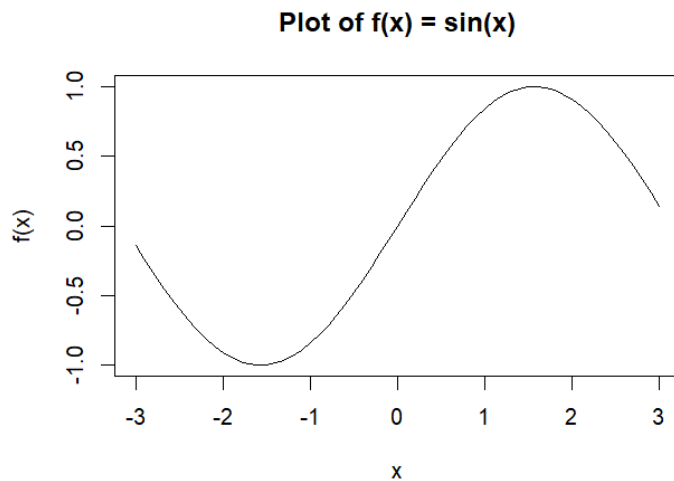


## Question 1

```
# Generate x values in the interval (-3, 3) with a step size of 0.1
x <- seq(-3, 3, 0.1)

# Compute the corresponding y values using the sin function
y <- sin(x)

# Plot the function
plot(x, y, type = "l", xlab = "x", ylab = "f(x)", main = "Plot of f(x) = sin(x)")
```



## Question 2

```
# Probability of suffering from the disease
p <- 0.2

# Number of workers
n <- 6

# Calculate the probability of four or more workers suffering from the disease
prob <- sum(dbinom(4:n, size = n, prob = p))

# Print the result
cat("The probability that four or more workers will suffer from the disease:", prob)
```

### Output:

The probability that four or more workers will suffer from the disease: 0.01696

### Question 3

*# Function to compute GCD using recursion*

```
gcd_recursive <- function(a, b) {  
  if (b == 0) {  
    return(a) # Base case: GCD is a when b is 0  
  } else {  
    return(gcd_recursive(b, a %% b)) # Recursive call with updated values  
  }  
}
```

*# Read input numbers from the user*

```
a <- as.integer(readline("Enter the first number: "))  
b <- as.integer(readline("Enter the second number: "))
```

*# Compute the GCD using recursion*

```
result <- gcd_recursive(a, b)
```

*# Print the result*

```
cat("The GCD of", a, "and", b, "is", result)
```

```
> source("~/active-rstudio-document")
```

```
Enter the first number: 12
```

```
Enter the second number: 15
```

```
The GCD of 12 and 15 is 3
```

### Question 4

*# Define the data points*

```
data <- matrix(c(2,10,2,5,8,4,5,8,7,5,6,4,1,2,4,9), ncol = 2, byrow = TRUE)
```

*# Define the initial centroids*

```
initial_centroids <- matrix(c(2,10,5,8,1,2), ncol = 2, byrow = TRUE)
```

*# Perform k-means clustering*

```
kmeans_result <- kmeans(data, centers = initial_centroids)
```

*# Get the cluster assignments*

```
cluster_assignments <- kmeans_result$cluster
```

*# Get the final centroids*

```
final_centroids <- kmeans_result$centers  
# Print the cluster assignments and final centroids  
cat("Cluster Assignments:")  
print(cluster_assignments)  
cat("\nFinal Centroids:")  
print(final_centroids)
```

```
Cluster Assignments:[1] 1 3 2 1 2 2 3 1
```

```
Final Centroids:      [,1]      [,2]  
1 3.666667 9.000000  
2 7.000000 4.333333  
3 1.500000 3.500000
```