Math 208: Creating Neural Network Functions

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```
library(palmerpenguins)
library(dplyr)
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
      filter, lag
## The following objects are masked from 'package:base':
##
##
      intersect, setdiff, setequal, union
library(neuralnet)
## Attaching package: 'neuralnet'
## The following object is masked from 'package:dplyr':
##
##
      compute
library(tidyverse)
## -- Attaching core tidyverse packages ---
                                            ----- tidyverse 2.0.0 --
## v forcats 1.0.0
                       v readr
                                    2.1.4
## v ggplot2 3.4.2
                                    1.5.0
                        v stringr
                     v tibble
## v lubridate 1.9.2
                                    3.2.1
## v purrr
              1.0.1
                        v tidyr
                                    1.3.0
## -- Conflicts ------ tidyverse_conflicts() --
## x neuralnet::compute() masks dplyr::compute()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                         masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
```

```
penguins_example <- penguins %>% drop_na %>% mutate(sex=ifelse(sex=="female",1,0))
```

Task 1:

write a function that requires three arguments: i. A data frame or tibble ii. A length 1 character vector indicating the name of the outcome column in the dataset. iii. A character vector of unspecified length containing the names of the input features to be selected and scaled. and returns a new data set which contains a tibble containing only the outcome vector which should be renamed outcome and the scaled feature vectors, each of which has been scaled using the scale function.

Solution:

```
outcome_func<-function(x_df, outcome, input_vec){</pre>
  outcome_1 <- x_df[,c(outcome, input_vec)] %>% mutate_at(~scale(.),.vars=vars(input_vec))
  names(outcome_1)[names(outcome_1)== outcome] <- "outcome"</pre>
  return(outcome_1)
  }
outcome_col<-c("sex")</pre>
z_vect_cols<-c("body_mass_g","bill_length_mm")</pre>
result_a <- outcome_func(penguins_example, outcome_col, z_vect_cols)
## Warning: Using an external vector in selections was deprecated in tidyselect 1.1.0.
## i Please use 'all_of()' or 'any_of()' instead.
##
     # Was:
##
     data %>% select(input_vec)
##
##
     # Now:
##
     data %>% select(all_of(input_vec))
##
## See <https://tidyselect.r-lib.org/reference/faq-external-vector.html>.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last lifecycle warnings()' to see where this warning was
## generated.
head(result_a)
```

```
## # A tibble: 6 x 3
##
     outcome body_mass_g[,1] bill_length_mm[,1]
##
       <dbl>
                        <dbl>
                                             <dbl>
## 1
           0
                       -0.568
                                            -0.895
## 2
           1
                       -0.506
                                            -0.822
                       -1.19
## 3
           1
                                            -0.675
## 4
           1
                       -0.940
                                            -1.33
## 5
           0
                       -0.692
                                            -0.858
## 6
                       -0.723
                                            -0.931
```

Task 2: Write a function to randomly split a data frame or tibble into Training and Test that requires two arguments: i. A data frame or tibble ii. The percentage of the total number of rows that should be from training and returns a list which has two elements, one that is the Training data and the other is the Test data.

Solution:

```
split_data<-function(x_df, percentage){</pre>
split_labels <- sample(c("Training", "Test"),</pre>
prob=c(percentage,1-percentage), replace=T,
 size=nrow(x_df))
Training_sample <- x_df %>%
filter(split labels=="Training")
Test_sample <- x_df %>%
filter(split_labels=="Test")
list(Training=Training_sample, Test=Test_sample)
result_b<-split_data(result_a,0.7)
glimpse(result_b)
## List of 2
## $ Training: tibble [231 x 3] (S3: tbl df/tbl/data.frame)
##
    ..$ outcome
                   : num [1:231] 0 1 1 1 1 0 1 1 0 1 ...
##
     ..$ body_mass_g : num [1:231, 1] -0.568 -0.506 -1.189 -0.94 -1.251 ...
     .. ..- attr(*, "scaled:center")= num 4207
##
     ....- attr(*, "scaled:scale")= num 805
##
     ..$ bill_length_mm: num [1:231, 1] -0.895 -0.822 -0.675 -1.334 -0.529 ...
##
     .. ..- attr(*, "scaled:center")= num 44
     ...- attr(*, "scaled:scale")= num 5.47
##
   $ Test
             : tibble [102 x 3] (S3: tbl_df/tbl/data.frame)
##
                       : num [1:102] 0 1 0 0 1 1 1 0 1 1 ...
##
    ..$ outcome
     ..$ body_mass_g : num [1:102, 1] -0.692 -0.723 0.581 -0.506 -0.506 ...
     .. ..- attr(*, "scaled:center")= num 4207
##
##
     .. ..- attr(*, "scaled:scale")= num 805
##
     ..$ bill_length_mm: num [1:102, 1] -0.858 -0.931 -0.876 -0.986 -1.59 ...
     .. ..- attr(*, "scaled:center")= num 44
##
     .. ..- attr(*, "scaled:scale")= num 5.47
##
head(result_b)
## $Training
## # A tibble: 231 x 3
      outcome body_mass_g[,1] bill_length_mm[,1]
        <dbl>
##
                       <dbl>
                                           <dbl>
## 1
           0
                       -0.568
                                          -0.895
## 2
            1
                       -0.506
                                          -0.822
## 3
            1
                       -1.19
                                          -0.675
## 4
                                          -1.33
                       -0.940
            1
## 5
            1
                                          -0.529
                       -1.25
## 6
            0
                       0.240
                                          -1.72
                                          -1.35
## 7
            1
                       -0.630
## 8
                       -0.940
                                          -0.968
            1
## 9
            0
                        0.364
                                          -0.273
            1
## 10
                       -1.10
                                          -1.75
## # i 221 more rows
##
```

```
## $Test
## # A tibble: 102 x 3
##
      outcome body_mass_g[,1] bill_length_mm[,1]
##
         <dbl>
                          <dbl>
                                                <dbl>
##
    1
             0
                         -0.692
                                              -0.858
##
    2
             1
                         -0.723
                                              -0.931
    3
             0
                          0.581
                                              -0.876
##
##
    4
             0
                         -0.506
                                              -0.986
##
    5
             1
                         -0.506
                                              -1.59
##
    6
             1
                         -1.25
                                              -0.639
##
    7
             1
                         -1.13
                                              -1.17
             0
                          0.550
                                              -0.767
##
    8
##
    9
             1
                         -1.37
                                              -1.46
                         -0.878
                                              -0.803
## 10
             1
## # i 92 more rows
```

Task 3:

Write a function that takes in the following arguments: i. A data frame or tibble with a column named outcome and other columns that are all scaled feature vectors ii. A vector of integers that can be used as the hidden argument to the neuralnet function, i.e. a list of numbers of nodes of the hidden layers of a neural network to return a neuralnet object that is the result of running the neuralnet function on the data frame/tibble with the hidden nodes specified from the second argument and the following other arguments: linear.output = FALSE, act.fct="logistic" and using the outcome variable as the outcome in the formula argument

```
fit_model<-function(x_df, hidden_nodes_vec){</pre>
neural_net = neuralnet(outcome~.,linear.output = FALSE, act.fct="logistic",data=x_df, hidden=hidden_no
return(neural_net)
outcome<-c("sex")
result_b[[1]]
  # A tibble: 231 x 3
##
##
      outcome body_mass_g[,1] bill_length_mm[,1]
##
        <dbl>
                          <dbl>
                                              <dbl>
##
    1
             0
                        -0.568
                                             -0.895
    2
##
             1
                        -0.506
                                             -0.822
##
    3
             1
                        -1.19
                                             -0.675
##
    4
             1
                        -0.940
                                             -1.33
##
    5
                        -1.25
                                             -0.529
             1
##
    6
             0
                         0.240
                                             -1.72
##
    7
             1
                        -0.630
                                             -1.35
##
    8
             1
                        -0.940
                                             -0.968
    9
             0
                         0.364
                                             -0.273
##
## 10
                        -1.10
                                             -1.75
             1
## # i 221 more rows
result_c<-fit_model(result_b[[1]], c(2,2))
```

Task 4:

plot(result_c)

Write a function that takes the following arguments: i. A neuralnet object ii. A data frame/tibble containing Training Data iii. A data frame/tibble containing Test Data and returns a vector containing the average training squared error and the average test squared error using the neuralnet object, where average squared error is as defined in the background section.

```
run_training_test<-function(model_obj, Training, Test){

train_predict <- predict(model_obj,newdata=Training)
Training_error<-Training %>% mutate(train_error_sq=(outcome-train_predict)^2) %>%
summarize(Avg_Error_train=mean(train_error_sq))

test_predict <- predict(model_obj,newdata=Test)
Test_error<-Test%>% mutate(test_error_sq=(outcome-test_predict)^2) %>%
summarize(Avg_Error_test=mean(test_error_sq))

c("Training_Error"=as.numeric(Training_error), "Test_Error"=as.numeric(Test_error))
}

run_training_test(result_c,result_b$Training,result_b$Test)
```

```
## Training_Error Test_Error
## 0.1045217 0.1227034
```

Task 5:

Write a function that takes the following arguments: i. A data frame or tibble ii. A length 1 character vector indicating the name of the outcome column in the dataset. iii. A character vector of unspecified length containing the names of the input features to be selected and scaled. iv. The percentage of the total number of rows in the data/frame or tibble that should be used in the training data. and returns a tibble where each row contains the Average Training and Average Test squared error for fitting a two-layer neural network at all possible combinations of numbers of hidden nodes at each layer.

Solution:

```
final_func<-function(x_df, outcome, input_vec, percentage){</pre>
Results <- as_tibble (expand.grid(`First layer`=c(1,2,3), `Second layer`=c(1,2,3), `Training error`=0, `Te
 scaled_data<-outcome_func(x_df,outcome,input_vec)</pre>
training_test_data<-split_data(scaled_data,percentage)</pre>
for(i in 1:9){
 current_mod<-fit_model(training_test_data$Training,</pre>
hidden_nodes_vec=as.numeric(Results[i,c("First layer","Second layer")])
)
 error <- run_training_test (current_mod,
 training_test_data$Training,
training_test_data$Test)
Results[i,c("Training error")]<-error[1]</pre>
Results[i,c("Test error")]<-error[2]</pre>
}
Results
}
```

```
set.seed(1)
my_results<-final_func(penguins_example,"sex", c("bill_length_mm","body_mass_g"), 0.7)
my_results</pre>
```

##	#	A tibble: 9 x	4		
##		'First layer'	'Second layer'	'Training error'	'Test error'
##		<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	1	1	0.186	0.221
##	2	2	1	0.183	0.195
##	3	3	1	0.154	0.233
##	4	1	2	0.180	0.215
##	5	2	2	0.143	0.205
##	6	3	2	0.102	0.184
##	7	1	3	0.135	0.193
##	8	2	3	0.0955	0.177
##	9	3	3	0.0545	0.240