

# HW07

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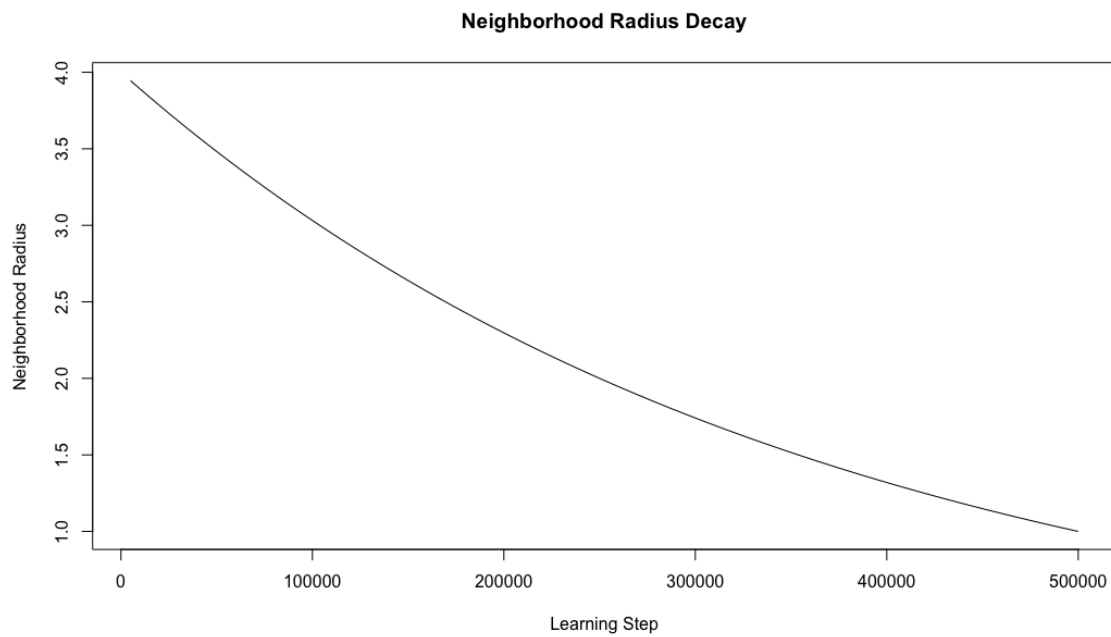
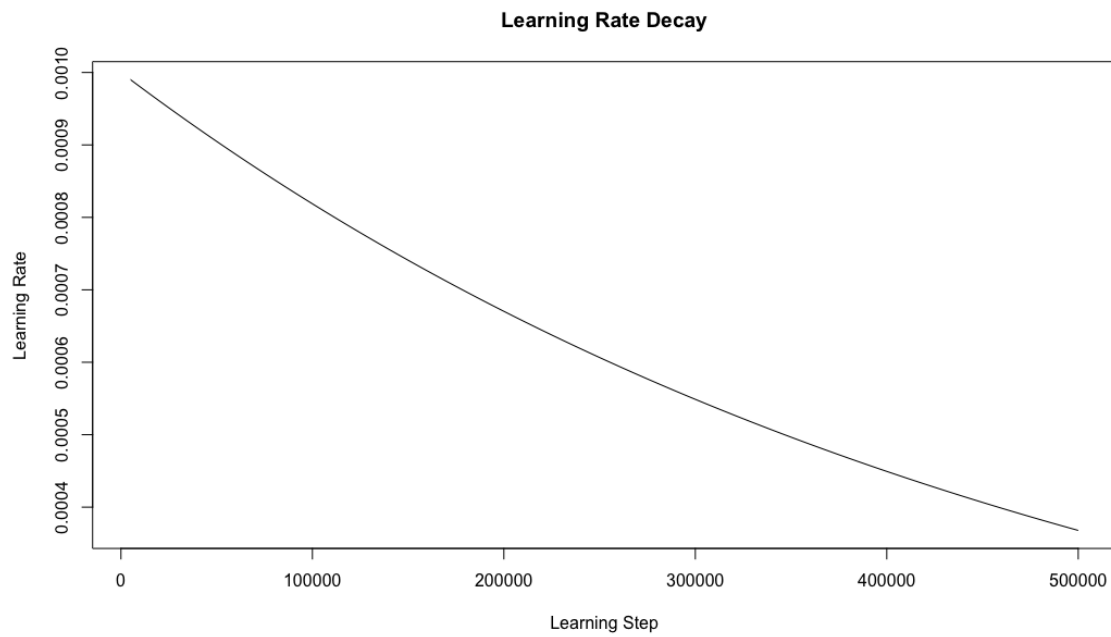
*3/21/2018*

## Problem 2a

### Parameters

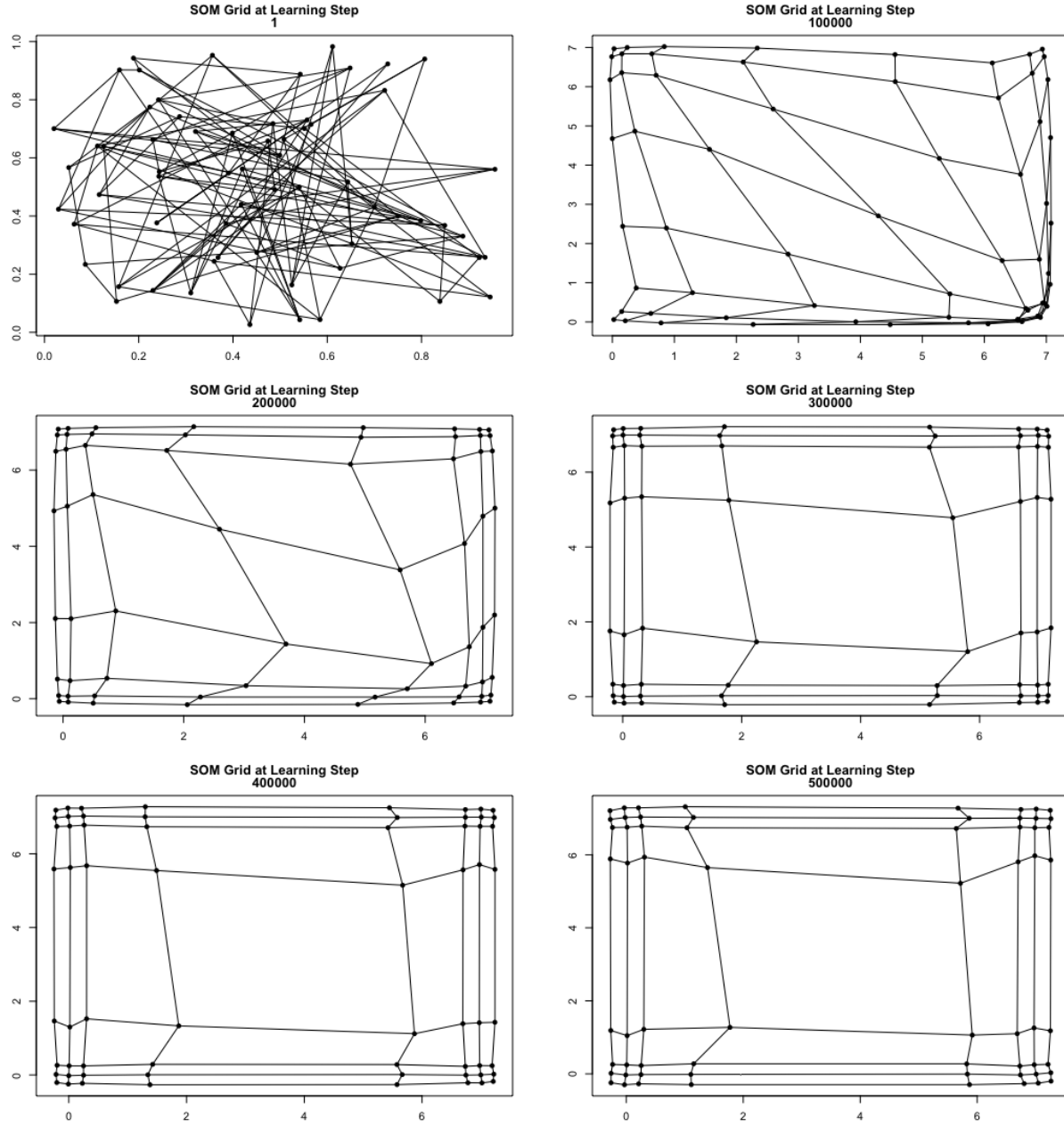
- Network Parameters
  - Topology: 64 PEs (8x8 2-dimensional square lattice)
- Learning Parameters
  - Initial Weights: Drawn from  $U[0, 1]$
  - Initial Learning Rate ( $\alpha_{init}$ ): 0.001
  - Learning Decay Rate:  $\alpha_{init} * e^{\frac{-i}{n}}$ 
    - \*  $i$  = current learning step
    - \*  $n$  = total number of learning steps
  - Initial Radius ( $\theta_{init}$ ): 4
  - Radius Decay Rate:  $\theta_{init} * e^{\frac{-i}{\mu}}$ 
    - \*  $i$  = current learning step
    - \*  $n$  = total number of learning steps
    - \*  $\mu = \frac{n}{\log(\theta_{init})}$
  - Stopping Criteria: 500,000 learning steps
- Input Data
  - 1000 Samples drawn from each (4000 total):
    - \* Normal[(7,7), 0.1]
    - \* Normal[(0,7), 0.1]
    - \* Normal[(7,0), 0.1]
    - \* Normal[(0,0), 0.1]

## Decay Graphs



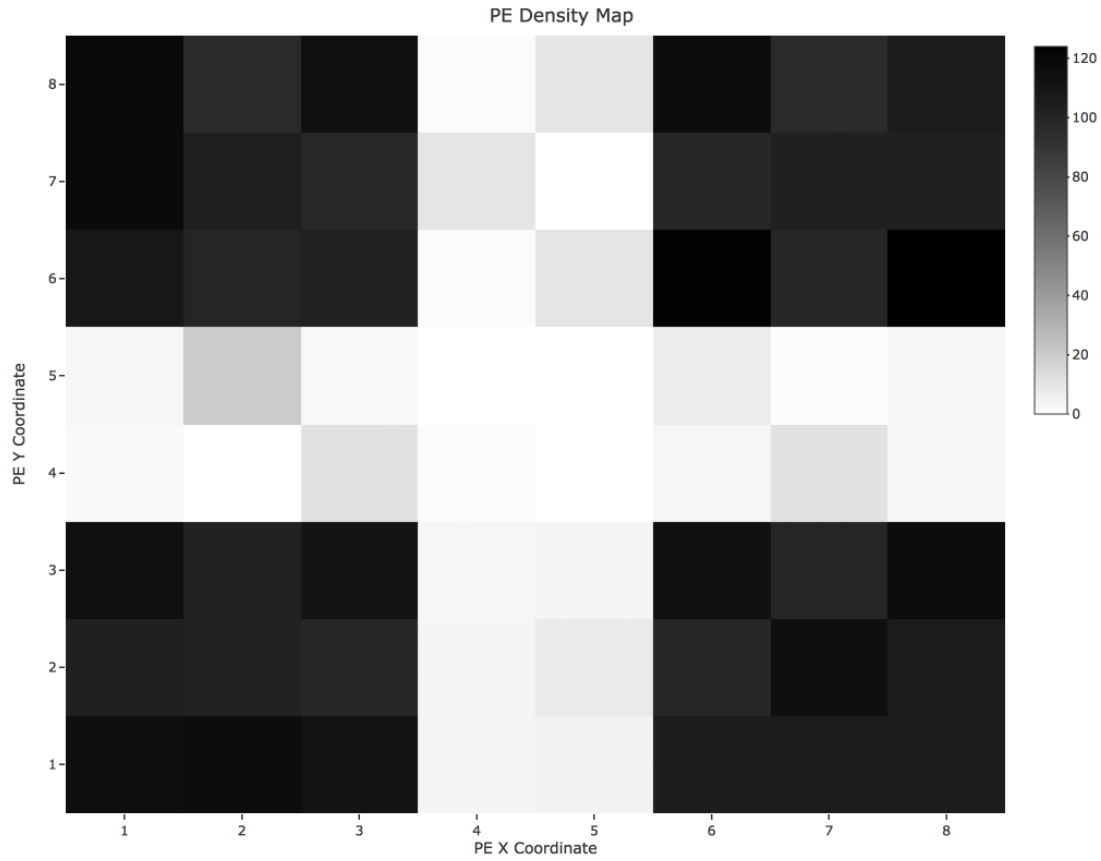
As we can see, we used an exponential decay function to decrease the neighborhood radius and learning rate over time as a function of the number of learning steps completed; this helped us to obtain very strong network results.

## Lattice Learning Graphs



As we can see, we achieved very good results on our data; we can see very clearly at the end of our learning that the prototypes have gravitated to the points  $(0,0)$ ,  $(0,7)$ ,  $(7,0)$  and  $(7,7)$ . We can also see that the “density” of the data at those locations are very similar and almost equal as the prototypes form a symmetry at the four quadrants (this makes sense as our data is equally distributed around those four points). The learning in the first 100,000 steps causes a massive change to the graph showing the prototypes mapped back to the input space, while as the learning continues it changes less drastically, both as a result of our parameter decay and the convergence of the network to our stable solution.

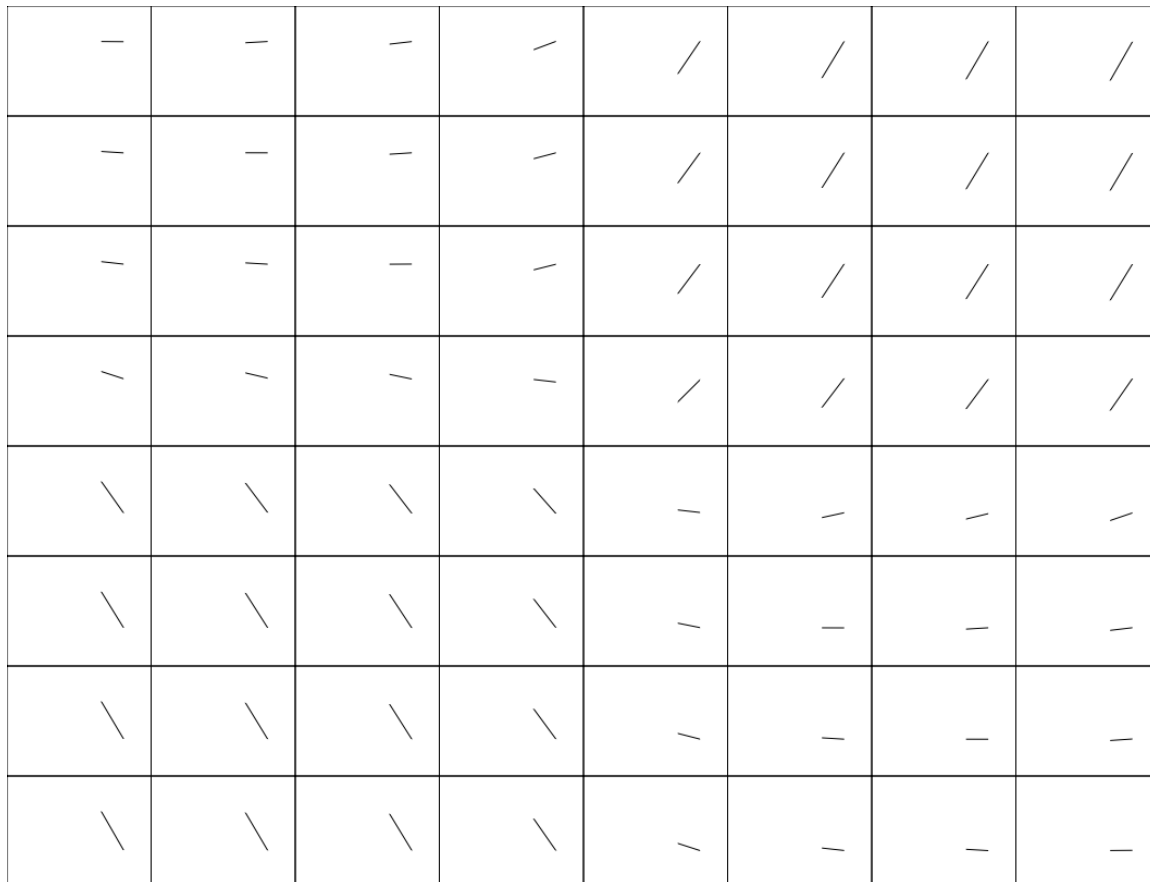
## Recalled PE Mapping



Our recalled density mapping shows great results as well. As we can clearly see in this grey-scale heatmap representation of the amount of data points that map to each of our 64 PEs, the clear majority of the data points map to one of the PEs in the corners represented by the prototypes that map to our data epicenters. We have very clear definition in the intermediary PEs showing that very few data points map to them; this is our expected result, since the learning process would have mapped most of our points to the prototypes in the four corners.

### Problem 3

## 2D Weight Vectors Plot



Our 2-dimension weight vector plots does an excellent job of showing how well our clustering relations are visualized. Each cell in this 8x8 visualization represents the corresponding PE in our lattice, where the x-axis is the sequential number of dimensions of that PE in in the input space (in this case, two dimension) and the y-axis represents the value at those dimensions. Even without the dead zones marked, we can see very clear and profound delineations between our four clusters.

### Problem 4

Blah blah.

### Problem 5

Blah blah.

## Code Appendix

```
## No scientific notation

options(scipen = 999)

## Load in the necessary libraries

library(plotly)

## Read in the necessary functions

source("~/Documents/Rice_University/Spring_2018/NML502/HW07/build_SOM.R")
source("~/Documents/Rice_University/Spring_2018/NML502/HW07/learn_map.R")
source("~/Documents/Rice_University/Spring_2018/NML502/HW07/plot_SOM.R")
source("~/Documents/Rice_University/Spring_2018/NML502/HW07/recall_PE.R")

##### Problem 2 #####

## Set the SOM lattice dimensions

matrix_dim <- 8

## Set the initial radius of influence on neighbors

radius <- matrix_dim / 2

## Set the number of iterations

num_iter <- 500000

## Set the initial learning rate

ler_rate <- 0.001

## Load in the input data

x_1 <- rnorm(n = 1000, mean = 7, sd = sqrt(0.1))
x_2 <- rnorm(n = 1000, mean = 7, sd = sqrt(0.1))

x_3 <- rnorm(n = 1000, mean = 0, sd = sqrt(0.1))
x_4 <- rnorm(n = 1000, mean = 7, sd = sqrt(0.1))

x_5 <- rnorm(n = 1000, mean = 7, sd = sqrt(0.1))
x_6 <- rnorm(n = 1000, mean = 0, sd = sqrt(0.1))

x_7 <- rnorm(n = 1000, mean = 0, sd = sqrt(0.1))
x_8 <- rnorm(n = 1000, mean = 0, sd = sqrt(0.1))
```

```

X_1 <- rbind(x_1, x_2)
X_2 <- rbind(x_3, x_4)
X_3 <- rbind(x_5, x_6)
X_4 <- rbind(x_7, x_8)

X <- cbind(X_1, X_2, X_3, X_4)

rownames(X) <- NULL

## Get the size of a typical input

input_size <- length(X[, 1])

## Build the SOM matrix

SOM_lattice <- build_SOM(input_size, matrix_dim)

## Perform the learning process

learn_results <- learn_map(SOM_lattice, X, num_iter, radius, ler_rate)

## Plot the results

par(mfrow = c(3, 2))
par(mar = c(3, 3, 3, 3))
plot_SOM(learn_results)

## Plot the decay of the radius and learning rate

plot(x = param_container[[1]], y = param_container[[2]], type = "l",
     xlab = "Learning Step",
     ylab = "Neighborhood Radius",
     main = "Neighborhood Radius Decay")

plot(x = param_container[[1]], y = param_container[[3]], type = "l",
     xlab = "Learning Step",
     ylab = "Learning Rate",
     main = "Learning Rate Decay")

## Perform the recall

recall_PE(learn_results, X)

##### Problem 3 #####

## Unwrap the final PE prototypes

x_lattice <- learn_results[[length(learn_results)]] [[2]] [[1]]
y_lattice <- learn_results[[length(learn_results)]] [[2]] [[2]]

```

```

## Scale the PEs down and vectorize for plotting purposes

scale_x <- scale(c(t(x_lattice)))
scale_y <- scale(c(t(y_lattice)))

## Generate the frame and add the plots

par(mfrow = c(matrix_dim, matrix_dim))
par(mar = c(0, 0, 0, 0))

for (i in 1:length(x_lattice)) {

  plot(x = 1:input_size,
       y = c(scale_x[i], scale_y[i]),
       type = "l",
       xaxt = "n",
       yaxt = "n",
       ann = FALSE,
       xlim = c(-3, 3),
       ylim = c(-3, 3))

}

## Build the function that learns the mapping

learn_map <- function(SOM_lattice, X, num_iter, radius, ler_rate) {

  ## Container for the current prototype lattice

  SOM_container <- list()

  ## Container for learning step, radius and learning rate

  param_container <- list()
  ler_steps <- numeric()
  radii <- numeric()
  ler_rates <- numeric()

  ## Save the initial parameters

  init_radius <- radius
  decay_constant <- num_iter / log(init_radius)

  init_ler_rate <- ler_rate

  ## Perform the iterated learning

  for (i in 1:num_iter) {

    ## Decay the learning rate

    ler_rate <- init_ler_rate * exp(-i / num_iter)

```



```

## Decay the radius

radius <- init_radius * exp(-i / decay_constant)

## Add the parameters states to the container

if (i %% 5000 == 0) {

  ler_steps[length(ler_steps) + 1] <- i
  radii[length(radii) + 1] <- radius
  ler_rates[length(ler_rates) + 1] <- ler_rate

}

## Randomly select an input vector

rand_ind <- sample(x = 1:dim(X)[2], size = 1)
x <- as.matrix(X[, rand_ind], ncol = 1)

## Calculate the matrix of differences

matrix_diffs <- list()
matrix_diffs[[1]] <- x[1, ] - SOM_lattice[[2]][[1]]
matrix_diffs[[2]] <- x[2, ] - SOM_lattice[[2]][[2]]

## Get the Euclidean distance

mat_a <- matrix_diffs[[1]]^2
mat_b <- matrix_diffs[[2]]^2
matrix_dist <- sqrt(mat_a + mat_b)

## Find the winning PE

min_neuron <- min(matrix_dist)
min_loc <- as.vector(which(matrix_dist == min(matrix_dist), arr.ind = TRUE))

## Calculate the Manhattan distance

mat_c <- abs(SOM_lattice[[1]][[1]] - min_loc[1])
mat_d <- abs(SOM_lattice[[1]][[2]] - min_loc[2])
man_dist <- mat_c + mat_d
neighbor_func <- exp(-((man_dist)/(radius))^2)

## Update the weights

SOM_lattice[[2]][[1]] <- SOM_lattice[[2]][[1]] + ler_rate *
  (neighbor_func * matrix_diffs[[1]])
SOM_lattice[[2]][[2]] <- SOM_lattice[[2]][[2]] + ler_rate *
  (neighbor_func * matrix_diffs[[2]])

## Add to the container

if ((i == 1) || (i %in% seq(from = 0, to = num_iter, length.out = 6))) {

```

```

        SOM_container[[length(SOM_container) + 1]] <- list(i, SOM_lattice[[2]])
    }

}

param_container[[1]] <- ler_steps
param_container[[2]] <- radii
param_container[[3]] <- ler_rates

param_container <- param_container

return(SOM_container)
}

## Build the function to recall each input
recall_PE <- function(learn_results, X) {

    ## Unwrap the final SOM lattice

    x_lattice <- learn_results[[length(learn_results)]] [[2]] [[1]]
    y_lattice <- learn_results[[length(learn_results)]] [[2]] [[2]]

    ## Container for which PE the input maps to

    neuron_map <- matrix(0, ncol = dim(x_lattice)[1], nrow = dim(x_lattice)[1])
    colnames(neuron_map) <- 1:dim(x_lattice)[1]
    rownames(neuron_map) <- 1:dim(x_lattice)[1]

    ## Loop through the input data for recall

    for (i in 1:dim(X)[2]) {

        ## Get the current input vector

        x <- matrix(X[, i], ncol = 1)

        ## Calculate the matrix of differences

        matrix_diffs <- list()
        matrix_diffs[[1]] <- x[1, ] - x_lattice
        matrix_diffs[[2]] <- x[2, ] - y_lattice

        ## Get the Euclidean distance

        mat_a <- matrix_diffs[[1]]^2
        mat_b <- matrix_diffs[[2]]^2
        matrix_dist <- sqrt(mat_a + mat_b)

        ## Find the winning PE

```

```

min_neuron <- min(matrix_dist)
min_loc <- as.vector(which(matrix_dist == min(matrix_dist), arr.ind = TRUE))

## Add the location to the neuron map

if (neuron_map[min_loc[1], min_loc[2]] == 0) {

  neuron_map[min_loc[1], min_loc[2]] <- 1

} else {

  neuron_map[min_loc[1], min_loc[2]] <- neuron_map[min_loc[1], min_loc[2]] + 1

}

}

s <- as.character(1:matrix_dim)

temp_vec <- as.vector(neuron_map)
temp_mat <- matrix(temp_vec, nrow = matrix_dim, ncol = matrix_dim)
temp_mat <- apply(temp_mat, 2, rev)

plot_ly(z = temp_mat, x = ~s, y = ~s, colors = colorRamp(c("white", "black")),
        type = "heatmap") %>%
  layout(title = "PE Density Map", xaxis = list(title = "PE X Coordinate"),
        yaxis = list(title = "PE Y Coordinate"))

}

## Function to build the SOM matrix

build_SOM <- function(input_size, matrix_dim) {

  ## Container for the matrices of weights

  SOM_weights <- list()

  ## Container for the indices

  SOM_indices <- list()

  ## Build the list of weight indices

  SOM_indices[[1]] <- matrix(rep(1:matrix_dim, times = matrix_dim),
                             nrow = matrix_dim, ncol = matrix_dim)

  SOM_indices[[2]] <- matrix(rep(1:matrix_dim, times = matrix_dim),
                             byrow = T, nrow = matrix_dim, ncol = matrix_dim)

  ## Build the list of weights

```

```

for (i in 1:input_size) {

  SOM_weights[[i]] <- matrix(runif(n = matrix_dim^2, min = 0, max = 1),
                             ncol = matrix_dim, nrow = matrix_dim)

}

## Return the results

return(list(SOM_indices, SOM_weights))

}

## Build the plot_SOM function

plot_SOM <- function(learn_results) {

  ## For loop to iterate over the results

  for (i in 1:length(learn_results)) {

    ## Unwrap the learning step and lattice at each step

    ler_step <- learn_results[[i]][[1]]
    x_lattice <- as.vector(learn_results[[i]][[2]][[1]])
    y_lattice <- as.vector(learn_results[[i]][[2]][[2]])

    ## Build the lattice

    plot_lattice <- cbind(x_lattice, y_lattice)

    ## Plot the lattice

    plot(plot_lattice, pch = 16, main = c("SOM Grid at Learning Step", ler_step),
         xlab = "", ylab = "")

    ## Add the horizontal neighbor lines

    for (i in 1:dim(plot_lattice)[1]) {

      if (i %% matrix_dim != 0) {

        segments(plot_lattice[i, 1], plot_lattice[i, 2], plot_lattice[i + 1, 1],
                  plot_lattice[i + 1, 2])

      }

    }

    ## Add the vertical neighbor lines

    for (i in 1:dim(plot_lattice)[1]) {

```

```
if (((dim(plot_lattice)[1] - i) - matrix_dim) >= 0) {  
    segments(plot_lattice[i, 1], plot_lattice[i, 2], plot_lattice[i + matrix_dim, 1],  
             plot_lattice[i + matrix_dim, 2])  
}  
  
}  
  
}  
  
}
```