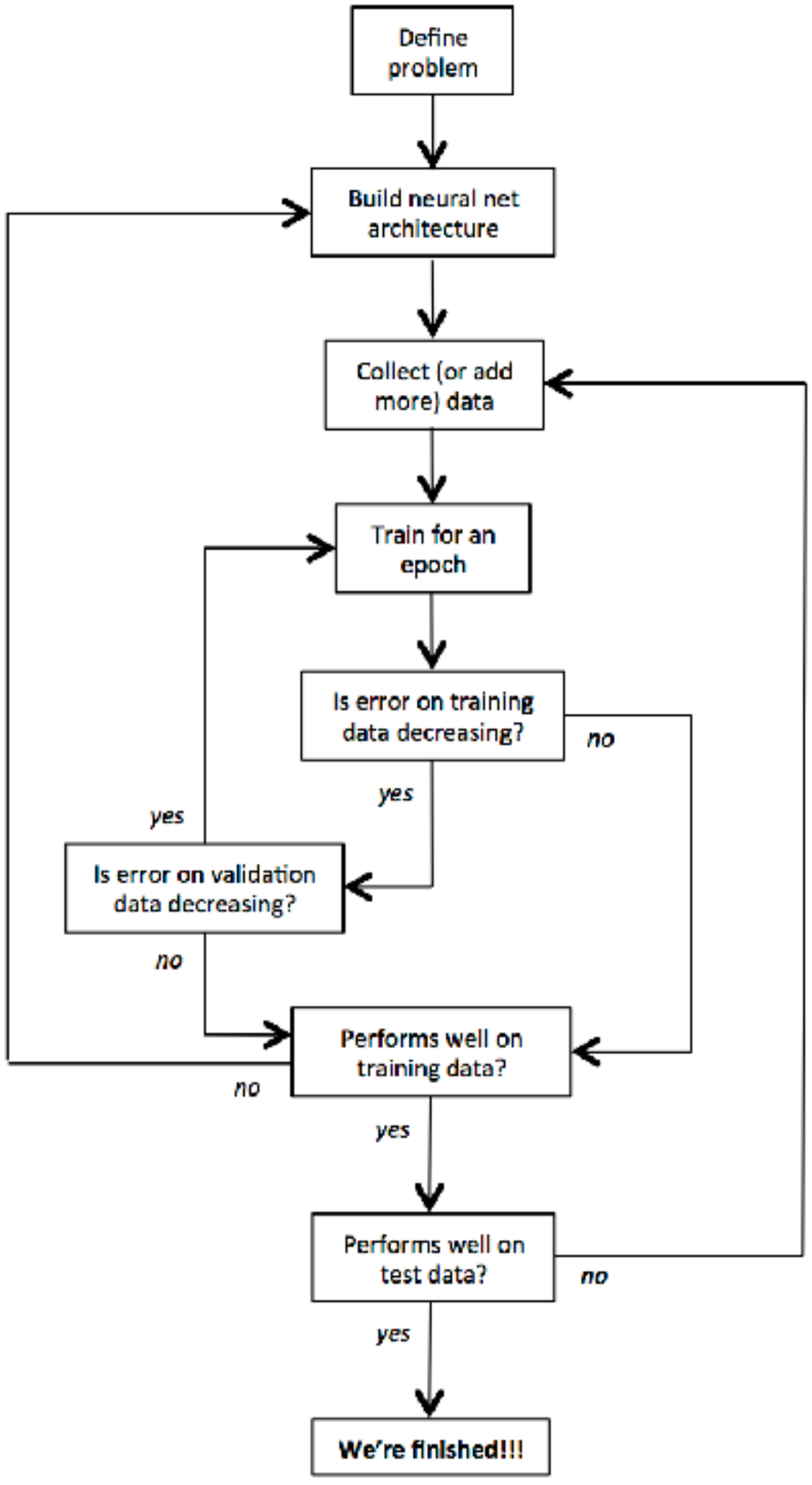
Learning Rate
Activation functions
Weight initalization strategies
Loss functions
Normalization
Layer size & number of layers

mini-batch size
Regularization
Momentum
Sparsity

Work Flow

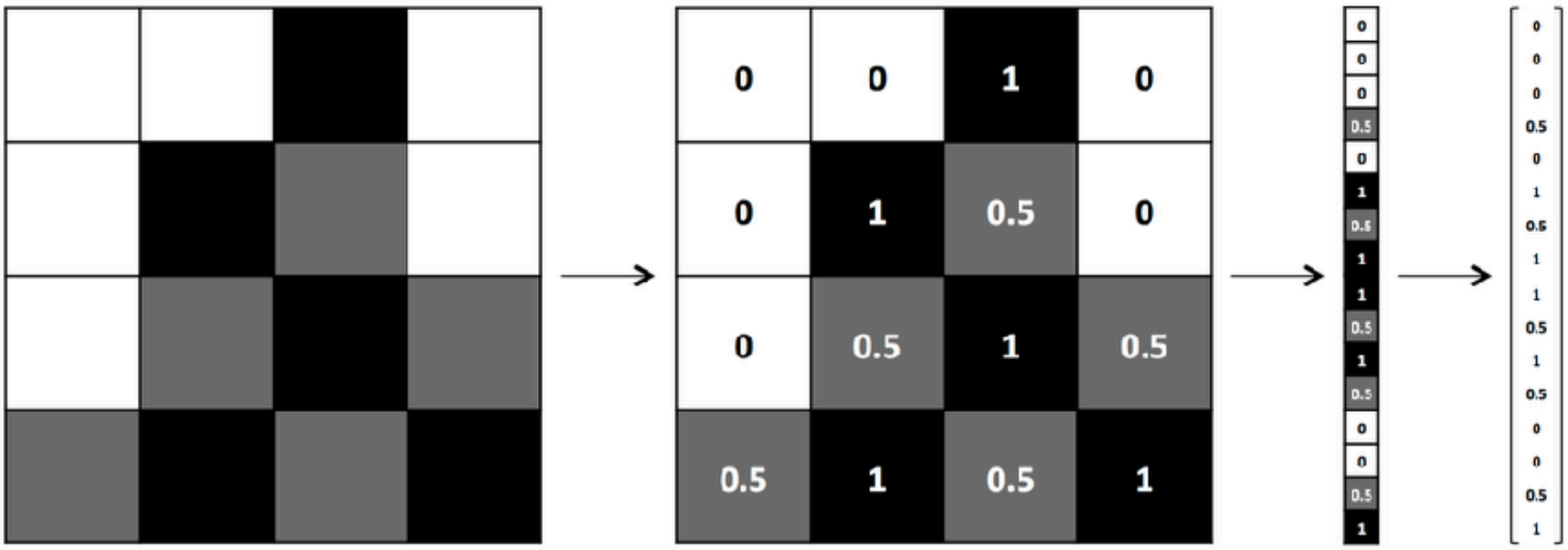
Full Dataset:





Input

Need to map input into vector



Images & Scaling

Image of 32 pixels by 32 pixels with 3 color channels (RGB)

Fully connected neuron needs 32*32*3 = 3,072 weights

Image of 200 pixels by 200 pixels with 3 color channels (RGB)

Fully connected neuron needs 200*200*3 = 120,000 weights

Image researchers use up to 150 layers

Deep Learning

More neurons than previous networks
More complex ways of connecting layers
Explosion of computing power to train
Automatic feature extraction

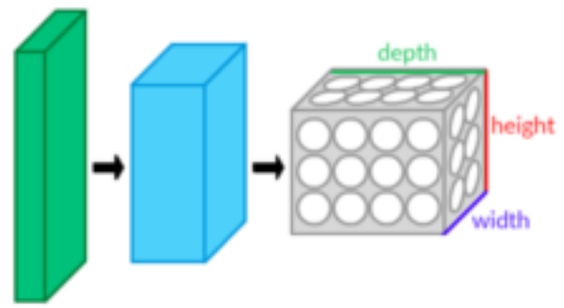
Some Deep Learning Networks

Unsupervised Pre-Trained Networks
Convolutional Neural Networks
  Common for image Analysis
Recurrent Neural Networks
  Time series analysis
Recursive Neural Networks

Convolutional Neural Network

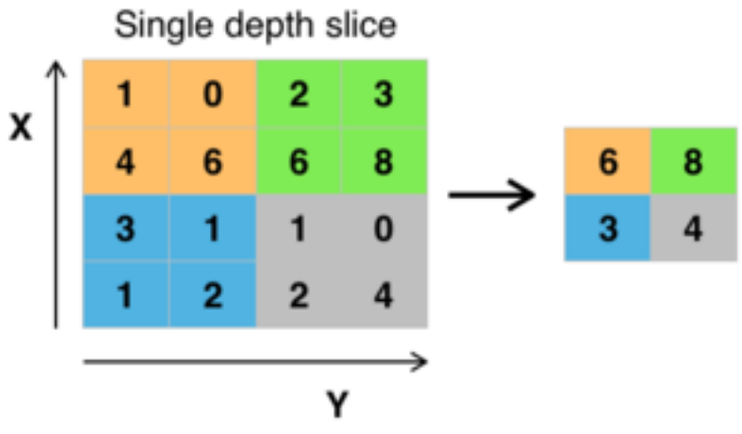
Convolutional Layer

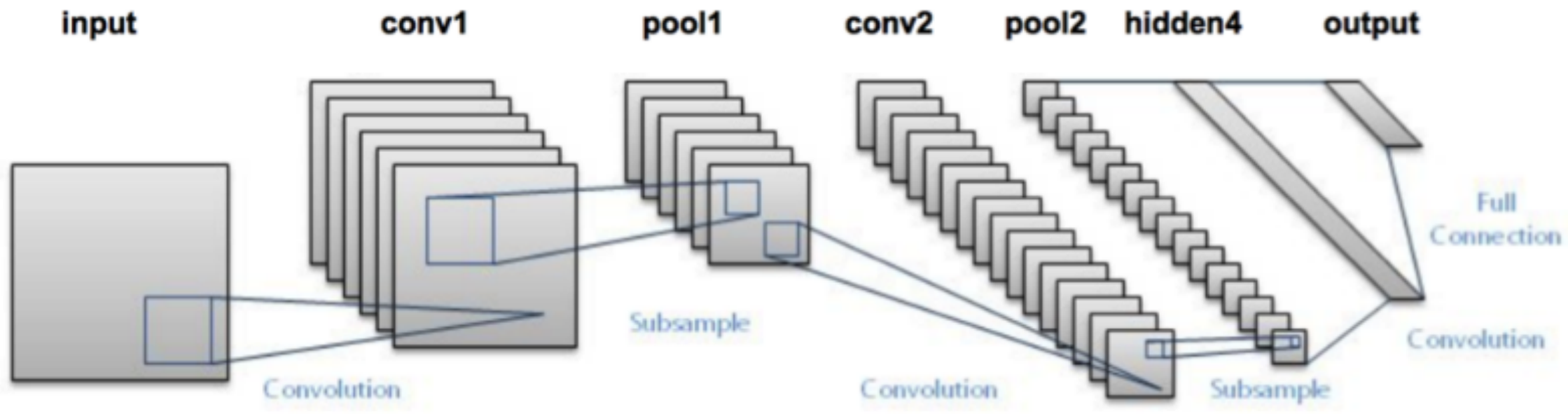
- 3-D network of neurons
- Only locally connected
- Each 2-D slice in depth share same weight



Pooling Layer

- Down-sampling layer




input
conv1
pool1
conv2
pool2
hidden4
output
Full
Connection
Convolution
Subsample
Convolution
Convolution
Subsample

Hello World of Deep Learning

Mixed National Institute of Standards & Technology database of handwritten digits

60,000 training images

Normalized to 20x20 pixels with grayscale

Different Methods with Error Rate

Type	Classifier	Error rate (%)
Linear classifier	Pairwise linear classifier	7.6
K-Nearest Neighbors	K-NN with non-linear deformation (P2DHMDM)	0.52
Neural network	2-layer 784-800-10	1.6
Neural network	2-layer 784-800-10	0.7
Deep neural network	6-layer 784-2500-2000-1500-1000-500-10	0.35
Convolutional neural network	Committee of 35 conv. net, 1-20-P-40-P-150-10	0.23

Spark DeepLearning

MultilayerPerceptronClassifier

feedforward artificial neural network

backpropagation for learning the model

Parameters

Number of Layers
Neurons per layer
Tolerance of iteration
Block size of the learning
Seed size
Max iteration number

```
iris = spark.read.format("csv"). \
        option("header",True).\
        option("inferschema",True).\
        load("iris.txt")

from pyspark.ml.feature import StringIndexer
from pyspark.ml.feature import VectorAssembler
from pyspark.ml import Pipeline

iris_indexer = StringIndexer(inputCol="species", outputCol="label").fit(iris)

iris_assembler = VectorAssembler(inputCols=["sepal_length","sepal_width", "petal_length",
"petal_width"], outputCol="features")

pipeline = Pipeline(stages=[iris_indexer, iris_assembler])
iris_formated = pipeline.fit(iris).transform(iris)

from pyspark.ml.classification import MultilayerPerceptronClassifier
from pyspark.ml.evaluation import MulticlassClassificationEvaluator

(train, test) = iris_formated.randomSplit([0.6, 0.4], 1234)
```

```
# specify layers for the neural network:
# input layer of size 4 (features), two intermediate of size 5 and 4
# and output of size 3 (classes)
layers = [4, 5, 4, 3]

# create the trainer and set its parameters
trainer = MultilayerPerceptronClassifier(maxIter=100, layers=layers, blockSize=128, seed=1234)

# train the model
model = trainer.fit(train)

result = model.transform(test)
predictionAndLabels = result.select("prediction", "label")
evaluator = MulticlassClassificationEvaluator(metricName="accuracy")
print("Test set accuracy = " + str(evaluator.evaluate(predictionAndLabels)))
```

```
Test set accuracy = 0.9607843137254902
```

```
result \
    .filter(result.label != result.prediction) \
    .select("label","rawPrediction","probability", "prediction") \
    .show()
```

```
+-----+--------------------+--------------------+----------+
|label|       rawPrediction|         probability|prediction|
+-----+--------------------+--------------------+----------+
|  1.0|[38.1539357915169...|[0.99999999999719...|       0.0|
|  1.0|[37.2094508219486...|[0.99999999999999...|       0.0|
+-----+--------------------+--------------------+----------+
```