#### **HUDM 6026: Computational Statistics**

#### **Artificial Neural Networks**

#### References

- *The Elements of Statistical Learning* (2009), by T. Hastie, R. Tibshirani, and J. Friedman.
- *Machine Learning: A Probabilistic Perspective* (2012), by K. Murphy

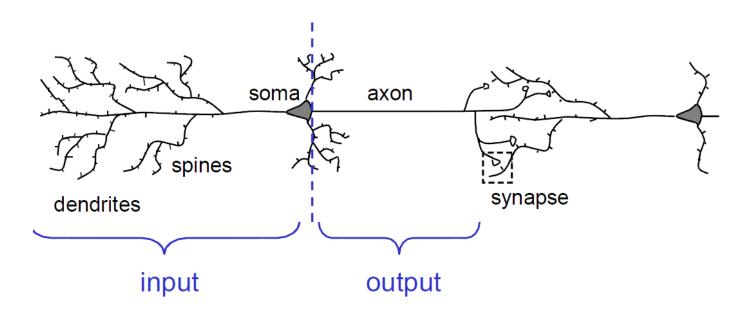
#### **Lesson Goals:**

• Understand artificial neural networks and the back-propagation algorithm for both regression and classification problems.

#### **Artificial Neural Networks**

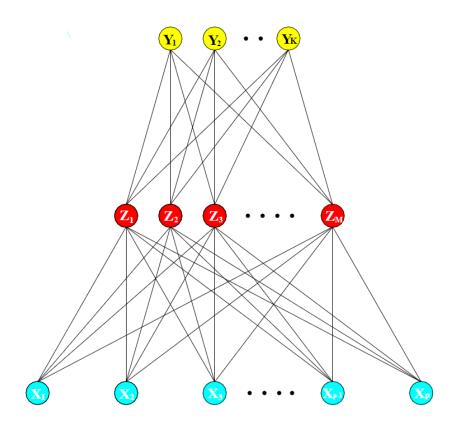
- The central idea of *artificial neural networks* (ANN) is to extract linear combinations of the inputs as derived features, and then model the target as a nonlinear function of these features.
- This machine learning method originated as an algorithm trying to mimic neurons in the brain.
- The brain has extraordinary computational power to represent and interpret complex environments.
- ANN attempts to capture this mode of computation.

• Specifically, ANN is a formalism for representing functions, inspired from biological systems and composed of parallel computing units, each computing a simple function.



Parts of the neuron: dendrites, soma (body), axon, synapses

- We will describe the most widely used ANN, the single layer perceptron (single hidden layer back-propagation network).
- A neural network is a two-stage regression or classification model, typically represented by a network diagram.



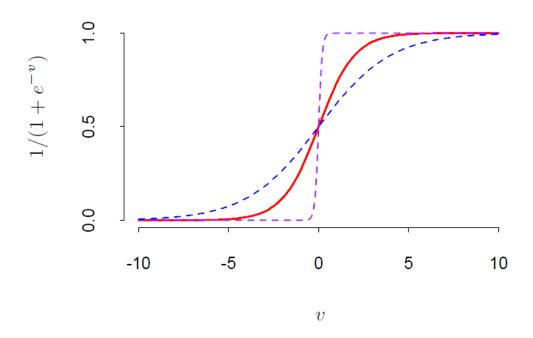
- In regression, typically K = 1 and there is only one output unit  $Y_1$  at the top of the network diagram.
- These networks, however, can handle multiple quantitative responses in a seamless fashion (multivariate regression).
- For K-class classification, there are K units at the top of the network diagram, with the k<sup>th</sup> unit modeling the probability of class k.
- There are K target measurements  $Y_k$ , k = 1, ..., K, each being coded as a 0 1 variable for the k<sup>th</sup> class.

• Derived features  $Z_m$  are created from linear combinations of the inputs X, and then the target  $Y_k$  is modeled as a function of the linear combinations of the  $Z_m$ :

$$Z_{m} = \sigma(\alpha_{0m} + \alpha_{m}^{T}X), \ m = 1, \dots, M,$$
$$T_{k} = \beta_{0k} + \beta_{k}^{T}Z, \ k = 1, \dots, K,$$
$$f_{k}(X) = g_{k}(T), \ k = 1, \dots, K,$$

where 
$$Z = (Z_1, Z_2, ..., Z_M)$$
 and  $T = (T_1, T_2, ..., T_K)$ 

• The activation function  $\sigma(v)$  is typically the sigmoid (logistic) function:  $\frac{1}{1+e^{-v}}$ 



**FIGURE 11.3.** Plot of the sigmoid function  $\sigma(v) = 1/(1 + \exp(-v))$  (red curve), commonly used in the hidden layer of a neural network. Included are  $\sigma(sv)$  for  $s = \frac{1}{2}$  (blue curve) and s = 10 (purple curve). The scale parameter s controls the activation rate, and we can see that large s amounts to a hard activation at v = 0. Note that  $\sigma(s(v - v_0))$  shifts the activation threshold from 0 to  $v_0$ .

- Neural network diagrams are sometimes drawn with an additional *bias* unit feeding into every unit in the hidden and output layers.
- Thinking of the constant "1" as an additional input feature, this bias unit captures the intercepts  $\alpha_{0m}$  and  $\beta_{0k}$  in the model.
- The output function  $g_k(T)$  allows a final transformation of the vector of outputs T.
- For regression, we typically choose the identity function  $g_k(T) = T_K$

• Early work in *K*-class classification also used the identity function, but this was later abandoned in favor of the *softmax* function:

$$g_k(T) = \frac{e^{T_k}}{\sum_{\ell=1}^K e^{T_\ell}}$$

- Note that this is exactly the transformation used in the multilogit model, which produces positive estimates that sum to one.
- The units in the middle of the network, computing the derived features  $Z_m$ , are called *hidden units* because these values are not directly observed.

- We can think of the  $Z_m$  as a basis expansion of the original inputs X; the neural network is then a standard linear model, or linear multilogit model, using these transformations as inputs.
- However, the parameters of the basis functions are learned from the data, which is an important enhancement over the basis expansion techniques.
- Notice that if  $\sigma$  is the identity function, then the entire model collapses to a linear model in the inputs.

- Thus, a neural network can be thought of as a nonlinear generalization of the linear model, both for regression and classification.
- By introducing the nonlinear transformation  $\sigma$ , it greatly enlarges the class of linear models.
- In general, there can be more than one hidden layers.
- The neural network model has unknown parameters, often called *weights*.

- We seek values for the weights that make the model fit the training data well.
- We denote the complete set of weights by  $\theta$ , which consists of:

$$\{\alpha_{0m}, \alpha_m; m = 1, 2, ..., M\}$$
  $M(p+1)$  weights,  
 $\{\beta_{0k}, \beta_k; k = 1, 2, ..., K\}$   $K(M+1)$  weights.

• For regression, we use RSS as our measure of fit (error function):

$$R(\theta) = \sum_{k=1}^{K} \sum_{i=1}^{N} (y_{ik} - f_k(x_i))^2.$$

• For classification, we use squared error or cross-entropy (deviance):

$$R(\theta) = -\sum_{i=1}^{N} \sum_{k=1}^{K} y_{ik} \log f_k(x_i),$$

and the corresponding classifier  $G(x) = \operatorname{argmax}_k f_k(x)$ .

• With the softmax activation function and the cross-entropy error function, the neural network model is exactly a linear logistic regression model in the hidden units, and all the parameters are estimated by maximum likelihood.

- Typically, we do not want the global minimizer  $R(\theta)$ , as this is likely to be an overfit solution.
- Instead, some regularization is need; this is achieved directly through a penalty term, or indirectly by early stopping.
- The generic approach to minimizing  $R(\theta)$  is by gradient descent, called *back-propagation* in this setting.
- The gradient can be easily derived using the chain rule for differentiation.

- This can be computed by a forward and backward sweep over the network, keeping track only of quantities local to each unit.
- Here is back-propagation in detail for squared error loss.
- Let  $z_{mi} = \sigma(\alpha_{0m} + \alpha_m^T x_i)$  and  $z_i = (z_{1i}, z_{2i}, ..., z_{Mi})$ . Then:

$$R(\theta) \equiv \sum_{i=1}^{N} R_i$$

$$= \sum_{i=1}^{N} \sum_{k=1}^{K} (y_{ik} - f_k(x_i))^2,$$

• With the following derivatives:

$$\frac{\partial R_i}{\partial \beta_{km}} = -2(y_{ik} - f_k(x_i))g_k'(\beta_k^T z_i)z_{mi},$$

$$\frac{\partial R_i}{\partial \alpha_{m\ell}} = -\sum_{k=1}^K 2(y_{ik} - f_k(x_i))g_k'(\beta_k^T z_i)\beta_{km}\sigma'(\alpha_m^T x_i)x_{i\ell}.$$

• Given these derivatives, a gradient descent update at the (r + 1)st iteration has the form (where  $\gamma_r$  is the *learning rate*):

$$\beta_{km}^{(r+1)} = \beta_{km}^{(r)} - \gamma_r \sum_{i=1}^{N} \frac{\partial R_i}{\partial \beta_{km}^{(r)}},$$
$$\alpha_{m\ell}^{(r+1)} = \alpha_{m\ell}^{(r)} - \gamma_r \sum_{i=1}^{N} \frac{\partial R_i}{\partial \alpha_{m\ell}^{(r)}},$$

• We can simplify the derivatives to:

$$\frac{\partial R_i}{\partial \beta_{km}} = \delta_{ki} z_{mi},$$
$$\frac{\partial R_i}{\partial \alpha_{m\ell}} = s_{mi} x_{i\ell}.$$

- The quantities  $\delta_{ki}$  and  $s_{mi}$  are "errors" from the current model at the output and hidden layer units, respectively. Note that the output layer errors  $\delta_{ki} = (\hat{f}_k(x_i) f_k(x_i))$
- From their definitions, these errors satisfy the *back-propagation* equations:

$$s_{mi} = \sigma'(\alpha_m^T x_i) \sum_{k=1} \beta_{km} \delta_{ki},$$

- Using this, the gradient descent updates can be implemented with a two-pass algorithm.
- In the forward pass, the current weights are fixed and the predicted values  $\hat{f}_k(x_i)$  are computed.
- In the *backward pass*, the errors  $\delta_{ki}$  are computed, and then backpropagated via the back-propagation equations to give the errors  $s_{mi}$
- Both sets of errors are then used to compute the gradients for the updates.

- This two-pass procedure is what is known as back-propagation.
- It has also been called the *delta rule*.
- The computational components for cross-entropy have the same form as those for the sum of squares function.
- The advantage of back-propagation are its simple, local nature. In the back-propagation algorithm, each hidden unit passes and receives information only to and from units that share a connection.

- Usually starting values for weights are chosen to be random values near zero.
- Hence, the model starts out nearly linear and becomes nonlinear as the weights increase.
- Use of exact zero weights leads to zero derivatives and perfect symmetry, and the algorithm never moves.
- Starting instead with large weights often leads to poor solutions.

- Often neural networks have too many weights and will overfit the data at the global minimum of *R*.
- An explicit method for regularization is *weight decay*, which is analogous to ridge regression used for linear models.
- We add a penalty to the error function  $R(\theta) + \lambda J(\theta)$ , where:

$$J(\theta) = \sum_{km} \beta_{km}^2 + \sum_{m\ell} \alpha_{m\ell}^2$$

and  $\lambda$  is a tuning parameter.

- Larger values of  $\lambda$  will tend to shrink the weights toward zero; typically cross-validation is used to estimate  $\lambda$ .
- Other forms for the penalty (such as weight elimination) are:

$$J(\theta) = \sum_{km} \frac{\beta_{km}^2}{1 + \beta_{km}^2} + \sum_{m\ell} \frac{\alpha_{m\ell}^2}{1 + \alpha_{m\ell}^2},$$

• Also, since the scaling of the inputs determines the effective scaling of the weights in the bottom layer, it can have a large effect on the quality of the final solution; thus, it is best to standardize all inputs.

- Generally, it is better to have too many hidden units than too few.
- With too few hidden units, the model might not have enough flexibility to capture the nonlinearities in the data.
- With too many hidden units, the extra weights can be shrunk toward zero if appropriate regularization is used.
- Typically the number of hidden units is somewhere in the range of 5 to 100, with the number increasing with the number of inputs and number of training cases.

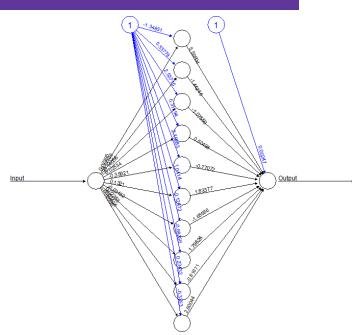
- The error function is nonconvex, possessing many local minima, so one must try a number of random starting configurations and the choose the solution giving the lowest error.
- Probably a better approach is to use the average predictions over the collection of networks as the final prediction.
- Another approach is via *bagging*, which averages the predictions of networks training from randomly perturbed version of the training data.

## ANN – Example 1 R Code

- In this first example, we use the R library "neuralnet" to train and build a neural network that is able to take a number and calculate the square root.
- The ANN will take a single input (the number you want square rooting) and produce a single output (the square root of the input).
- The ANN will contain 10 hidden neurons to be trained within each layer.
- The ANN does a reasonable job at finding the square root!

# ANN – Example 1 R Code (cont.)

```
## Example 1: Create a ANN to perform square rooting
library(neuralnet) # For Neural Network
# Generate 50 random numbers uniformly distributed between 0 and 100
# and store them as a dataframe
traininginput <- as.data.frame(runif(50, min = 0, max = 100))
trainingoutput <- sqrt(traininginput)</pre>
# Column bind the data into one variable
trainingdata <- cbind(traininginput,trainingoutput)</pre>
colnames(trainingdata) <- c("Input","Output")</pre>
# Train the neural network
# Going to have 10 hidden lavers
# Threshold is a numeric value specifying the threshold for the partial
# derivatives of the error function as stopping criteria.
net.sqrt <- neuralnet(Output ~ Input, trainingdata, hidden = 10, threshold = 0.01)</pre>
print(net.sqrt)
# Plot the neural network
plot(net.sqrt)
# Test the neural network on some training data
testdata <- as.data.frame((1:10)^2) # generate some squared numbers
net.results <- compute(net.sqrt, testdata) # run them through the neural network</pre>
# Let's see what properties net.sqrt has
ls(net.results)
# Let's see the results
print(net.results$net.result)
# Let's display a better version of the results
cleanoutput <- cbind(testdata,sgrt(testdata), as.data.frame(net.results$net.result))</pre>
colnames(cleanoutput) <- c("Input", "Expected Output", "Neural Net Output")
print(cleanoutput)
```



	Input	Expected	Output	Neural Net Output
1	1		1	1.027993874
2	4		2	2.005192003
3	9		3	3.001567330
4	16		4	3.986023794
5	25		5	5.003264027
6	36		6	6.011652248
7	49		7	6.990255178
8	64		8	8.000067508
9	81		9	9.016304667
10	100		10	9.960483101

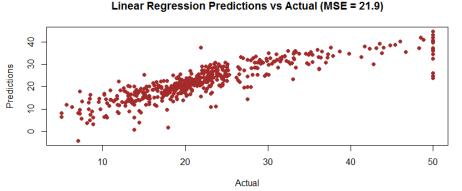
#### ANN – Example 2 R Code

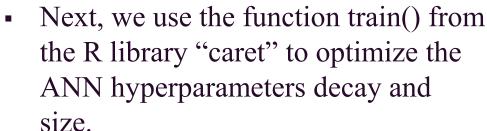
- In this second example, we use the R library "nnet" to train and build a neural network that is able to predict median value of owner-occupied homes.
- This example of regression with neural networks is compared with multiple linear regression.
- The data set is housing data for 506 census tracts of Boston from the 1970 census.
- We see that the ANN outperforms linear regression in terms of lower MSE.

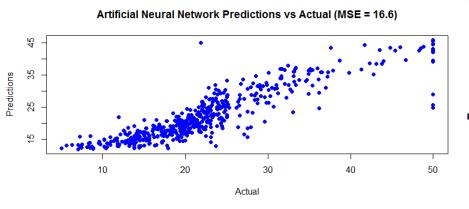
#### ANN – Example 2 R Code (cont.)

```
# Example 2: Create a ANN for prediction
# We give a brief example of regression with neural networks and comparison with
# multivariate linear regression. The data set is housing data for 506 census tracts of
# Boston from the 1970 census. The goal is to predict median value of owner-occupied homes.
# Load the data and insepct the range (which is 1 - 50)
library(mlbench)
data(BostonHousing)
summary(BostonHousing$medv)
# Build the multiple linear regression model
lm.fit <- lm(medv \sim ... data = BostonHousing)
lm.predict <- predict(lm.fit)</pre>
# Calculate the MSE and plot
mean((lm.predict - BostonHousing\$medv)^2) # MSE = 21.89483
par(mfrow = c(2,1))
plot(BostonHousing$medv, lm.predict, main = "Linear Regression Predictions vs Actual (MSE = 21.9)",
    xlab = "Actual", ylab = "Predictions", pch = 19, col = "brown")
# Build the feed-forward ANN (w/ one hidden layer)
library(nnet)
                    # For Neural Network
nnet.fit <- nnet(medv/50 ~ ., data=BostonHousing, size = 2) # scale inputs: divide by 50 to get 0-1 range
nnet.predict <- predict(nnet.fit)*50 # multiply 50 to restore original scale</pre>
# Calculate the MSE and plot
mean((nnet.predict - BostonHousing$medv)^2) # MSE = 16.56974
plot(BostonHousing$medv, nnet.predict, main = "Artificial Neural Network Predictions vs Actual (MSE = 16.6)".
    xlab = "Actual", ylab = "Predictions", pch = 19, col = "blue")
```

## ANN – Example 2 R Code (cont.)







- This library also performs resampling to give a better estimate of the error.
- Note that we scale linear regression by the same value, so the error statistics are directly comparable.

## ANN – Example 2 R Code (cont.)

- Tuned ANN RMSE = 0.0818
- Linear Regression RMSE = 0.0985

```
Neural Network
506 samples
13 predictors
No pre-processing
Resampling: Bootstrapped (25 reps)
Summary of sample sizes: 506, 506, 506, 506, 506, 506, ...
Resampling results across tuning parameters:
  decay size RMSE
                      Rsquared RMSE SD Rsquared SD
              0.083
                      0.8
                                0.00856 0.0378
  0.1
              0.0827 0.801
                                0.00894 0.0416
  0.1
              0.0818 0.806
                                0.00866 0.0382
  0.5
              0.0921 0.762
                                0.00904 0.0405
  0.5
              0.0908 0.77
                                0.00837 0.0386
              0.0901 0.771
RMSE was used to select the optimal model using the smallest value.
The final values used for the model were size = 6 and decay = 0.1.
```

```
Linear Regression

506 samples
13 predictors

No pre-processing
Resampling: Bootstrapped (25 reps)

Summary of sample sizes: 506, 506, 506, 506, 506, 506, ...

Resampling results

RMSE Rsquared RMSE SD Rsquared SD
0.0985 0.717 0.00911 0.0473
```

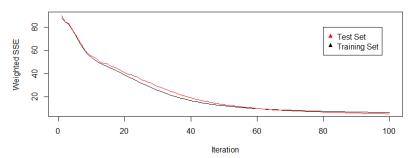
## ANN – Example 3 R Code

- In this third example, we use the R library "RSNNS" to train and build a neural network that is able to predict the identification of Iris plant Species.
- The ANN performs the classification on the basis of plant attribute measurements: Sepal Length, Sepal Width, Petal Length, Petal Width.
- This example of classification with neural networks uses the standard back-propagation algorithm.

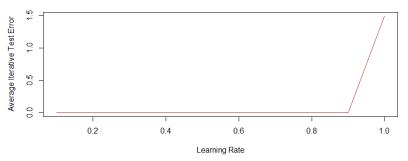
# ANN – Example 3 R Code (cont.)

```
# Example 3: Create a ANN for classification
# Predict the identification of Iris plant Species on the basis of plant
# attribute measurements: Sepal.Length, Sepal.Width, Petal.Length, Petal.Width
library(RSNNS)
# Load and store the 'iris' data
data(iris)
# Generate a sample from the 'iris' data set
irisSample <- iris[sample(1:nrow(iris), length(1:nrow(iris))), 1:ncol(iris)]
irisValues <- irisSample[, 1:4]
head(irisValues)
irisTargets <- irisSample[, 5]</pre>
head(irisTargets)
# Generate a binary matrix from an integer-valued input vector representing class labels
irisDecTargets <- decodeClassLabels(irisTargets)</pre>
head(irisDecTargets)
# Split the data into the training and testing set, and then normalize
irisSample <- splitForTrainingAndTest(irisValues, irisDecTargets, ratio = 0.15)
irisSample <- normTrainingAndTestSet(irisSample)
# Train the Neural Network (Multi-Layer Perceptron)
nn3 <-- mlp(iris5ample$inputsTrain, iris5ample$targetsTrain, size = 2, learnFuncParams = 0.1, maxit = 100,
 inputsTest = irisSample$inputsTest, targetsTest = irisSample$targetsTest)
print(nn3)
# Predict using the testing data
testPred6 <- predict(nn3, irisSample$inputsTest)
# Calculate the Confusion Matrices for the Training and Testing Sets
confusionMatrix(irisSample$targetsTrain, fitted.values(nn3))
confusionMatrix(irisSample$targetsTest, testPred6)
# Calculate the Weights of the Newly Trained Network
weightMatrix(nn3)
# Plot the Iterative Error of both training (black) and test (red) error
# This shows hows the Number of Iterations Affects the Weighted SSE
plotIterativeError(nn3, main = "# of Iterations vs. Weighted SSE")
legend(80, 80, legend = c("Test Set", "Training Set"), col = c("red", "black"), pch = 17)
# See how changing the Learning Rate Affects the Average Test Error
err <- vector(mode = "numeric", length = 10)
learnRate = seq(0.1, 1, length.out = 10)
for (i in 10){
 fit <- mlp(irisSample$inputsTrain, irisSample$targetsTrain, size = 2, learnFuncParams = learnRate[i], maxit = 50,
   inputsTest = irisSample$inputsTest, targetsTest = irisSample$targetsTest)
  err[i] <- mean(fit$IterativeTestError)
# Plot the Effect of Learning Rate vs. Average Iterative Test Error
plot learnRate, err, xlab = "Learning Rate", ylab = "Average Iterative Test Error",
  main = "Learning Rate vs. Average Test Error", type = "1", col = "brown")
```

#### # of Iterations vs. Weighted SSE



#### Learning Rate vs. Average Test Error



We can see the iterative error of both training and test sets, and we see how changing the learning rate affects the average test error!