

# ANN with different Sets of Layers Darshan

2024-08-01

```
# Install and load required packages
install.packages(c('neuralnet', 'keras', 'tensorflow'), dependencies = TRUE)

##
## The downloaded binary packages are in
## /var/folders/bb/9352ds8s1g5cscthpcw8v4t40000gn/T//Rtmpv9cJ1J/downloaded_packages

install.packages("tidyverse")

##
## The downloaded binary packages are in
## /var/folders/bb/9352ds8s1g5cscthpcw8v4t40000gn/T//Rtmpv9cJ1J/downloaded_packages

install.packages("cowplot")

##
## The downloaded binary packages are in
## /var/folders/bb/9352ds8s1g5cscthpcw8v4t40000gn/T//Rtmpv9cJ1J/downloaded_packages

library(neuralnet)
library(tidyverse)

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr   1.5.1
## v ggplot2    3.5.1      v tibble    3.2.1
## v lubridate  1.9.3      v tidyr     1.3.1
## v purrr      1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::compute() masks neuralnet::compute()
## x dplyr::filter()  masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors

library(cowplot)

##
## Attaching package: 'cowplot'
##
## The following object is masked from 'package:lubridate':
##
## stamp
```

```
# Prepare the iris dataset
```

```
iris <- iris %>% mutate_if(is.character, as.factor)  
summary(iris)
```

```
##   Sepal.Length   Sepal.Width   Petal.Length   Petal.Width  
##   Min.    :4.300   Min.    :2.000   Min.    :1.000   Min.    :0.100  
##   1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300  
##   Median :5.800   Median :3.000   Median :4.350   Median :1.300  
##   Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199  
##   3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800  
##   Max.    :7.900   Max.    :4.400   Max.    :6.900   Max.    :2.500  
##           Species  
##   setosa      :50  
##   versicolor:50  
##   virginica   :50  
##  
##  
##
```

```
# Set the seed and split the data into training and test sets
```

```
set.seed(254)  
data_rows <- floor(0.80 * nrow(iris))  
train_indices <- sample(c(1:nrow(iris)), data_rows)  
train_data <- iris[train_indices, ]  
test_data <- iris[-train_indices, ]
```

```
# Define a function to train and evaluate the neural network model
```

```
train_and_evaluate <- function(hidden_layers) {  
  model <- neuralnet(Species ~ Sepal.Length + Sepal.Width + Petal.Length + Petal.Width,  
                     data = train_data, hidden = hidden_layers, linear.output = FALSE)  
  pred <- predict(model, test_data)  
  labels <- c("setosa", "versicolor", "virginica")  
  prediction_label <- data.frame(max.col(pred)) %>%  
    mutate(pred = labels[max.col(pred)]) %>%  
    select(pred) %>%  
    unlist()  
  check <- as.numeric(test_data$Species) == max.col(pred)  
  accuracy <- (sum(check) / nrow(test_data)) * 100  
  list(model = model, accuracy = accuracy)  
}
```

```
# Train and evaluate models with different hidden layer configurations
```

```
hidden_layers_list <- list(c(4, 2), c(25, 9), c(40, 18))  
results <- data.frame(Hidden_Layers = character(), Accuracy = numeric(), stringsAsFactors = FALSE)  
models <- list()
```

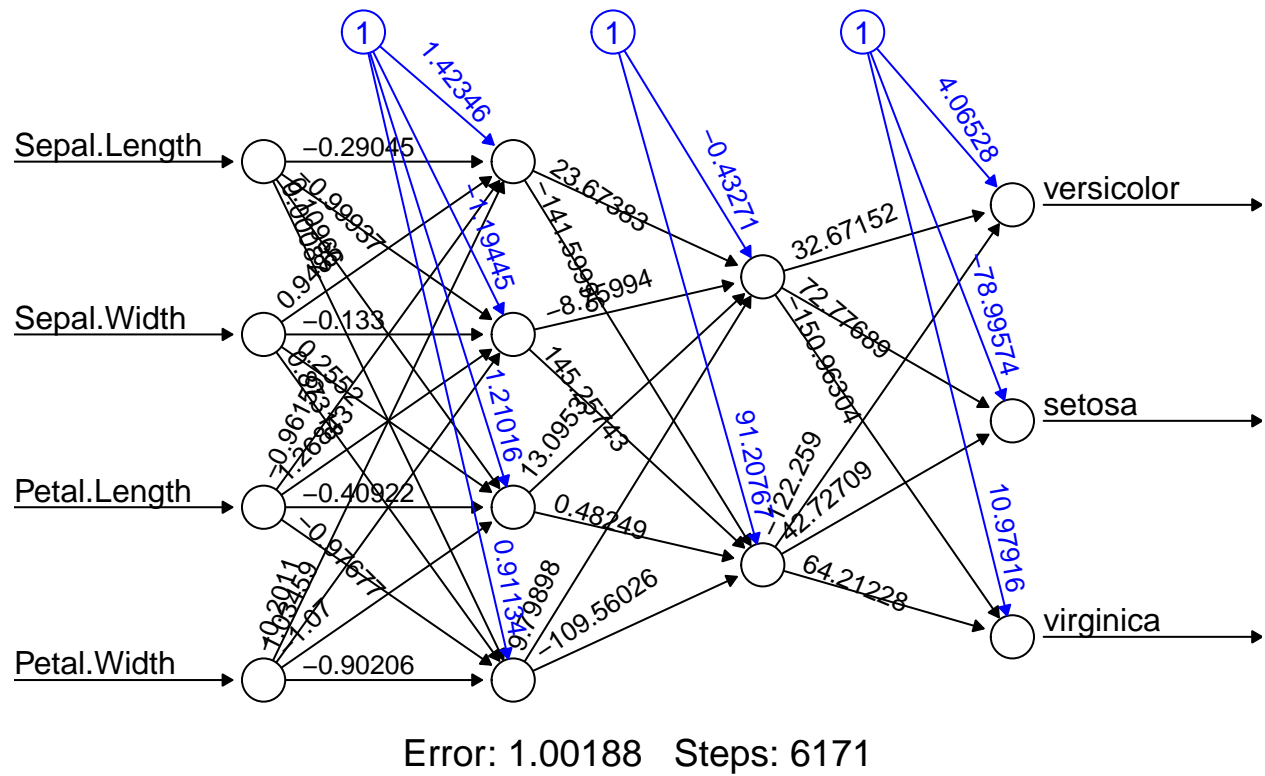
```
for (hidden_layers in hidden_layers_list) {  
  result <- train_and_evaluate(hidden_layers)  
  models <- c(models, list(result$model))  
  results <- rbind(results, data.frame(Hidden_Layers = paste(hidden_layers, collapse = ", "), Accuracy = result$accuracy))  
}
```

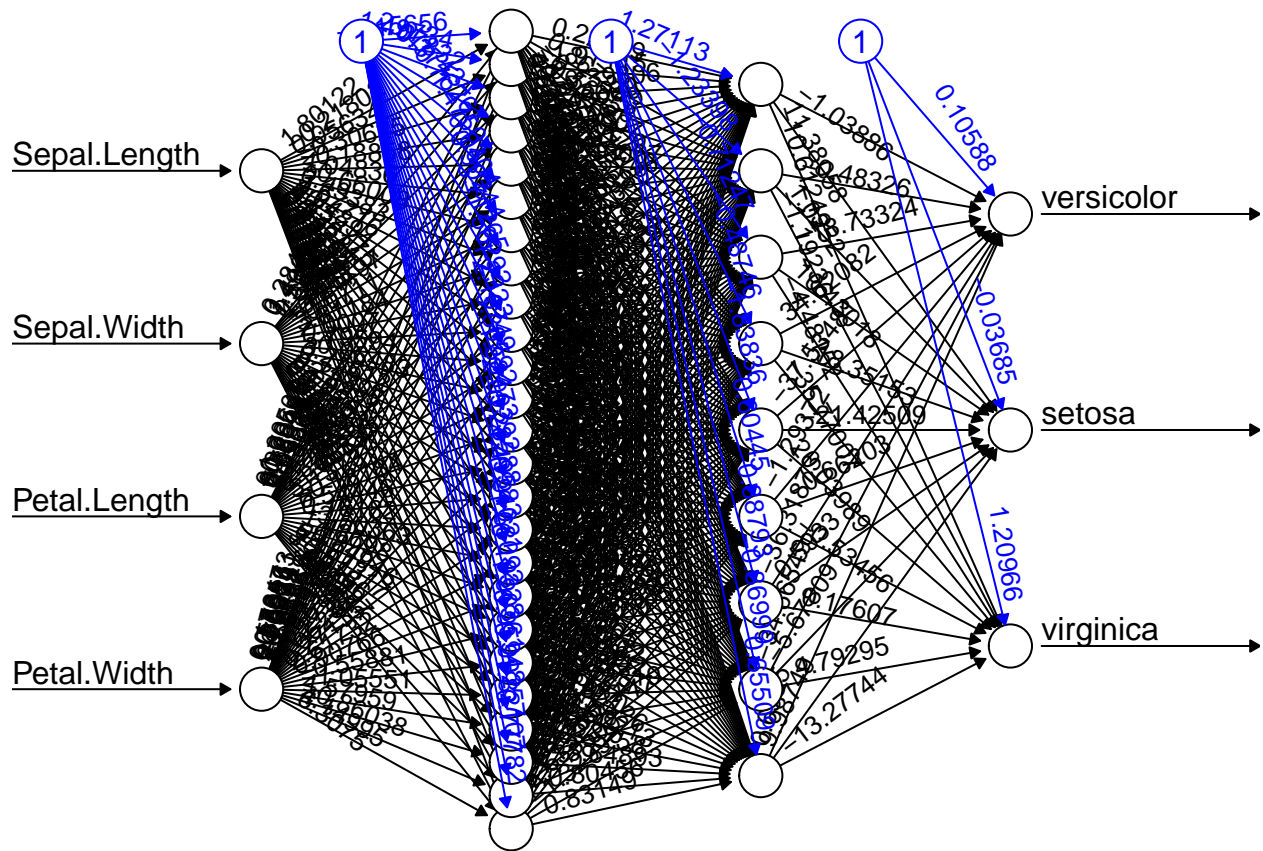
```
# Create plots for the models
```

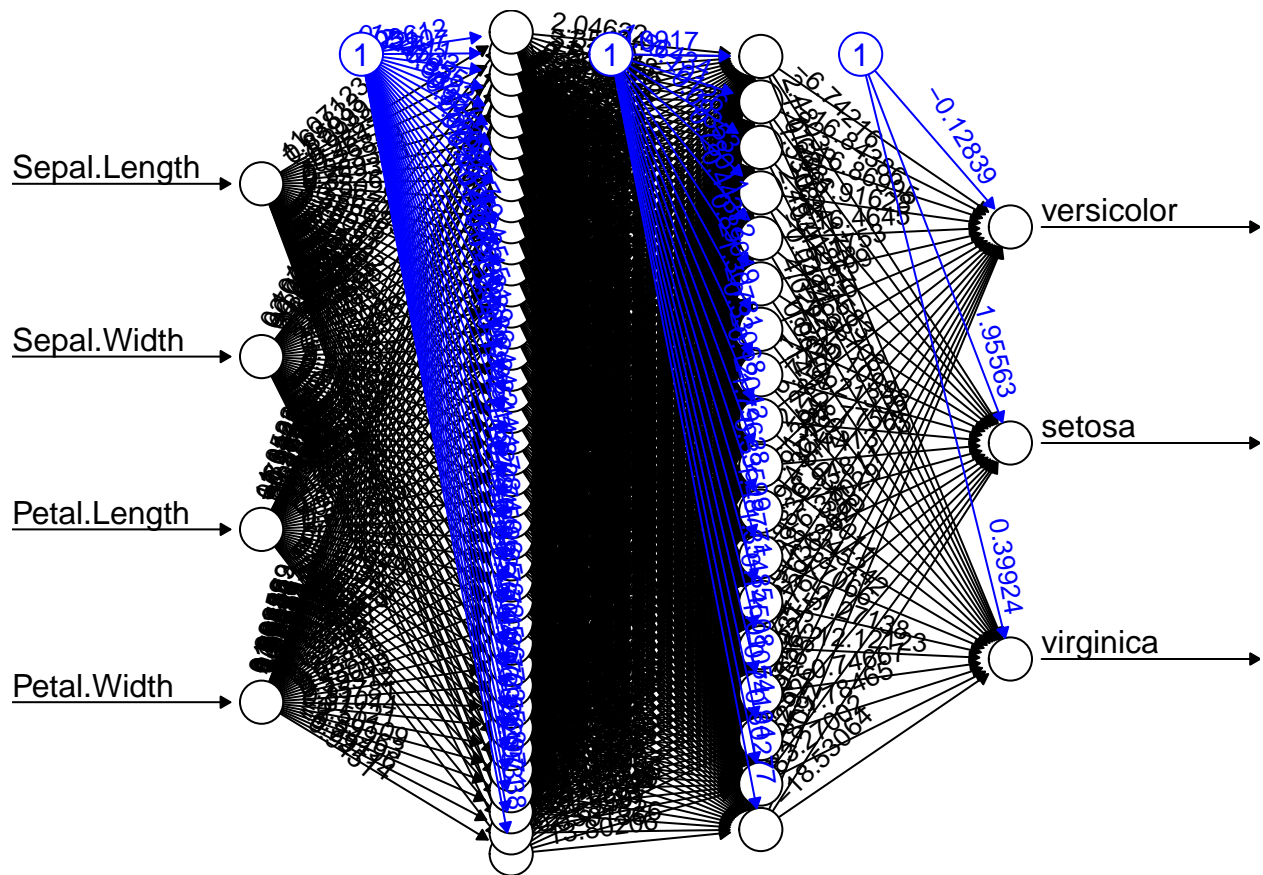
```

model_plots <- lapply(1:length(models), function(i) {
  plot(models[[i]], rep = 'best', main = paste("Model:", paste(hidden_layers_list[[i]], collapse = ", ")
})

```



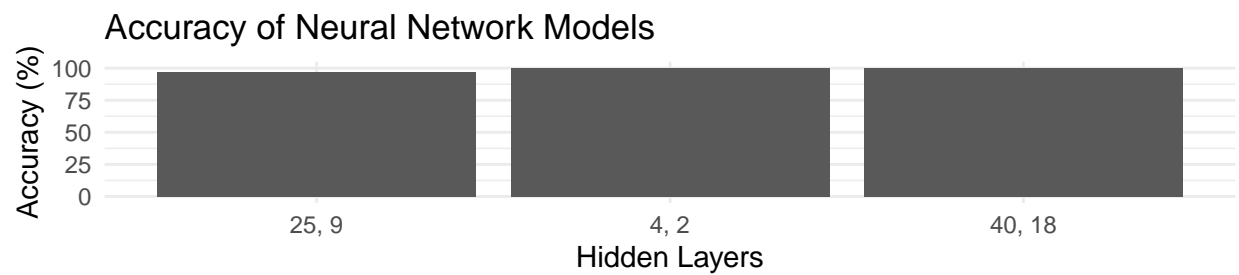




```
# Plot the accuracy bar graph
accuracy_plot <- ggplot(results, aes(x = Hidden_Layers, y = Accuracy)) +
  geom_bar(stat = "identity") +
  labs(title = "Accuracy of Neural Network Models", x = "Hidden Layers", y = "Accuracy (%)") +
  theme_minimal()

# Combine the model plots and accuracy plot using cowplot
top_row <- plot_grid(plotlist = model_plots, nrow = 1)
bottom_row <- plot_grid(accuracy_plot)
final_plot <- plot_grid(top_row, bottom_row, ncol = 1, rel_heights = c(2, 1))

# Display the final combined plot
print(final_plot)
```



```
# Print the accuracy of the models  
print(results)
```

```
##  Hidden_Layers  Accuracy  
## 1           4, 2 100.00000  
## 2          25, 9  96.66667  
## 3          40, 18 100.00000
```