perceptron

Joyce Robbins 8/13/2018

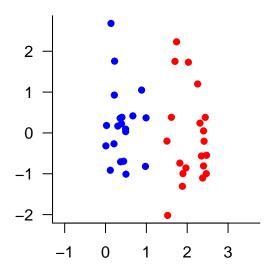
The perceptron is a simple algorithm that learns to classify inputs into two classes by adjusting the weights (w) in the equation $y_i = \text{sign}(w_i x_i)$ until all inputs in a training set are correctly classified. Here the steps of algorithm will be presented visually in two-dimensional space.

The basics

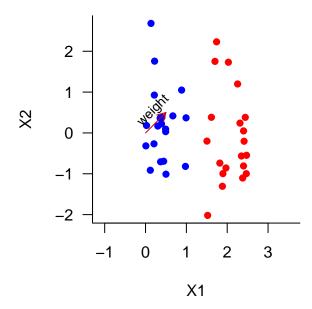
We start by plotting (x_1, x_2) , coloring each point by class. Note that the points can be separated by a line; if this is not the case, the algorithm won't work.

```
#devtools::install_github("jtr13/perceptron")
library(perceptron)
set.seed(9)
X <- matrix(c(runif(20), runif(20)+1.5, rnorm(40)), nrow = 40)
Y <- c(rep(-1, 20), rep(1, 20))

X <- cbind(rep(1, length(Y)), X)
W <- c(0, .5, .5)
weight_matrix <- matrix(W, 1)
i <- 1
converged <- FALSE
set.seed(1)
par(xpd = TRUE)
draw_points(X[,2], X[,3], Y, axes = TRUE)</pre>
```



We start with an arbitrary weight vector, (w0, w1, w2). Often (0, 0, 0) is used, but we'll start with (0, 0.5, 0.5) so we can visualize it:

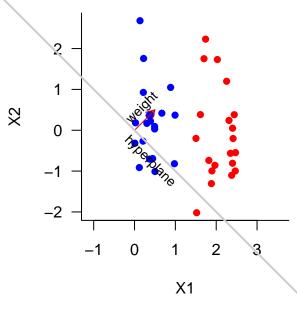


The decision boundary, or hyperplane, is the line orthogonal to the weight vector. For points on the line, the sign of $(w_i x_i)$ equals zero. On one side of the line, the sign of $(w_i x_i)$ is greater than zero whereas on the other side the sign of $(w_i x_i)$ is less than zero; hence the line serves to divide all points into two classes according to the perceptron logic.

```
all.mis.points \leftarrow X[sign(W \% * \% t(X)) != Y, , drop = F]
if (length(all.mis.points) == 0) converged <- TRUE</pre>
if (!converged) {
  all.mis.points.Y <- Y[sign(W %*% t(X)) != Y]</pre>
  index <- sample(nrow(all.mis.points), 1)</pre>
  mis.point <- all.mis.points[index,]</pre>
  mis.point.Y <- all.mis.points.Y[index]</pre>
}
s <- shift(W)
draw_points(X[,2], X[,3], Y, axes = TRUE)
mtext("X1", side = 1, line = 3)
mtext("X2", side = 2, line = 3)
draw_weight_vector(W)
label_vector(s["x"], s["y"], W[2] + s["x"],
              W[3] + s["y"], "weight")
draw_hyperplane(W)
e <- get_endpoints(W)
if (sum(is.na(e)) == 0) {
  label_vector(e[1], e[2], e[3], e[4],
                label = "hyperplane",
                "below")
}
all.mis.points \leftarrow X[sign(W \%*\% t(X)) != Y, , drop = F]
if (length(all.mis.points) == 0) converged <- TRUE</pre>
```

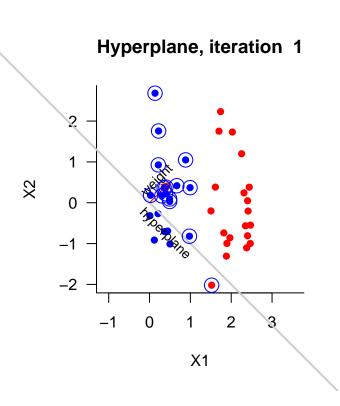
```
if (!converged) {
   all.mis.points.Y <- Y[sign(W %*% t(X)) != Y]
   index <- sample(nrow(all.mis.points), 1)
   mis.point <- all.mis.points[index,]
   mis.point.Y <- all.mis.points.Y[index]
}
if (!converged) {
   title(paste("Hyperplane, iteration ", i))
} else {
     title(paste("\nConverged! Iteration ", i))
}</pre>
```

Hyperplane, iteration 1



Note the circled points – these are the misclassified points – the ones for which $y_i \neq \text{sign}(w_i x_i)$.

```
if (length(all.mis.points) == 0) converged <- TRUE
if (!converged) {
    all.mis.points.Y <- Y[sign(W %*% t(X)) != Y]
    index <- sample(nrow(all.mis.points), 1)
    mis.point <- all.mis.points[index,]
    mis.point.Y <- all.mis.points.Y[index]
}
if (!converged) {
    title(paste("Hyperplane, iteration ", i))
} else {
        title(paste("\nConverged! Iteration ", i))
}
points(all.mis.points[,2],
        all.mis.points[,3],
        col = "blue", cex = 2)</pre>
```



The Algorithm

The perceptron algorithm works by updating the weight vector based on a randomly selected misclassified point, calculating the new hyperplane, and repeating until the hyperplane separates all points into the two classes.

The formula for the new weight vector is:

```
w_{t+1} = w_t + \eta y_i x_i, where x_i = the misclassified point y_i = the true label of the misclassified point (-1 or 1)
```

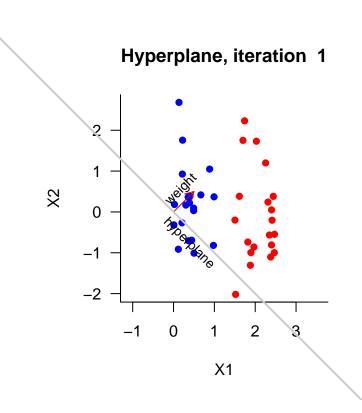
 $\eta =$ the learning rate, which we'll set to 1 for the sake of simplicity

Visually, the new weight vector, w_{t+1} , is determined by adding $y_i x_i$ to w_t and then shifting by the offset $w_0/||w||_2$.

We'll go through the algorithm one step at a time.

We begin with our original weight vector and hyperplane:

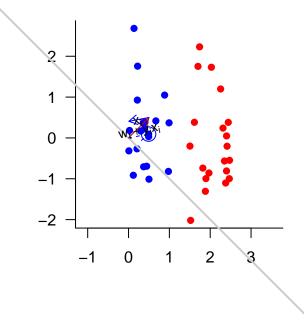
```
s <- shift(W)
draw_points(X[,2], X[,3], Y, axes = TRUE)
mtext("X1", side = 1, line = 3)
mtext("X2", side = 2, line = 3)
draw_weight_vector(W)
label_vector(s["x"], s["y"], W[2] + s["x"],
             W[3] + s["y"], "weight")
draw_hyperplane(W)
e <- get_endpoints(W)</pre>
if (sum(is.na(e)) == 0) {
  label_vector(e[1], e[2], e[3], e[4],
               label = "hyperplane",
                "below")
}
all.mis.points <- X[sign(W \%*\% t(X)) != Y, , drop = F]
if (length(all.mis.points) == 0) converged <- TRUE</pre>
if (!converged) {
  all.mis.points.Y <- Y[sign(W %*% t(X)) != Y]</pre>
  index <- sample(nrow(all.mis.points), 1)</pre>
  mis.point <- all.mis.points[index,]</pre>
  mis.point.Y <- all.mis.points.Y[index]</pre>
if (!converged) {
  title(paste("Hyperplane, iteration ", i))
} else {
    title(paste("\nConverged! Iteration ", i))
}
```



Next we randomly select a misclassified point. In the diagram below, x_i is shown as a **dashed blue arrow**, and $y_i x_i$ added to w_t as a **solid blue arrow**:

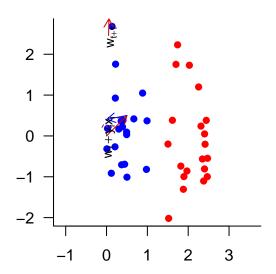
```
draw_points(X[,2], X[,3], Y, axes = TRUE)
title("1) Select a misclassified point")
draw_weight_vector(W)
draw_hyperplane(W)
circle_point(mis.point)
draw_mis_vector(mis.point)
label_vector(0, 0, mis.point[2],
             mis.point[3],
             expression(x[i]))
draw_mis_vector_added(W, mis.point,
                      mis.point.Y)
s <- shift(W)
label_vector(W[2] + s["x"],
             W[3] + s["y"],
             W[2] + s["x"] + mis.point.Y *
               mis.point[2],
             W[3] + s["y"] + mis.point.Y *
               mis.point[3],
             expression(w[t] + y[i]*x[i]),
             "below")
```





Next we determine the new weight vector by shifting the vector sum by $w_0/||w||_2$:

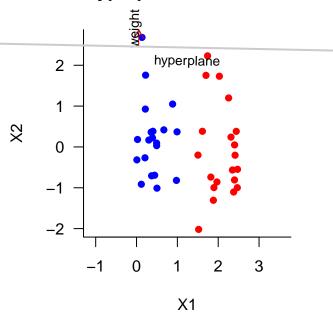
```
draw_points(X[,2], X[,3], Y, axes = TRUE)
title("2) Draw new weight vector")
draw_weight_vector(W)
draw_mis_vector_added(W, mis.point, mis.point.Y)
draw_new_weight_vector(W, mis.point, mis.point.Y)
label_vector(s["x"],
             s["y"],
             W[2] + s["x"] + mis.point.Y * mis.point[2],
             W[3] + s["y"] + mis.point.Y * mis.point[3], expression(w[t] + y[i]*x[i]), "below")
# calculate new weight vector
W <- W + mis.point.Y %*% mis.point
weight_matrix <- rbind(weight_matrix, matrix(W, 1))</pre>
i <- i + 1
draw_weight_vector(W)
s <- shift(W)
label_vector(s["x"], s["y"], W[2] + s["x"],
             W[3] + s["y"], expression(w[t+1]),
             "below")
```



Finally, we draw the new hyperplane:

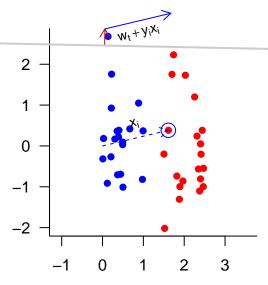
```
s <- shift(W)
draw_points(X[,2], X[,3], Y, axes = TRUE)
mtext("X1", side = 1, line = 3)
mtext("X2", side = 2, line = 3)
draw_weight_vector(W)
label_vector(s["x"], s["y"], W[2] + s["x"],
             W[3] + s["y"], "weight")
draw_hyperplane(W)
e <- get_endpoints(W)</pre>
if (sum(is.na(e)) == 0) {
  label_vector(e[1], e[2], e[3], e[4],
               label = "hyperplane",
               "below")
all.mis.points <- X[sign(W %*% t(X)) != Y, , drop = F]
if (length(all.mis.points) == 0) converged <- TRUE</pre>
if (!converged) {
  all.mis.points.Y <- Y[sign(W ** t(X)) != Y]
  index <- sample(nrow(all.mis.points), 1)</pre>
  mis.point <- all.mis.points[index,]</pre>
  mis.point.Y <- all.mis.points.Y[index]</pre>
if (!converged) {
  title(paste("Hyperplane, iteration ", i))
} else {
    title(paste("\nConverged! Iteration ", i))
}
```

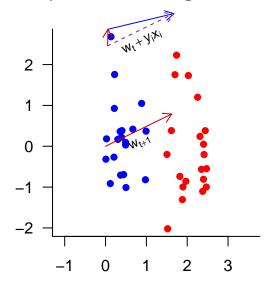
Hyperplane, iteration 2



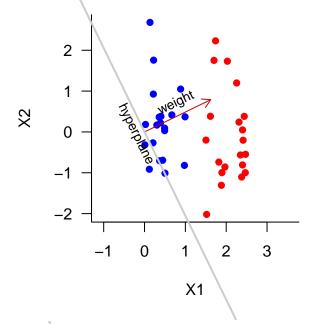
```
while (!converged) {
  draw_points(X[,2], X[,3], Y, axes = TRUE)
  title("1) Select a misclassified point")
  draw weight vector(W)
  draw_hyperplane(W)
  circle_point(mis.point)
  draw_mis_vector(mis.point)
  label_vector(0, 0, mis.point[2],
               mis.point[3],
               expression(x[i]))
  draw_mis_vector_added(W, mis.point,
                        mis.point.Y)
  s <- shift(W)
  label_vector(W[2] + s["x"],
               W[3] + s["y"],
               W[2] + s["x"] + mis.point.Y *
                 mis.point[2],
               W[3] + s["y"] + mis.point.Y *
                 mis.point[3],
               expression(w[t] + y[i]*x[i]),
               "below")
  draw_points(X[,2], X[,3], Y, axes = TRUE)
  title("2) Draw new weight vector")
  draw_weight_vector(W)
  draw_mis_vector_added(W, mis.point, mis.point.Y)
  draw_new_weight_vector(W, mis.point, mis.point.Y)
  label_vector(s["x"],
               s["y"],
               W[2] + s["x"] + mis.point.Y * mis.point[2],
```

```
W[3] + s["y"] + mis.point.Y * mis.point[3], expression(w[t] + y[i]*x[i]), "below")
  # calculate new weight vector
  W <- W + mis.point.Y ** mis.point
  weight_matrix <- rbind(weight_matrix, matrix(W, 1))</pre>
  i <- i + 1
  draw_weight_vector(W)
  s <- shift(W)
  label_vector(s["x"], s["y"], W[2] + s["x"],
               W[3] + s["y"], expression(w[t+1]),
                "below")
  s <- shift(W)
  draw_points(X[,2], X[,3], Y, axes = TRUE)
  mtext("X1", side = 1, line = 3)
  mtext("X2", side = 2, line = 3)
  draw_weight_vector(W)
  label_vector(s["x"], s["y"], W[2] + s["x"],
               W[3] + s["y"], "weight")
  draw_hyperplane(W)
  e <- get_endpoints(W)</pre>
  if (sum(is.na(e)) == 0) {
    label_vector(e[1], e[2], e[3], e[4],
                  label = "hyperplane",
                  "below")
  }
  all.mis.points <- X[sign(W %*% t(X)) != Y, , drop = F]
  if (length(all.mis.points) == 0) converged <- TRUE</pre>
  if (!converged) {
    all.mis.points.Y <- Y[sign(W %*% t(X)) != Y]</pre>
    index <- sample(nrow(all.mis.points), 1)</pre>
    mis.point <- all.mis.points[index,]</pre>
    mis.point.Y <- all.mis.points.Y[index]</pre>
  if (!converged) {
    title(paste("Hyperplane, iteration ", i))
      title(paste("\nConverged! Iteration ", i))
  }
}
```

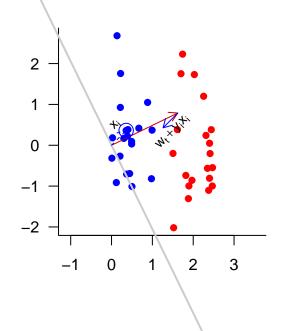


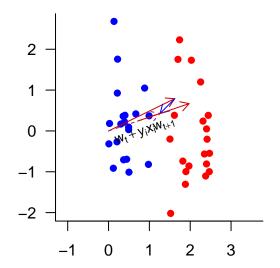


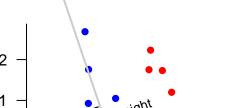
Hyperplane, iteration 3



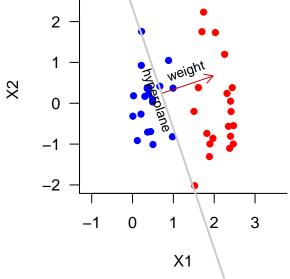
1) Select a misclassified point

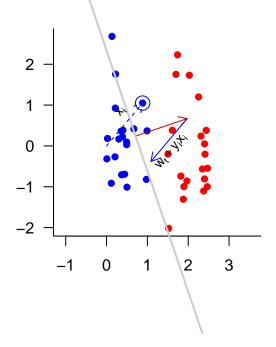


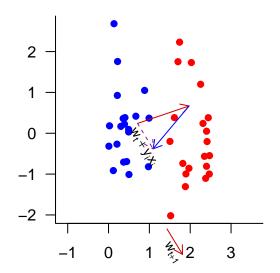




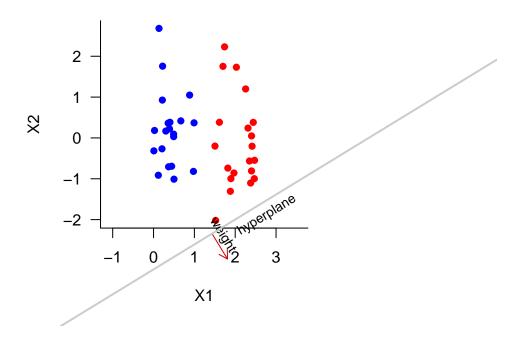
Hyperplane, iteration 4



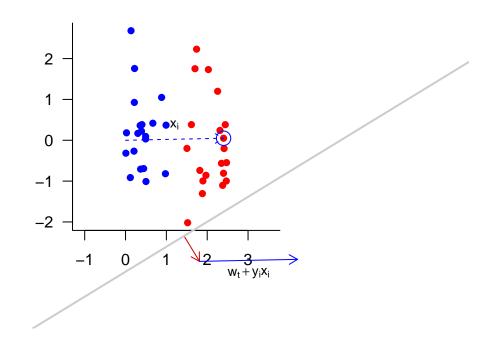


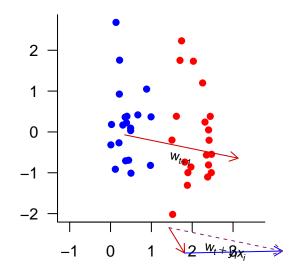


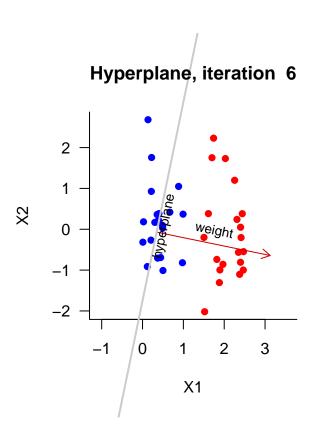
Hyperplane, iteration 5

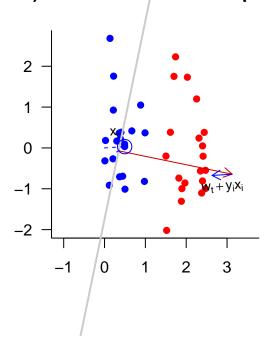


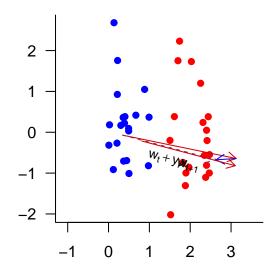
1) Select a misclassified point

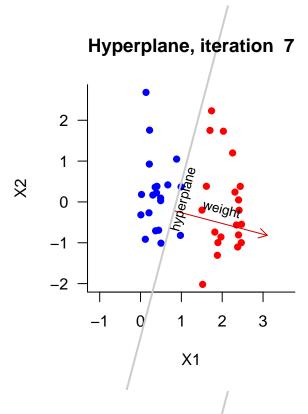


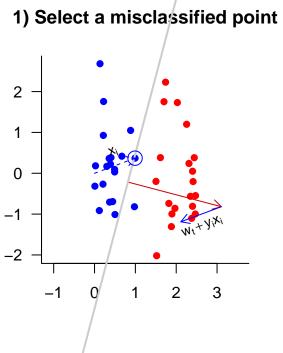


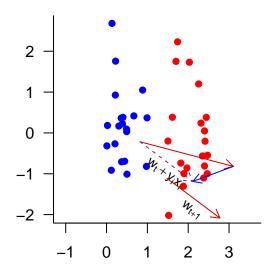


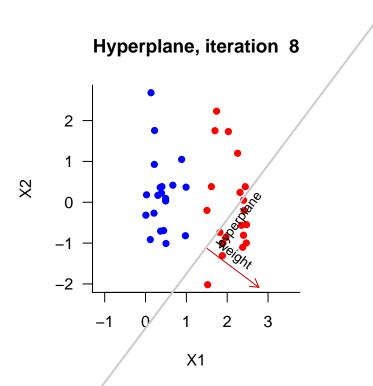


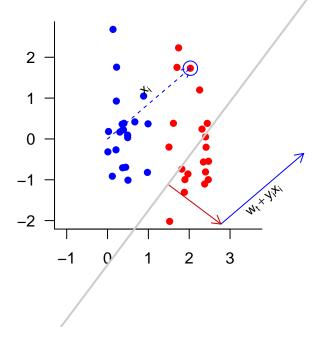


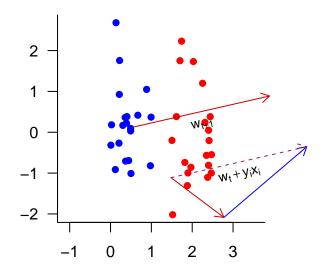


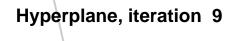


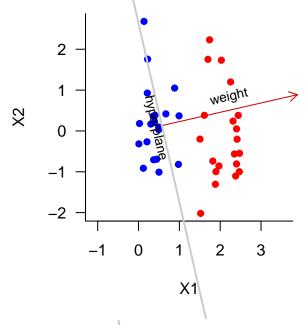


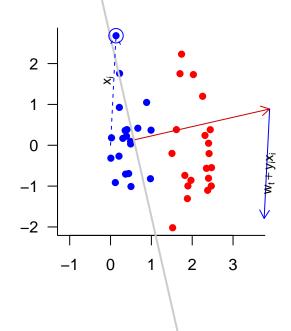


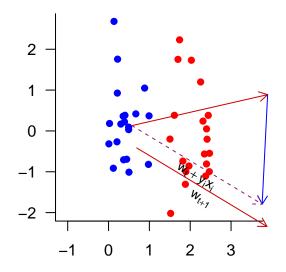


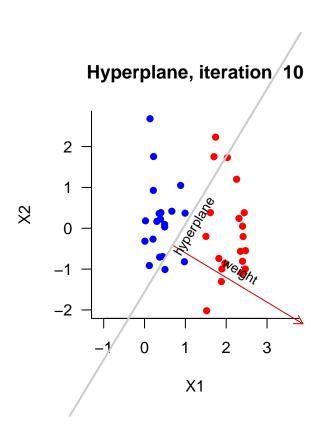


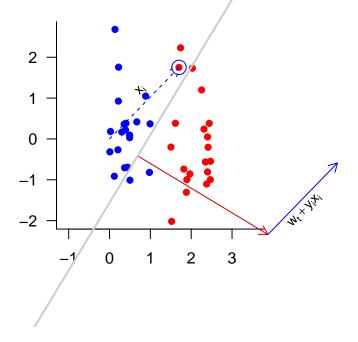


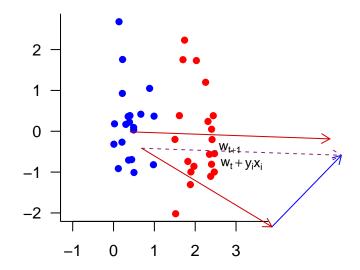


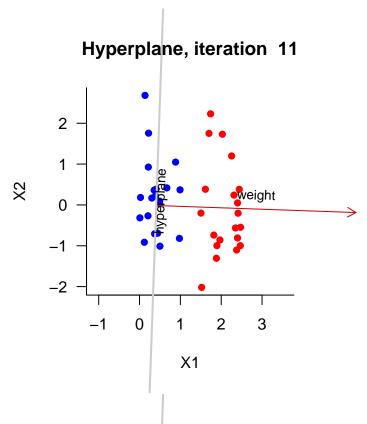




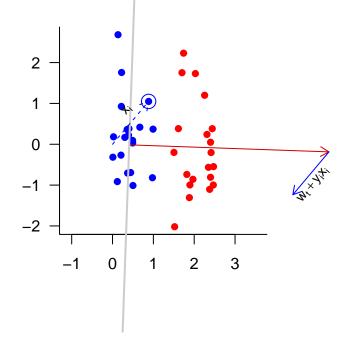


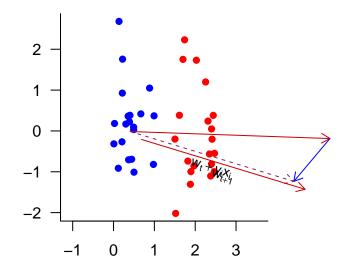


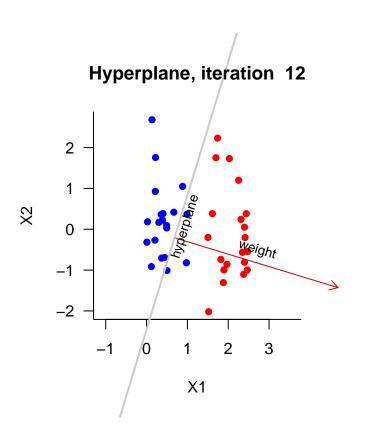


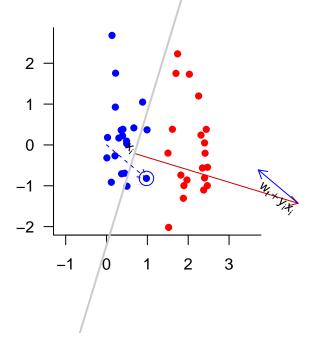


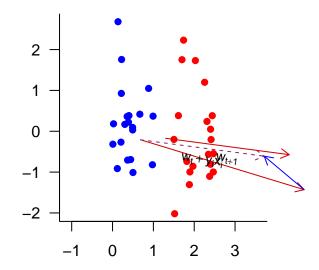


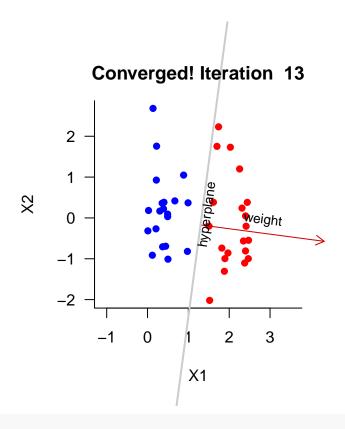












```
p <- par()$usr</pre>
```

Summary of 13 iterations:

```
library(tidyverse)
library(scales)
save(weight_matrix, file = "w.rds")
colnames(X) <- c("X0", "X1", "X2")</pre>
x <- as.data.frame(X) %>% mutate(Y = Y)
df <- data.frame(slope = apply(weight_matrix, 1, slope),</pre>
                  intercept = apply(weight_matrix, 1, intercept))
df <- df %>%
  mutate(iteration = as.factor(1:nrow(df)))
\# need to split up x since ggplot2 can only handle one color scale, and I need it for geom\_abline
group1 <- x[Y == -1,]
group2 \leftarrow x[Y == 1,]
purples <- brewer_pal(palette = "Purples")(9)</pre>
ggplot(group1, aes(X1, X2)) +
  geom_point(size = 2, color = "blue") +
 geom_point(data = group2, size = 2, color = "red") +
```

