

Advertising_EDA_Report

2025-01-30

Libraries and Imports

Data Import

```
#load staging data from SQL analysis
path <- "C:\\Users\\jbeas\\OneDrive\\Desktop\\Projects\\Advertising\\ad_staging.csv"
orig <- read.csv(path)

#additional stats from SQL analysis
stats_path <- "C:\\Users\\jbeas\\OneDrive\\Desktop\\Projects\\Advertising\\summary_stats.csv"
freq_path <- "C:\\Users\\jbeas\\OneDrive\\Desktop\\Projects\\Advertising\\freq_dist.csv"
mvals_path <- "C:\\Users\\jbeas\\OneDrive\\Desktop\\Projects\\Advertising\\missing_vals.csv"

data_stats <- read.csv(stats_path)
data_freq <- read.csv(freq_path)
data_mvals <- read.csv(mvals_path)

# verify data imported
head(orig)
```

	Ad_ID	Ad_Type	Visual_Complexity	Clicks	Time_Spent	Engagement_Score	Age_Group
## 1	1	AR	High	238	41	52	18-24
## 2	2	2D	Medium	116	44	87	35-44
## 3	3	AR	Medium	300	23	90	25-34
## 4	4	3D	High	65	120	61	18-24
## 5	5	AR	Low	92	65	93	25-34
## 6	6	2D	High	273	63	80	25-34

	Gender	Device_Type	Conversion_Rate	Bounce_Rate	CTR	Frame_Data
## 1	F	Desktop	8.72	25.92	5.09	frame_5
## 2	M	Mobile	7.99	9.05	9.02	frame_5
## 3	F	Mobile	1.65	11.32	7.57	frame_10
## 4	M	Desktop	6.22	39.11	3.64	frame_4
## 5	F	Tablet	8.31	15.94	14.97	frame_8
## 6	M	Desktop	8.06	25.38	2.92	frame_4

	User_Movement_Data	Age_Group_Numeric	Movement_Numeric
## 1	no movement		1
## 2	gaze, movement		3
## 3	movement only		2
## 4	movement only		2
## 5	no movement		1
## 6	no movement		1

```
## Visual_Complexity_Numeric Ad_Type_Numeric
## 1 3 3
## 2 2 1
## 3 2 3
## 4 3 2
## 5 1 3
## 6 3 1
```

```
#create copy of data
data <- orig
```

EDA

Summary Stats

```
#check for missing vals
data_mvals
```

```
## Total_rows Missing_Ad_ID Missing_Ad_Type Missing_Visual_Complexity
## 1 1000 0 0 0
## Missing_Clicks Missing_Time_Spent Missing_Engagement_Score Missing_Age_Group
## 1 0 0 0 0
## Missing_Gender Missing_Device_Type Missing_Conversion_Rate
## 1 0 0 0
## Missing_Bounce_Rate Missing_CTR Missing_Frame_Data Missing_User_Movement_Data
## 1 0 0 0 0
```

```
#summary stats
data_stats
```

	Metric	Mean	Std_Dev	Min_Value	Q1	Median	Q3	Max_Value
## 1	Bounce_Rate	22.73	9.91	5.00	14.21	22.90	31.48	39.99
## 2	Clicks	154.91	82.56	10.00	85.00	156.00	224.00	300.00
## 3	Conversion_Rate	5.47	2.60	1.00	3.17	5.48	7.71	9.98
## 4	CTR	7.97	3.93	1.03	4.64	8.03	11.09	14.99
## 5	Engagement_Score	74.87	14.80	50.00	62.00	75.00	88.00	100.00
## 6	Time_Spent	65.81	31.86	10.00	39.00	65.00	94.00	120.00

```
data_freq
```

	Category	Value	Frequency	Percentage
## 1	Ad_Type	3D	418	41.80%
## 2	Ad_Type	AR	380	38.00%
## 3	Ad_Type	2D	202	20.20%
## 4	Age_Group	25-34	410	41.00%
## 5	Age_Group	18-24	291	29.10%
## 6	Age_Group	35-44	210	21.00%
## 7	Age_Group	45-54	59	5.90%
## 8	Age_Group	55+	30	3.00%

## 9	Device_Type	Mobile	601	60.10%
## 10	Device_Type	Desktop	296	29.60%
## 11	Device_Type	Tablet	103	10.30%
## 12	Gender	F	503	50.30%
## 13	Gender	M	497	49.70%
## 14	User_Movement_Data	movement only	403	40.30%
## 15	User_Movement_Data	gaze, movement	384	38.40%
## 16	User_Movement_Data	no movement	213	21.30%
## 17	Visual_Complexity	Medium	504	50.40%
## 18	Visual_Complexity	High	293	29.30%
## 19	Visual_Complexity	Low	203	20.30%

```

#create copies for plotting
stats <- data_stats
freq <- data_freq

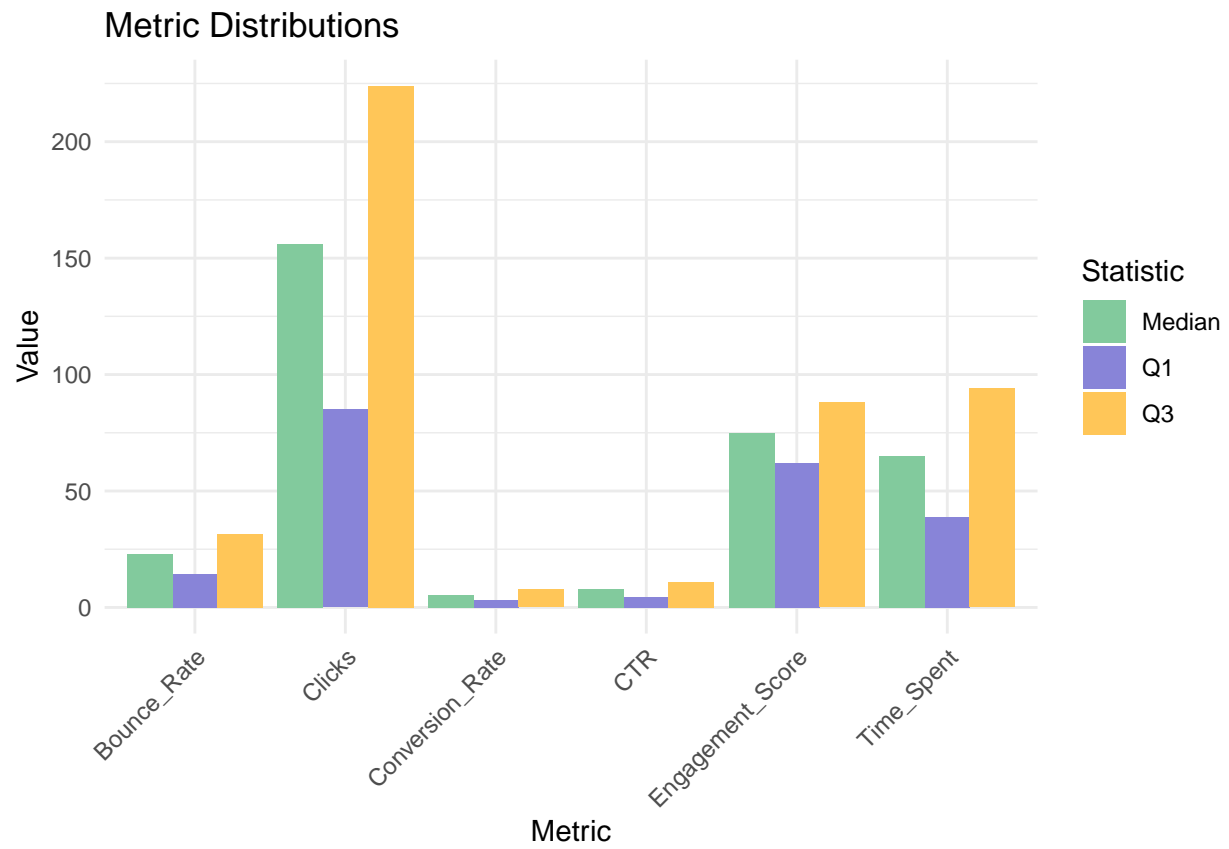
#metric distribution
metrics_long <- data_stats %>%
  dplyr::select(dplyr::all_of(c("Metric", "Q1", "Median", "Q3"))) %>%
  tidyr::pivot_longer(
    cols = dplyr::all_of(c("Q1", "Median", "Q3")),
    names_to = "Statistic",
    values_to = "Value"
  )

p1 <- ggplot(metrics_long, aes(x = Metric, y = Value, fill = Statistic)) +
  geom_bar(stat = "identity", position = "dodge") +
  theme_minimal() +
  labs(title = "Metric Distributions",
       x = "Metric",
       y = "Value") +
  theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
  scale_fill_manual(values = c("#82CA9D", "#8884D8", "#FFC658"))

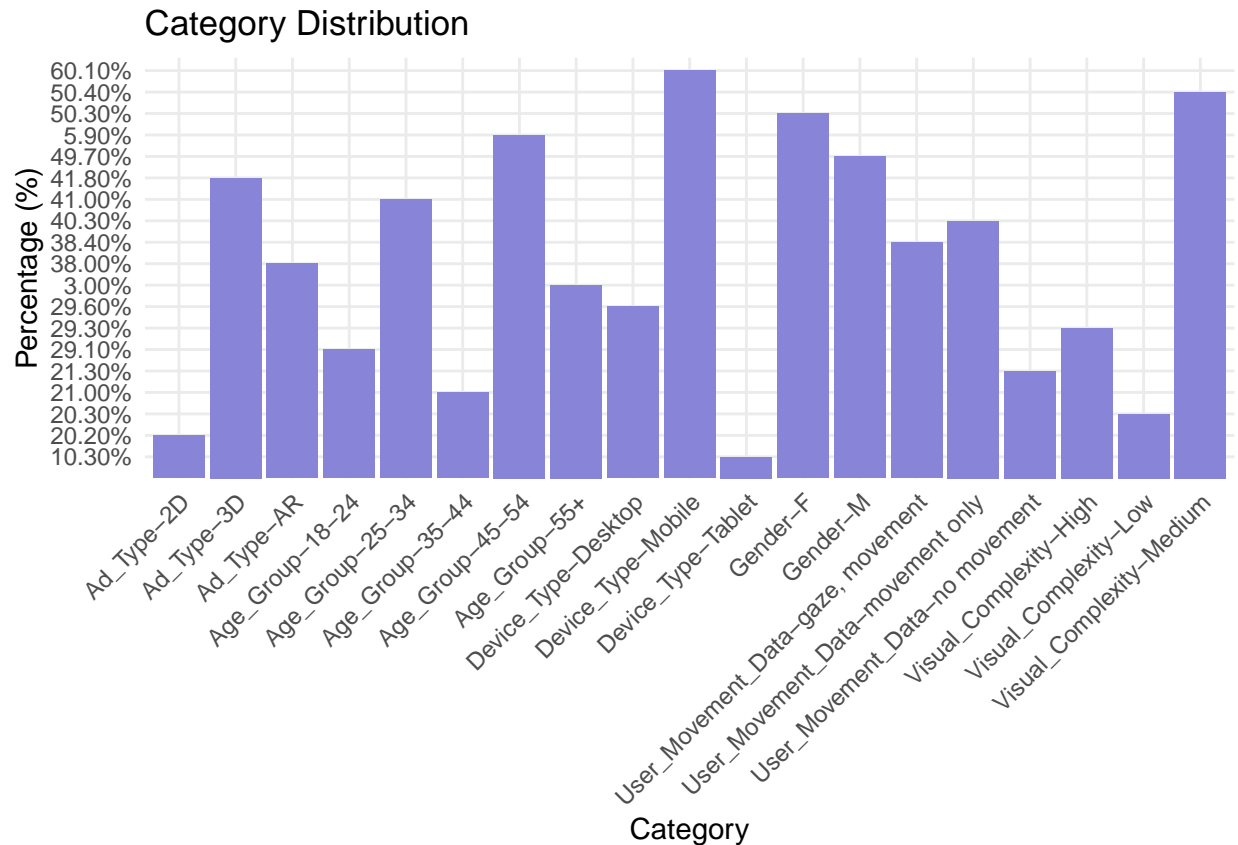
#frequency distribution
p2 <- ggplot(freq, aes(x = paste(Category, Value, sep = "-"), y = Percentage)) +
  geom_bar(stat = "identity", fill = "#8884D8") +
  theme_minimal() +
  labs(title = "Category Distribution",
       x = "Category",
       y = "Percentage (%)") +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))

# Print plots
print(p1)

```



```
print(p2)
```



```
#save
ggsave("visualizations\\metric_distributions.png", p1, width = 10, height = 6,)
ggsave("visualizations\\category_distribution.png", p2, width = 10, height = 6)
```

At a glance:

- Bounce Rate, Conversion Rate, CTR and Engagement Score all have fairly even distributions
- Clicks and Time Spent have a much more variance (higher Q1, Q3 values compared to median)
- We can use 'Q' values as a *'performance benchmark'* for future campaigns
 - If a campaign's metrics fall *below the Q1 value*, it is **underperforming**
 - If a campaign's metrics fall *above the Q3 value*, it is **successful**
 - anything in between means the ad is performing as expected

Areas to Investigate:

- **METRICS:** Clicks, Time Spent, Engagement Score, Conversion Rate, Bounce Rate, CTR
- 3D and AR ads are the most common Ad Types, but are they more successful than 2D?
- What impact does Visual Complexity have on our performance metrics?
- What is the relationship between Visual Complexity and User Movement, does this effect our metrics?

- What characteristics of an ad yields the highest metrics?
 - What characteristics effect Clicks, Time Spent Conversion Rate Bounce Rate and CTR the most?

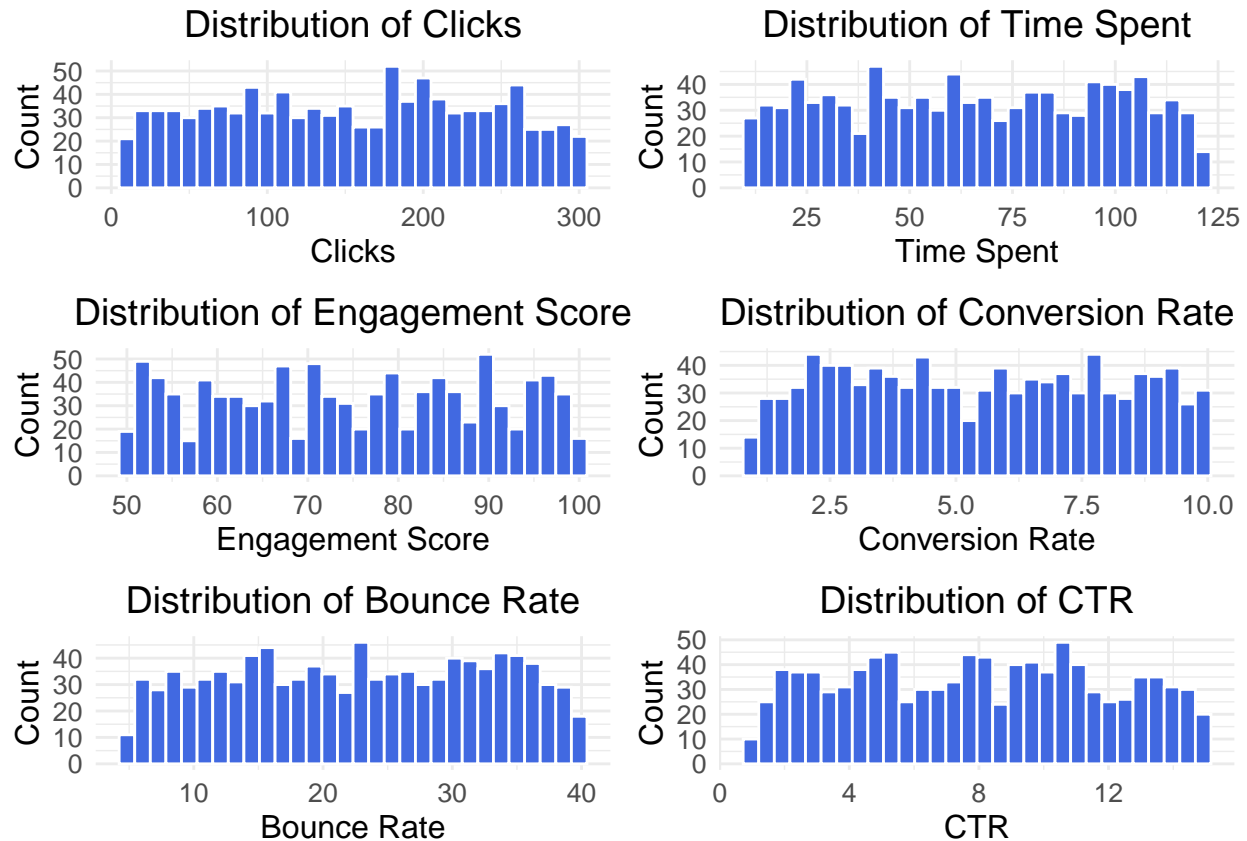
Bar Plots

```
#numerical data
# Create a barplot function
create_barplot <- function(data, column_name) {
  ggplot(data, aes(x = !!sym(column_name))) +
    geom_histogram(fill = "#4169E1", color = "white", bins = 30) +
    theme_minimal() +
    labs(
      title = paste("Distribution of", gsub("_", " ", column_name)),
      x = gsub("_", " ", column_name),
      y = "Count"
    ) +
    theme(
      plot.title = element_text(hjust = 0.5, size = 14),
      axis.title = element_text(size = 12),
      axis.text = element_text(size = 10)
    )
}

numerical_columns <- c(
  "Clicks",
  "Time_Spent",
  "Engagement_Score",
  "Conversion_Rate",
  "Bounce_Rate",
  "CTR"
)

plots <- list()
for (col in numerical_columns) {
  plots[[col]] <- create_barplot(data, col)
}

#plot
grid.arrange(
  plots[["Clicks"]],
  plots[["Time_Spent"]],
  plots[["Engagement_Score"]],
  plots[["Conversion_Rate"]],
  plots[["Bounce_Rate"]],
  plots[["CTR"]],
  ncol = 2
)
```



```
#save
ggsave("visualizations\\numerical_distributions.png", width = 10, height = 8)

# Function to calculate goodness-of-fit statistics
calculate_gof <- function(x, dist_name, params) {
  # Ensure x is positive for certain distributions
  x_adj <- if(min(x) <= 0) x - min(x) + 0.01 else x

  # Add small random noise to break ties
  x_jitter <- x + rnorm(length(x), 0, sd(x)/1000)
  x_adj_jitter <- x_adj + rnorm(length(x_adj), 0, sd(x_adj)/1000)

  # Kolmogorov-Smirnov test with jittered data
  if(dist_name == "normal") {
    ks_test <- suppressWarnings(ks.test(x_jitter, "pnorm", mean = params$mean, sd = params$sd))
  } else if(dist_name == "lognormal") {
    ks_test <- suppressWarnings(ks.test(x_adj_jitter, "plnorm", meanlog = params$meanlog, sdlog = params$sdlog))
  } else if(dist_name == "weibull") {
    ks_test <- suppressWarnings(ks.test(x_adj_jitter, "pweibull", shape = params$shape, scale = params$scale))
  } else if(dist_name == "uniform") {
    ks_test <- suppressWarnings(ks.test(x_jitter, "punif", min = params$min, max = params$max))
  }

  # Return test statistics
  return(list(
    ks_stat = ks_test$statistic,

```

```

    ks_p = ks_test$p.value
  })
}

# Enhanced distribution analysis function
analyze_distribution <- function(data, column_name) {
  # Get the data from the column
  x <- data[[column_name]]

  # Basic error checking
  if(is.null(x)) {
    stop(paste("Column", column_name, "not found in data"))
  }

  # Calculate basic statistics
  mu <- mean(x, na.rm = TRUE)
  sigma <- sd(x, na.rm = TRUE)

  # Create sequence for curves
  x_range <- seq(min(x, na.rm = TRUE), max(x, na.rm = TRUE), length.out = 200)

  # Fit distributions and calculate goodness-of-fit
  tryCatch({
    # Normal
    normal_params <- list(mean = mu, sd = sigma)
    normal_y <- dnorm(x_range, mean = mu, sd = sigma)
    normal_gof <- calculate_gof(x, "normal", normal_params)

    # Log-normal
    x_lognorm <- if(min(x) <= 0) x - min(x) + 0.01 else x
    lognorm_fit <- fitdistr(x_lognorm, "lognormal")
    lognorm_params <- list(
      meanlog = lognorm_fit$estimate[1],
      sdlog = lognorm_fit$estimate[2]
    )
    lognorm_y <- dlnorm(x_range, meanlog = lognorm_params$meanlog,
                       sdlog = lognorm_params$sdlog)
    lognorm_gof <- calculate_gof(x, "lognormal", lognorm_params)

    # Weibull
    weibull_fit <- fitdistr(x - min(x) + 0.01, "weibull")
    weibull_params <- list(
      shape = weibull_fit$estimate[1],
      scale = weibull_fit$estimate[2]
    )
    weibull_y <- dweibull(x_range - min(x_range) + 0.01,
                        shape = weibull_params$shape,
                        scale = weibull_params$scale)
    weibull_gof <- calculate_gof(x, "weibull", weibull_params)

    # Uniform
    uniform_params <- list(min = min(x), max = max(x))
    uniform_y <- dunif(x_range, min = min(x), max = max(x))
  }, error = function(e) {
    stop(e$message)
  })
}

```



```

uniform_gof <- calculate_gof(x, "uniform", uniform_params)

# Create dataframe for all distributions
dist_df <- data.frame(
  x = rep(x_range, 4),
  y = c(normal_y, lognorm_y, weibull_y, uniform_y),
  Distribution = factor(rep(c("Normal", "Log-normal", "Weibull", "Uniform"),
    each = length(x_range)))
)

# Find best fitting distribution
gof_stats <- data.frame(
  Distribution = c("Normal", "Log-normal", "Weibull", "Uniform"),
  KS_stat = c(normal_gof$ks_stat, lognorm_gof$ks_stat,
    weibull_gof$ks_stat, uniform_gof$ks_stat),
  P_value = c(normal_gof$ks_p, lognorm_gof$ks_p,
    weibull_gof$ks_p, uniform_gof$ks_p)
)
best_fit <- gof_stats[which.max(gof_stats$P_value), "Distribution"]

# Create goodness-of-fit results dataframe
gof_results <- data.frame(
  Distribution = c("Normal", "Log-normal", "Weibull", "Uniform"),
  KS_stat = c(normal_gof$ks_stat, lognorm_gof$ks_stat,
    weibull_gof$ks_stat, uniform_gof$ks_stat),
  P_value = c(normal_gof$ks_p, lognorm_gof$ks_p,
    weibull_gof$ks_p, uniform_gof$ks_p)
)

# Print results to console
cat("\nGoodness-of-fit Test Results for", column_name, "\n")
cat("=====\n")
print(gof_results)
cat("\nBest fitting distribution:", best_fit, "\n")

# Create formatted text for plot annotation
gof_text <- paste(
  "Goodness-of-fit Statistics",
  sprintf("%-12s D      p-value", "Distribution"),
  sprintf("%-12s %.3f  %.2e", "Normal", normal_gof$ks_stat, normal_gof$ks_p),
  sprintf("%-12s %.3f  %.2e", "Log-normal", lognorm_gof$ks_stat, lognorm_gof$ks_p),
  sprintf("%-12s %.3f  %.2e", "Weibull", weibull_gof$ks_stat, weibull_gof$ks_p),
  sprintf("%-12s %.3f  %.2e", "Uniform", uniform_gof$ks_stat, uniform_gof$ks_p),
  sprintf("\nBest fit: %s", best_fit),
  sep = "\n"
)

# Create the plot with error handling
p <- tryCatch({
  ggplot() +
    # Histogram with density
    geom_histogram(data = data.frame(x = x), aes(x = x, y = ..density..),
      fill = "grey80", color = "white", alpha = 0.7,

```

```

        bins = 30) +
# Add theoretical distributions
geom_line(data = dist_df,
          aes(x = x, y = y, color = Distribution, linetype = Distribution),
          linewidth = 1) +
# Customize colors and linetypes
scale_color_manual(values = c("Normal" = "#FF4444",
                              "Log-normal" = "#4169E1",
                              "Weibull" = "#228B22",
                              "Uniform" = "#FFA500")) +
scale_linetype_manual(values = c("Normal" = "solid",
                                 "Log-normal" = "dashed",
                                 "Weibull" = "dotdash",
                                 "Uniform" = "dotted")) +

# Add statistics box
# Add statistics in a text box using annotate
annotate("text",
        x = min(x) + (max(x) - min(x)) * 0.2, # Position at 20% of x range
        y = max(density(x)$y) * 0.8,          # Position at 80% of max height
        label = gof_text,
        hjust = 0,
        vjust = 1,
        size = 3,
        family = "mono",
        color = "black",
        box.color = "black",
        box.padding = unit(0.5, "lines"),
        box.margin = unit(0.5, "lines")) +

# Theme and labels
theme_minimal() +
labs(
  title = paste("Distribution Analysis of", gsub("_", " ", column_name)),
  subtitle = "Comparing Theoretical Distributions with Goodness-of-Fit Tests",
  x = gsub("_", " ", column_name),
  y = "Density"
) +
theme(
  plot.title = element_text(hjust = 0.5, size = 14),
  plot.subtitle = element_text(hjust = 0.5, size = 10),
  axis.title = element_text(size = 12),
  axis.text = element_text(size = 10),
  legend.position = "right",
  legend.title = element_text(size = 10),
  legend.text = element_text(size = 9)
)

}, error = function(e) {
  message("Error in plot creation: ", e$message)
  return(NULL)
})

if (!is.null(p)) {
  return(p)
}

```

```

} else {
  # If plot creation failed, create a simple error plot
  return(ggplot() +
    annotate("text", x = 0.5, y = 0.5,
      label = "Error creating distribution plot",
      size = 5) +
    theme_void() +
    xlim(0, 1) + ylim(0, 1))
}

}, error = function(e) {
  message("Error in distribution fitting: ", e$message)
  return(NULL)
})
}

# Function to plot one column
plot_one <- function(data, column_name) {
  p <- analyze_distribution(data, column_name)
  if (!is.null(p)) {
    print(p)
    ggsave(
      filename = paste0("visualizations\\distribution_analysis_", tolower(column_name), ".png"),
      plot = p,
      width = 12,
      height = 7,
      dpi = 300
    )
  }
}

```

```
plot_one(data, "Clicks")
```

```

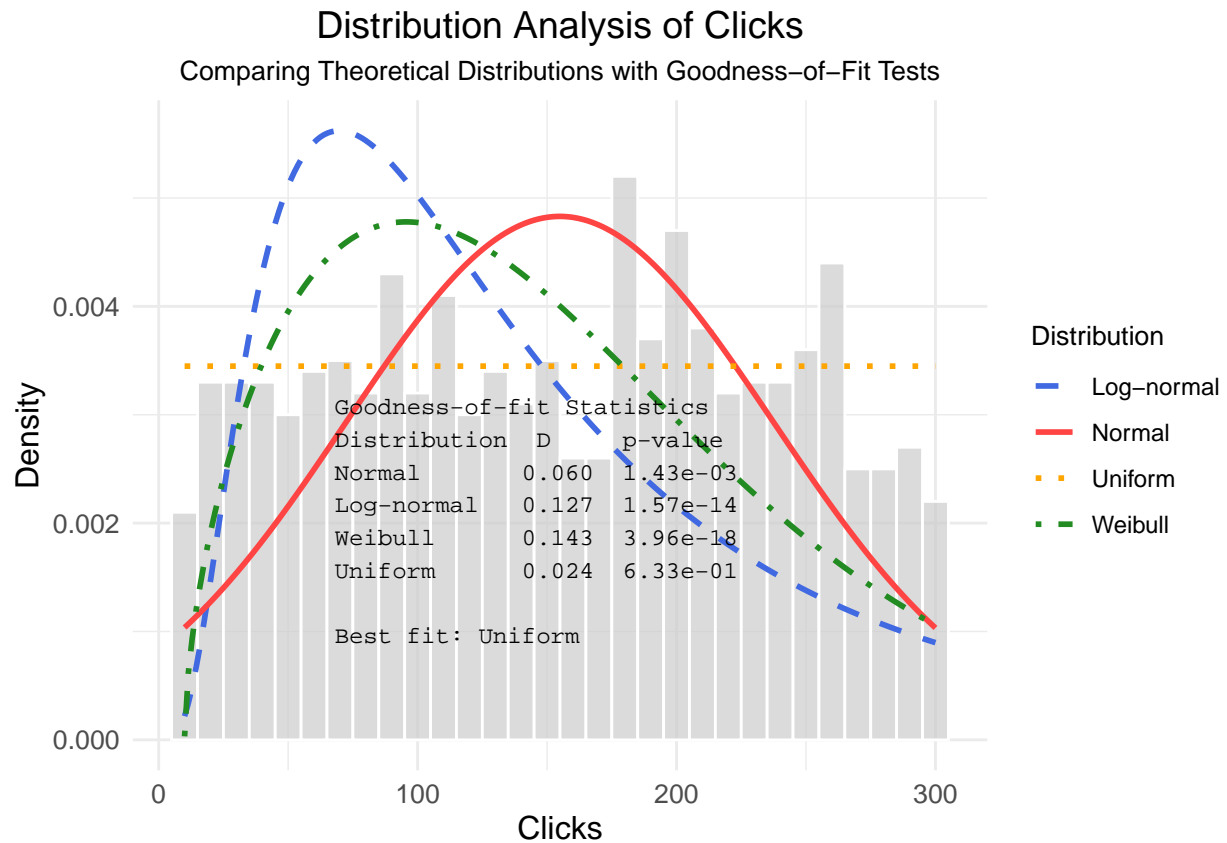
##
## Goodness-of-fit Test Results for Clicks
## =====
##   Distribution    KS_stat      P_value
## 1      Normal 0.06017627 1.431230e-03
## 2  Log-normal 0.12743358 1.569417e-14
## 3    Weibull 0.14276566 3.957503e-18
## 4     Uniform 0.02360576 6.330958e-01
##
## Best fitting distribution: Uniform

## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'

## Warning: The dot-dot notation ('..density..') was deprecated in ggplot2 3.4.0.
## i Please use 'after_stat(density)' instead.

```

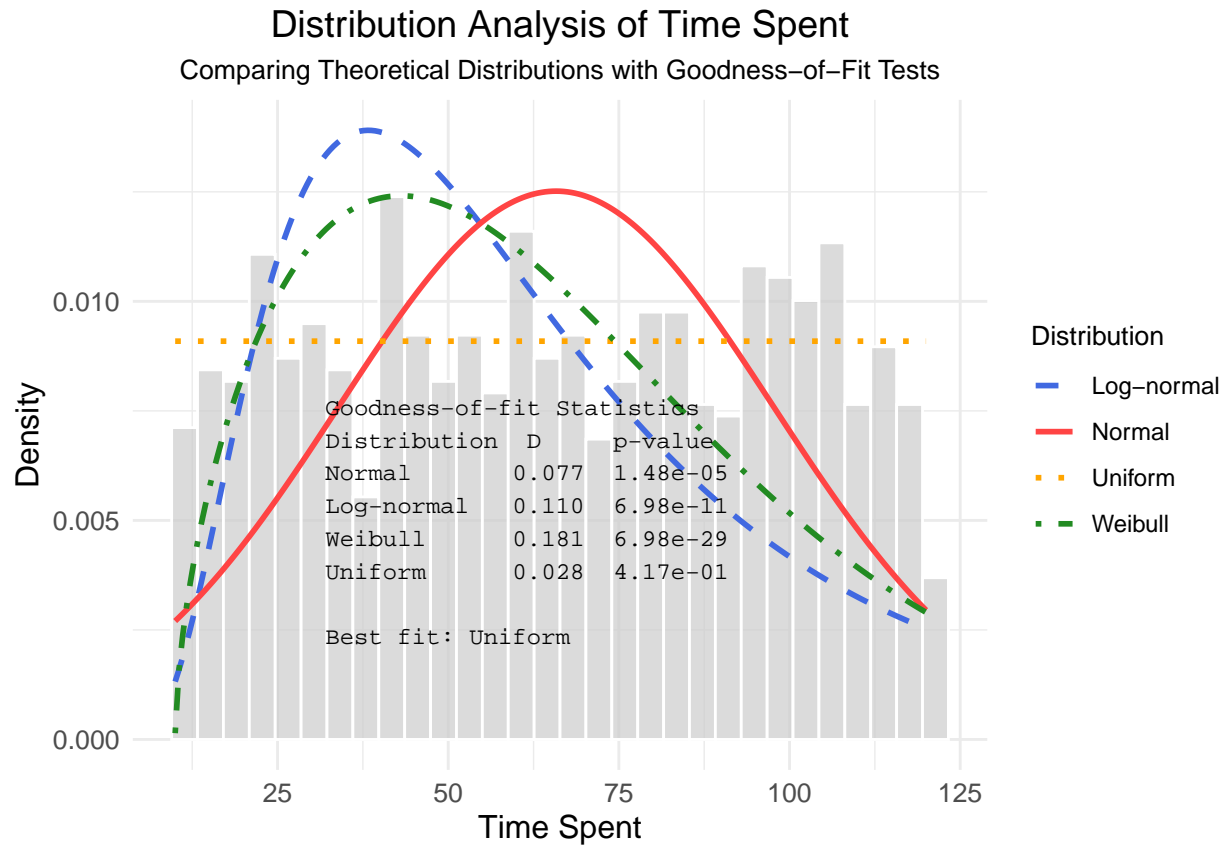
```
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```



```
plot_one(data, "Time_Spent")
```

```
##
## Goodness-of-fit Test Results for Time_Spent
## =====
##   Distribution    KS_stat    P_value
## 1      Normal 0.07685794 1.479623e-05
## 2  Log-normal 0.10972501 6.975675e-11
## 3      Weibull 0.18100459 6.978216e-29
## 4      Uniform 0.02791024 4.172009e-01
##
## Best fitting distribution: Uniform

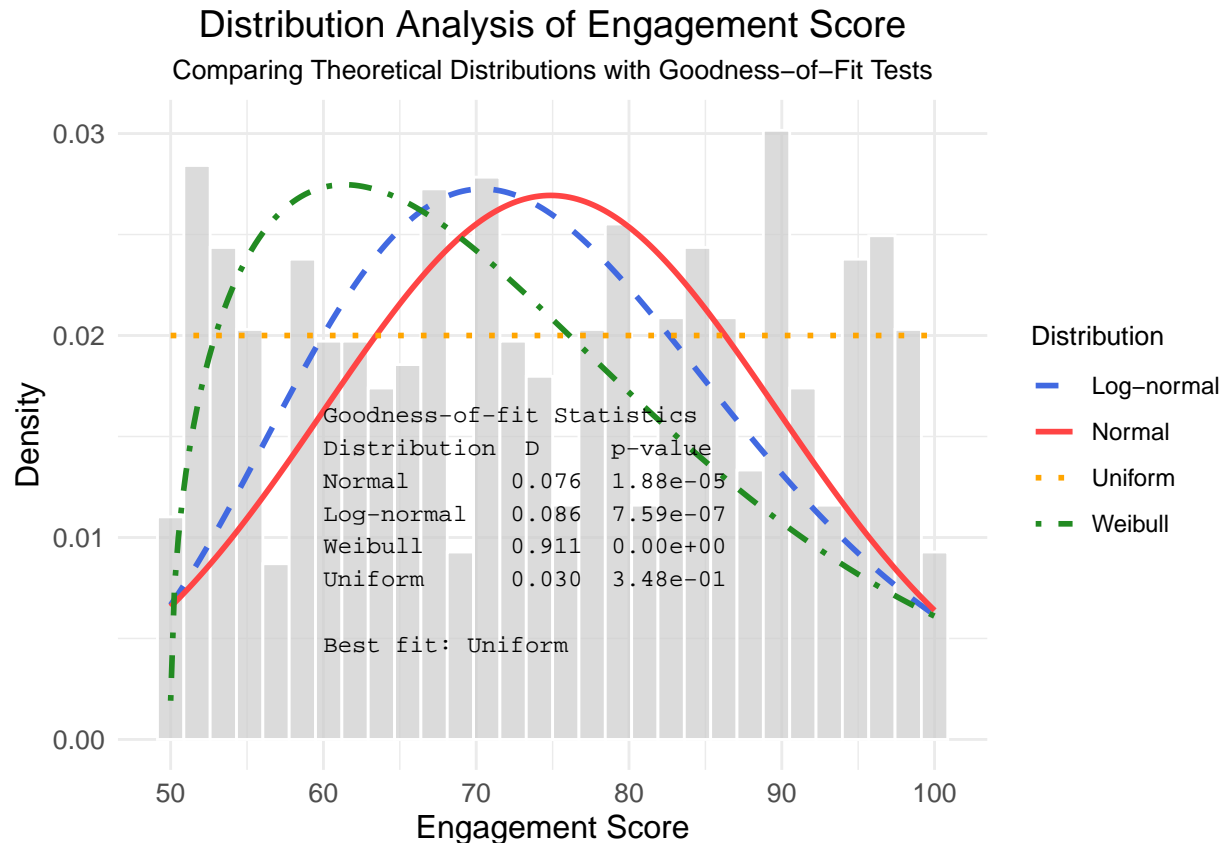
## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'
```



```
plot_one(data, "Engagement_Score")
```

```
##
## Goodness-of-fit Test Results for Engagement_Score
## =====
##   Distribution    KS_stat    P_value
## 1      Normal 0.07607241 1.881464e-05
## 2  Log-normal 0.08597759 7.590914e-07
## 3      Weibull 0.91107447 0.000000e+00
## 4      Uniform 0.02951776 3.482603e-01
##
## Best fitting distribution: Uniform

## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'
```

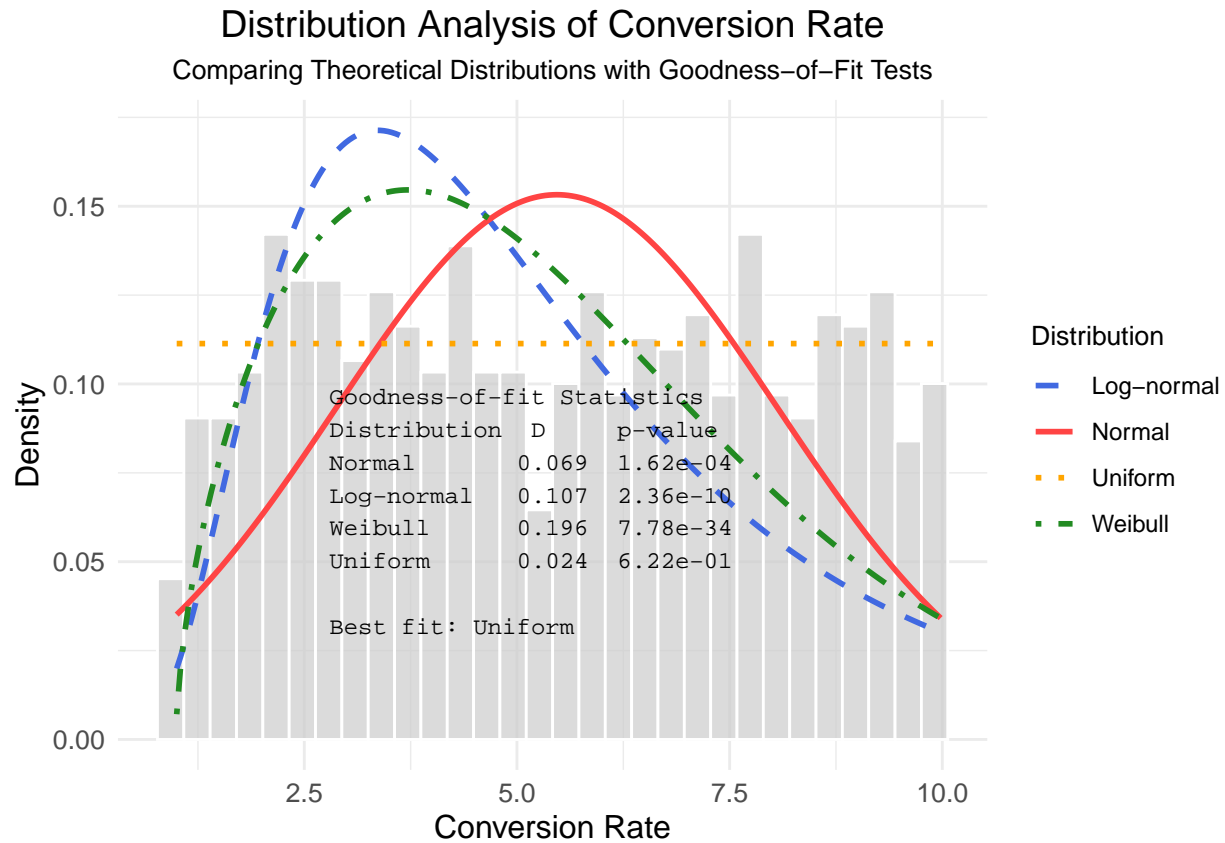


```
plot_one(data, "Conversion_Rate")
```

```
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
```

```
##
## Goodness-of-fit Test Results for Conversion_Rate
## =====
##   Distribution    KS_stat    P_value
## 1      Normal 0.06862982 1.621552e-04
## 2  Log-normal 0.10691421 2.357737e-10
## 3      Weibull 0.19612376 7.783288e-34
## 4      Uniform 0.02381761 6.218180e-01
##
## Best fitting distribution: Uniform
```

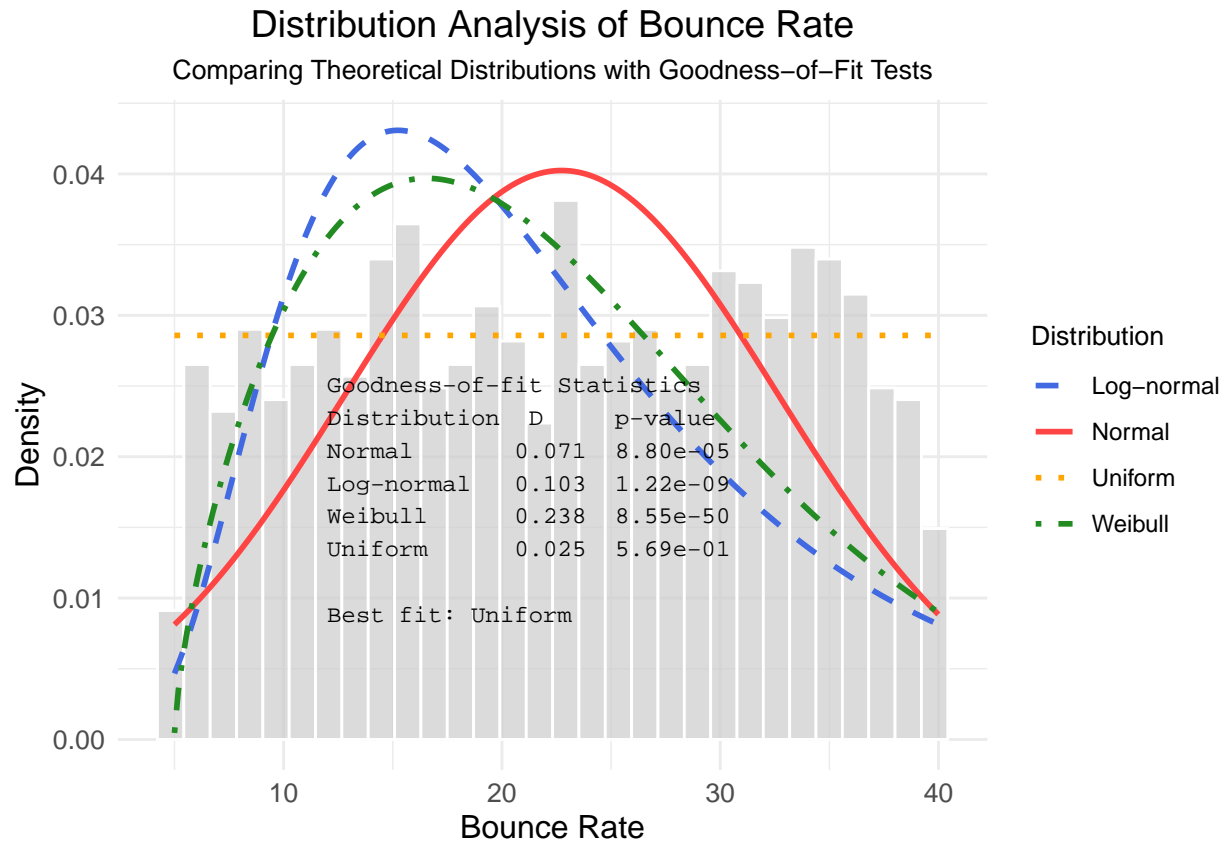
```
## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'
```



```
plot_one(data, "Bounce_Rate")
```

```
##
## Goodness-of-fit Test Results for Bounce_Rate
## =====
##   Distribution    KS_stat    P_value
## 1      Normal 0.07082074 8.801474e-05
## 2  Log-normal 0.10300658 1.216166e-09
## 3      Weibull 0.23840827 8.545416e-50
## 4      Uniform 0.02482560 5.686393e-01
##
## Best fitting distribution: Uniform

## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'
```

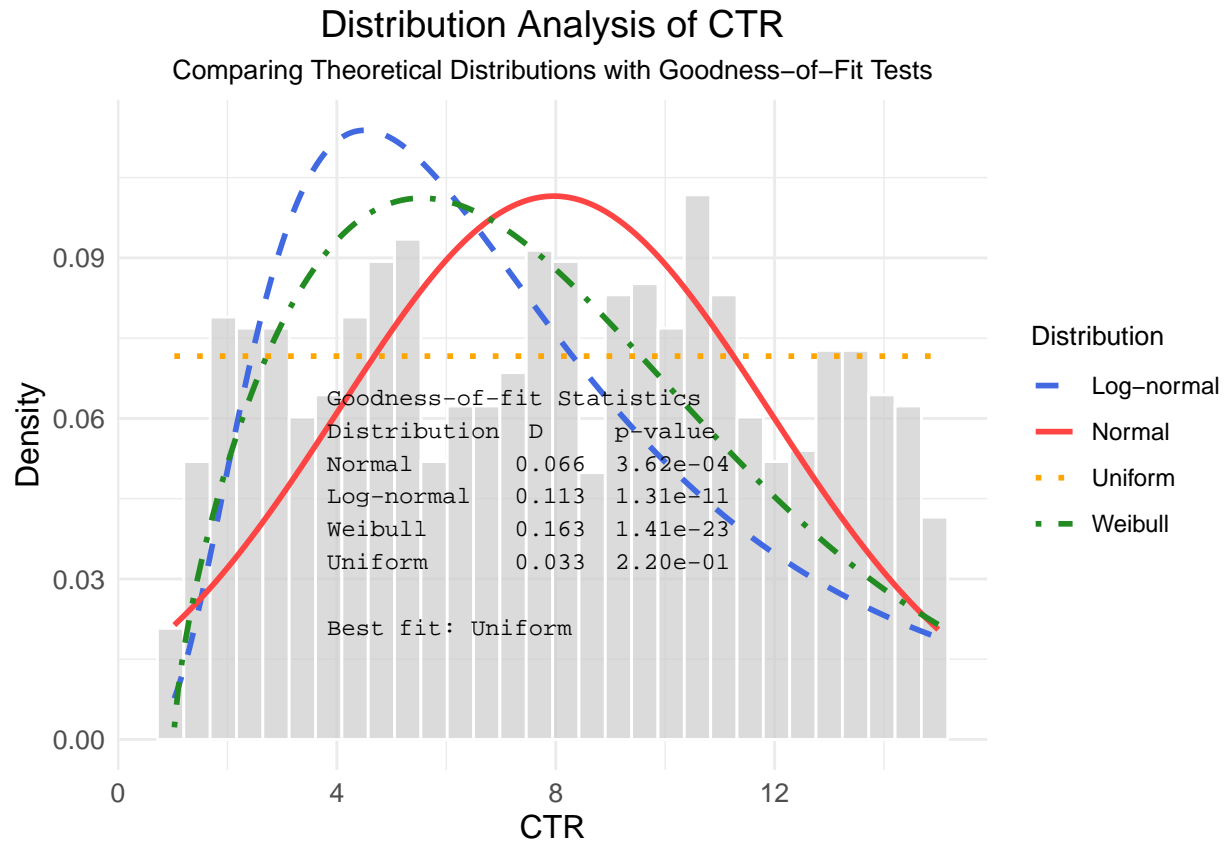


```
plot_one(data, "CTR")
```

```
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
## Warning in densfun(x, parm[1], parm[2], ...): NaNs produced
```

```
##
## Goodness-of-fit Test Results for CTR
## =====
##   Distribution    KS_stat      P_value
## 1      Normal 0.06563953 3.619707e-04
## 2  Log-normal 0.11346766 1.312278e-11
## 3      Weibull 0.16326103 1.410967e-23
## 4      Uniform 0.03322416 2.196164e-01
##
## Best fitting distribution: Uniform
```

```
## Warning in annotate("text", x = min(x) + (max(x) - min(x)) * 0.2, y =
## max(density(x)$y) * : Ignoring unknown parameters: 'box.colour', 'box.padding',
## and 'box.margin'
```

As we see from our Distribution Analysis, our data is Uniformly Distributed

Correlation Plot

```
numerical_columns <- c("Clicks", "Time_Spent", "Engagement_Score",
                       "Conversion_Rate", "Bounce_Rate", "CTR")

numerical_data <- data %>%
  dplyr::select(dplyr::all_of(numerical_columns))

# Calculate correlation matrix
cor_matrix <- cor(numerical_data)

# Print matrix for inspection
print("Correlation Matrix:")

## [1] "Correlation Matrix:"

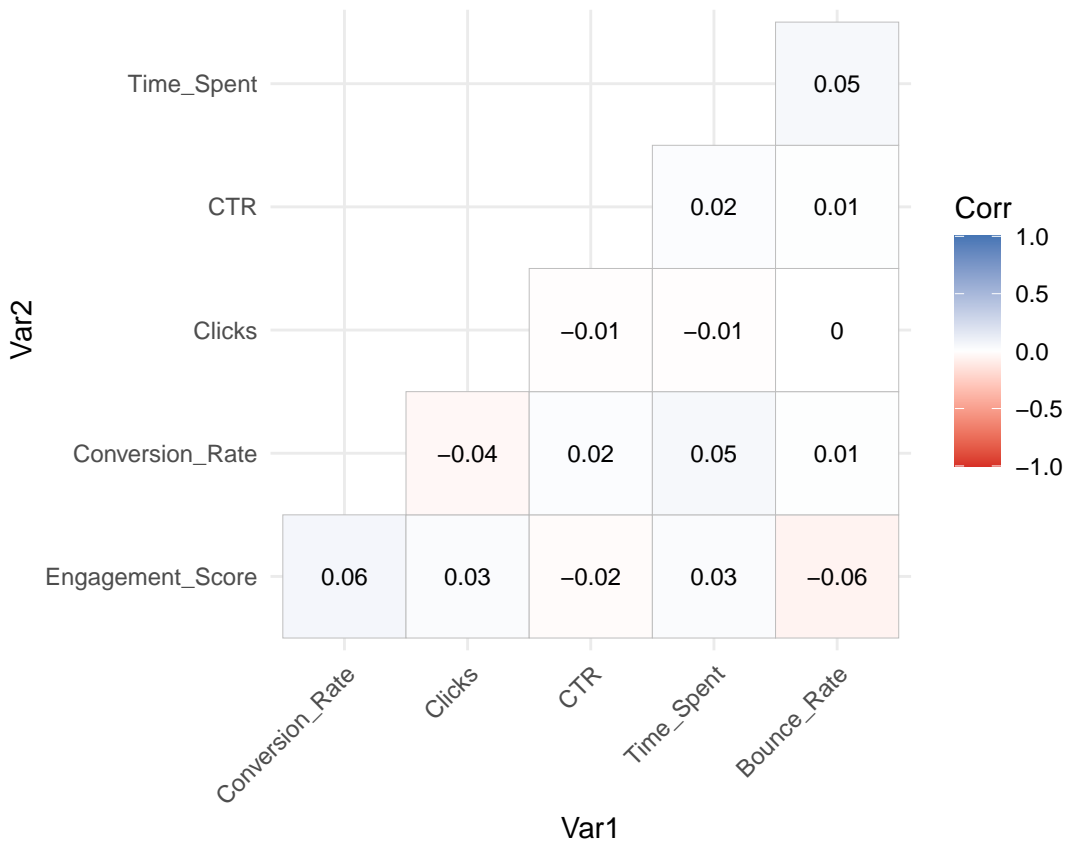
print(cor_matrix)

##           Clicks  Time_Spent Engagement_Score Conversion_Rate
## Clicks      1.000000000 -0.006126274      0.03056959  -0.044432411
## Time_Spent  -0.006126274  1.000000000      0.02564127   0.052200269
```

```
## Engagement_Score 0.030569585 0.025641273 1.00000000 0.055550416
## Conversion_Rate -0.044432411 0.052200269 0.05555042 1.000000000
## Bounce_Rate 0.004451510 0.045156588 -0.06488862 0.007921938
## CTR -0.006370837 0.015421482 -0.02109692 0.019890959
## Bounce_Rate CTR
## Clicks 0.004451510 -0.006370837
## Time_Spent 0.045156588 0.015421482
## Engagement_Score -0.064888619 -0.021096916
## Conversion_Rate 0.007921938 0.019890959
## Bounce_Rate 1.000000000 0.012645491
## CTR 0.012645491 1.000000000
```

```
# Create correlation plot
cor_plot <- ggcorrplot(cor_matrix,
  hc.order = TRUE,
  type = "lower",
  lab = TRUE,
  lab_size = 3,
  colors = c("#D73027", "white", "#4575B4")) +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))

# Display plot
print(cor_plot)
```



```

# Save plot
ggsave("visualizations/correlation_matrix.png",
      plot = cor_plot,
      width = 10,
      height = 8)

# Optional: Return correlation statistics
correlation_summary <- data.frame(
  Variable1 = character(),
  Variable2 = character(),
  Correlation = numeric(),
  stringsAsFactors = FALSE
)

```

Looking at the Barplots and Correlation matrix:

- Our data is not normally distributed, so linear modeling/analysis will not be that useful
- There are no strong correlations between any of the numerical columns
- We do not need to investigate the interactions between these columns further

Non-Linear Relationships

Ad Type vs. Engagement Score

Kruskall Test

- Null Hypothesis: 3D and AR ads are more effective than 2D ads
- Alternative Hypothesis: 3D and AR ads are not more successful than 2D ads

```

# 2. Kruskal-Wallis Test with detailed interpretation
e_score <- kruskal.test(Engagement_Score ~ Ad_Type, data = data)
c_score <- kruskal.test(Clicks ~ Ad_Type, data = data)
t_score <- kruskal.test(Time_Spent ~ Ad_Type, data = data)
ctr_score <- kruskal.test(CTR ~ Ad_Type, data = data)

# Create results dataframe with more detailed information
results <- data.frame(
  Metric = c("Engagement Score", "Clicks", "Time Spent", "CTR"),
  p_value = c(e_score$p.value, c_score$p.value, t_score$p.value, ctr_score$p.value),
  statistic = c(e_score$statistic, c_score$statistic, t_score$statistic, ctr_score$statistic)
)

# Add interpretation columns
results <- results %>%
  mutate(
    Significance = case_when(
      p_value < 0.01 ~ "Highly Significant",
      p_value < 0.05 ~ "Significant",
      TRUE ~ "Not Significant"
    )
  )

```

```

    ),
    Interpretation = case_when(
      p_value < 0.05 ~ "There are significant differences between ad types",
      TRUE ~ "No significant differences between ad types"
    )
  ) %>%
  mutate(
    p_value = round(p_value, 4),
    statistic = round(statistic, 2)
  )

# Print formatted results
print("Analysis of Ad Type Effects on Performance Metrics")

```

```
## [1] "Analysis of Ad Type Effects on Performance Metrics"
```

```
print("-----")
```

```
## [1] "-----"
```

```

for(i in 1:nrow(results)) {
  cat(sprintf("\nMetric: %s", results$Metric[i]))
  cat(sprintf("\n- P-value: %f", results$p_value[i]))
  cat(sprintf("\n- Chi-squared statistic: %f", results$statistic[i]))
  cat(sprintf("\n- Result: %s", results$Significance[i]))
  cat(sprintf("\n- Interpretation: %s\n", results$Interpretation[i]))
}

```

```

##
## Metric: Engagement Score
## - P-value: 0.213800
## - Chi-squared statistic: 3.090000
## - Result: Not Significant
## - Interpretation: No significant differences between ad types
##
## Metric: Clicks
## - P-value: 0.574300
## - Chi-squared statistic: 1.110000
## - Result: Not Significant
## - Interpretation: No significant differences between ad types
##
## Metric: Time Spent
## - P-value: 0.050900
## - Chi-squared statistic: 5.960000
## - Result: Not Significant
## - Interpretation: No significant differences between ad types
##
## Metric: CTR
## - P-value: 0.433200
## - Chi-squared statistic: 1.670000
## - Result: Not Significant
## - Interpretation: No significant differences between ad types

```

```

# Create summary statement
significant_metrics <- results$Metric[results$p_value < 0.05]
cat("\nSummary:\n")

##
## Summary:

if(length(significant_metrics) > 0) {
  cat("The following metrics show significant differences between ad types:\n")
  cat(paste("-", significant_metrics, collapse = "\n"))
  cat("\n\nThis suggests that ad type does influence these performance metrics.")
} else {
  cat("None of the metrics showed significant differences between ad types.\n")
  cat("This suggests that ad type may not be a determining factor in ad performance.")
}

## None of the metrics showed significant differences between ad types.
## This suggests that ad type may not be a determining factor in ad performance.

```

Boxplots

- Visualize our findings

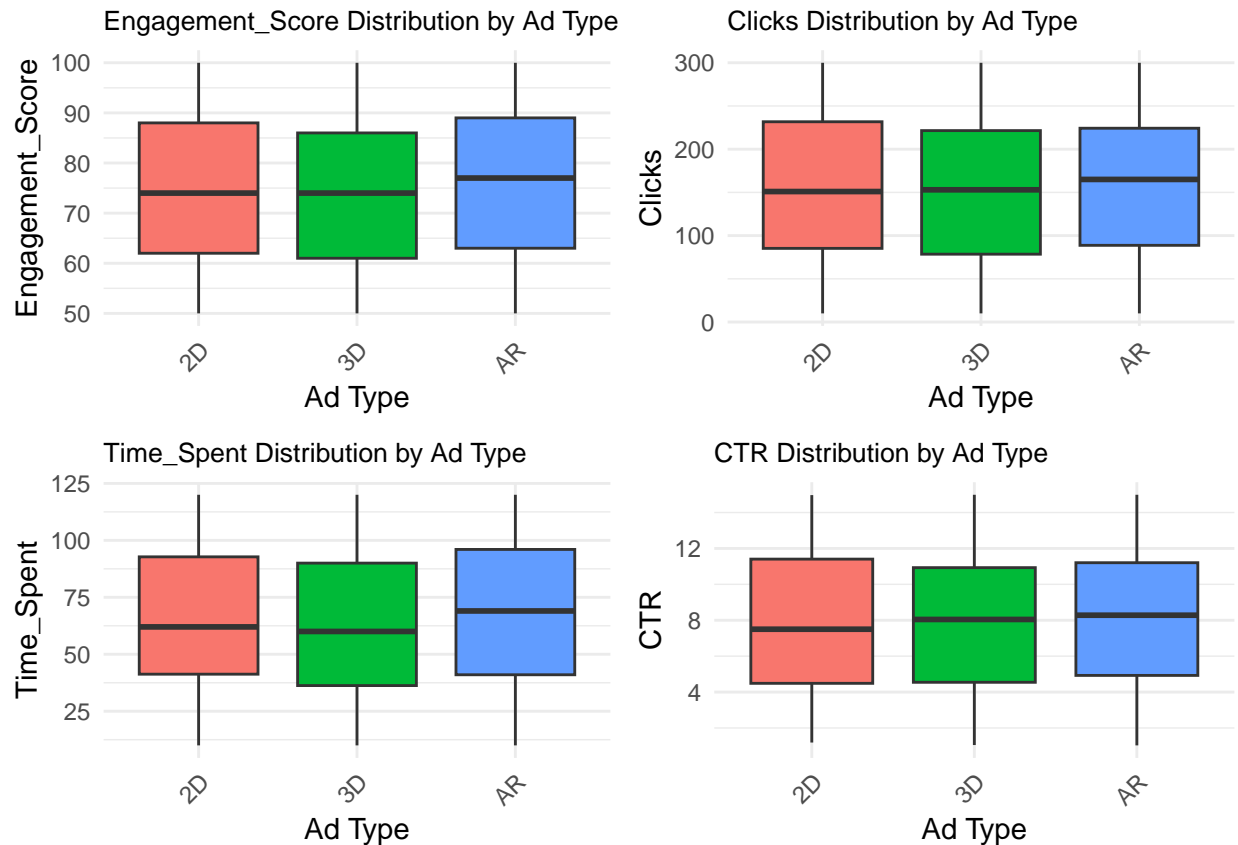
```

#function to create a boxplot for a given metric
create_boxplot <- function(data, metric_name) {
  ggplot(data, aes(x = Ad_Type, y = .data[[metric_name]])) +
    geom_boxplot(aes(fill = Ad_Type)) +
    theme_minimal() +
    labs(title = paste(metric_name, "Distribution by Ad Type"),
         x = "Ad Type",
         y = metric_name) +
    theme(
      axis.text.x = element_text(angle = 45, hjust = 1),
      plot.title = element_text(size = 10),
      legend.position = "none"
    )
}

#compare across all metrics
metrics <- c("Engagement_Score", "Clicks", "Time_Spent", "CTR")
plot_list <- lapply(metrics, function(metric) create_boxplot(data, metric))

#2x2 grid for plots
grid.arrange(
  plot_list[[1]], plot_list[[2]],
  plot_list[[3]], plot_list[[4]],
  ncol = 2
)

```



```
#save
ggsave("visualizations\\ad_type_metrics_comparison.png",
  arrangeGrob(
    plot_list[[1]], plot_list[[2]],
    plot_list[[3]], plot_list[[4]],
    ncol = 2
  ),
  width = 12, height = 10)
```

- Based on our results, we see that while Ad Type does not have any significant effect on our target metrics, we still do know that 3D ads and AR ads are more popular based on the barplot from earlier.

Interactions between Ad Type and Visual Complexity

```
# First, convert Visual_Complexity to factor with specified order
data$Visual_Complexity <- factor(data$Visual_Complexity,
  levels = c("Low", "Medium", "High"))

# Function to create interaction plot for a given metric
create_interaction_plot <- function(data, metric_name) {
  ggplot(data, aes(x = Visual_Complexity, y = .data[[metric_name]],
    color = Ad_Type, group = Ad_Type)) +
```

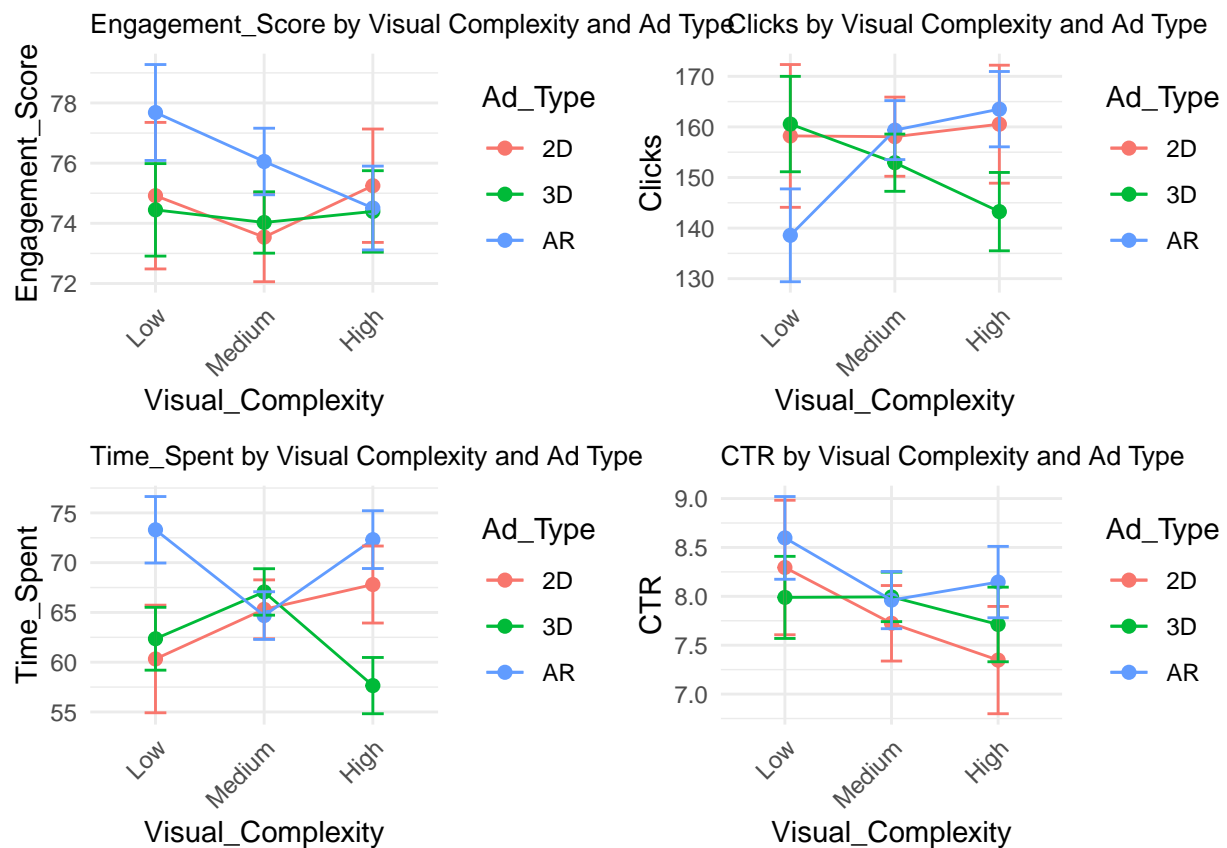
```

stat_summary(fun = mean, geom = "point", size = 2) +
stat_summary(fun = mean, geom = "line") +
stat_summary(fun.data = mean_se, geom = "errorbar", width = 0.2) +
theme_minimal() +
labs(title = paste(metric_name, "by Visual Complexity and Ad Type"),
     y = metric_name) +
theme(
  axis.text.x = element_text(angle = 45, hjust = 1),
  plot.title = element_text(size = 10)
)
}

# Create plots for all metrics
metrics <- c("Engagement_Score", "Clicks", "Time_Spent", "CTR")
interaction_plots <- lapply(metrics, function(metric) create_interaction_plot(data, metric))

# Arrange plots in a 2x2 grid
grid.arrange(
  interaction_plots[[1]], interaction_plots[[2]],
  interaction_plots[[3]], interaction_plots[[4]],
  ncol = 2
)

```



```

#save
ggsave("visualizations\\interaction_plots.png",
       arrangeGrob(
         interaction_plots[[1]], interaction_plots[[2]],
         interaction_plots[[3]], interaction_plots[[4]],
         ncol = 2
       ),
       width = 12, height = 10)

# Create empty list to store results
results <- list()

# Run ANOVA for each metric and store results
for (metric in metrics) {
  # Create formula and run ANOVA
  formula <- as.formula(paste(metric, "~ Ad_Type * Visual_Complexity"))
  model <- aov(formula, data = data)

  # Store results
  results[[metric]] <- summary(model)[[1]]
}

# Print results in a clear format
cat("Interaction Analysis Results:\n")

```

```
## Interaction Analysis Results:
```

```
cat("=====\n\n")
```

```
## =====
```

```

for (metric in metrics) {
  cat(sprintf("Metric: %s\n", metric))
  cat("-----\n")

  # Extract interaction p-value
  p_val <- results[[metric]]["Ad_Type:Visual_Complexity", "Pr(>F)"]
  f_val <- results[[metric]]["Ad_Type:Visual_Complexity", "F value"]

  cat(sprintf("F-value: %.3f\n", f_val))
  cat(sprintf("p-value: %.4f\n", p_val))
  cat(sprintf("Significant: %s\n\n", ifelse(p_val < 0.05, "Yes", "No")))
}

```

```

## Metric: Engagement_Score
## -----
## F-value: 0.473
## p-value: 0.7558
## Significant: No
##
## Metric: Clicks
## -----

```



```
## F-value: 1.663
## p-value: 0.1563
## Significant: No
##
## Metric: Time_Spent
## -----
## F-value: 3.536
## p-value: 0.0071
## Significant: Yes
##
## Metric: CTR
## -----
## F-value: 0.351
## p-value: 0.8434
## Significant: No
```

- This test tells us that how long users spend with an ad (Time Spent) depends on both the Ad Type AND Visual Complexity working together, the other metrics are not affected by Visual Complexity

Exploring the effects of Ad Type and Visual Complexity on Time Spent

```
# 1. Focused Time Spent Interaction Plot with Enhanced Details
time_spent_plot <- ggplot(data, aes(x = Visual_Complexity, y = Time_Spent,
                                     color = Ad_Type, group = Ad_Type)) +
  stat_summary(fun = mean, geom = "point") +
  stat_summary(fun = mean, geom = "line") +
  stat_summary(fun.data = mean_se, geom = "errorbar", width = 0.2) +
  theme_minimal() +
  labs(title = "Interaction Effect: Ad Type and Visual Complexity on Time Spent",
       subtitle = "Error bars represent standard error of the mean",
       y = "Time Spent (seconds)") +
  scale_color_brewer(palette = "Set2") +
  theme(
    legend.position = "right",
    plot.title = element_text(size = 12, face = "bold"),
    axis.title = element_text(size = 10),
    legend.title = element_text(size = 10)
  )

# 2. Post-hoc analysis
# Perform Tukey's HSD test
tukey_model <- aov(Time_Spent ~ Ad_Type * Visual_Complexity, data = data)
tukey_results <- TukeyHSD(tukey_model, which = "Ad_Type:Visual_Complexity")

# Filter significant comparisons (p < 0.05)
sig_comparisons <- as.data.frame(tukey_results$`Ad_Type:Visual_Complexity`)
sig_comparisons$comparison <- rownames(sig_comparisons)
sig_comparisons <- sig_comparisons[sig_comparisons$p_adj < 0.05, ]

# Print significant differences
cat("\nSignificant differences in Time Spent:\n")
```

```
##
## Significant differences in Time Spent:

cat("=====\n")

## =====

if(nrow(sig_comparisons) > 0) {
  for(i in 1:nrow(sig_comparisons)) {
    cat(sprintf("\n%s:\n", sig_comparisons$comparison[i]))
    cat(sprintf("Difference: %.2f seconds\n", sig_comparisons$diff[i]))
    cat(sprintf("Adjusted p-value: %.4f\n", sig_comparisons$p_adj[i]))
  }
} else {
  cat("No pairwise comparisons were significant at p < 0.05\n")
}

##
## 3D:High-AR:Low:
## Difference: -15.66 seconds
## Adjusted p-value: 0.0197
##
## AR:High-3D:High:
## Difference: 14.67 seconds
## Adjusted p-value: 0.0115

# Additional insight: Check effect sizes within each Visual Complexity level
cat("\nEffect Sizes within Visual Complexity Levels:\n")

##
## Effect Sizes within Visual Complexity Levels:

cat("=====\n")

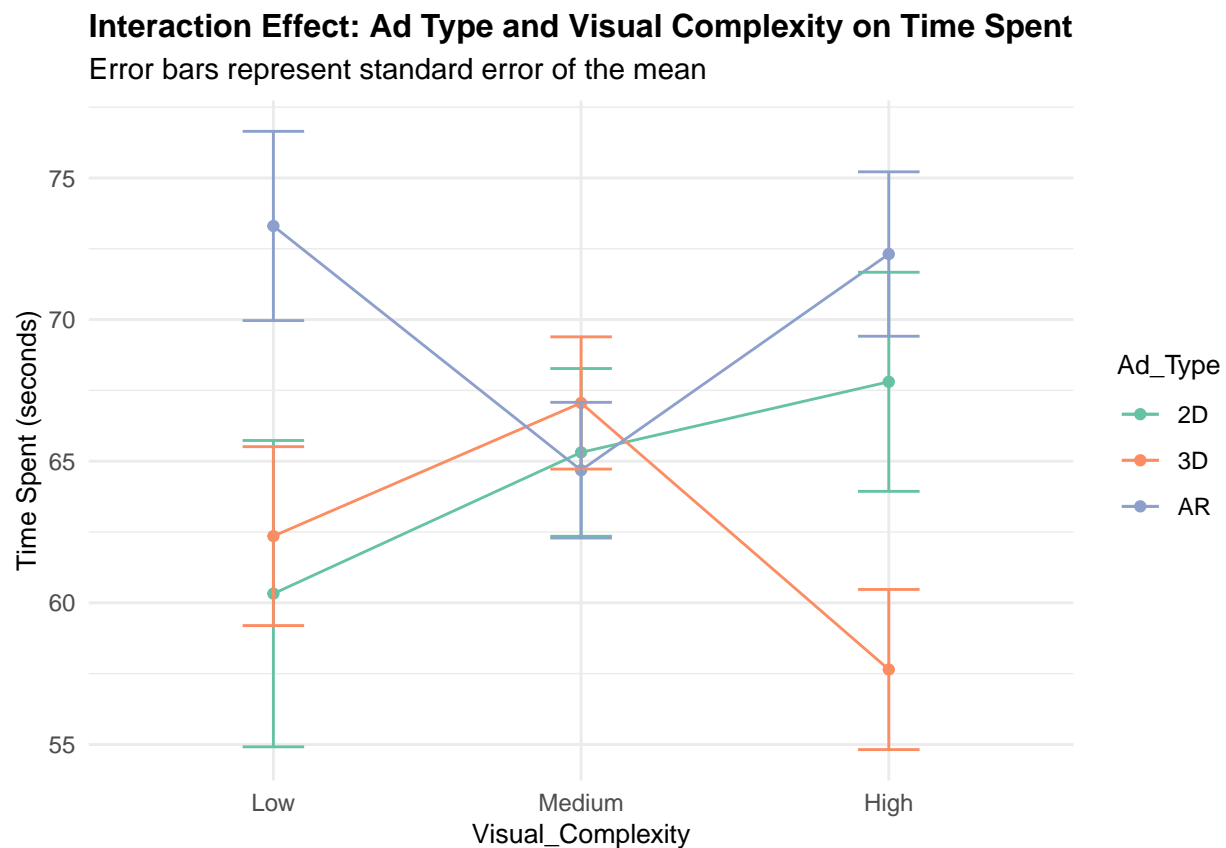
## =====

for(complexity in levels(data$Visual_Complexity)) {
  subset_data <- data[data$Visual_Complexity == complexity, ]
  effect_size <- summary(aov(Time_Spent ~ Ad_Type, data = subset_data))[[1]]
  cat(sprintf("\nVisual Complexity: %s\n", complexity))
  cat(sprintf("F-value: %.3f\n", effect_size$`F value`[1]))
  cat(sprintf("p-value: %.4f\n", effect_size$`Pr(>F)`[1]))
}

##
## Visual Complexity: Low
## F-value: 3.593
## p-value: 0.0293
##
## Visual Complexity: Medium
## F-value: 0.274
```

```
## p-value: 0.7603
##
## Visual Complexity: High
## F-value: 6.928
## p-value: 0.0012
```

```
# Display the focused plot
print(time_spent_plot)
```



```
#save
ggsave("visualizations\\time_spent_interaction_plot.png", time_spent_plot, width = 10, height = 6)
```

- Looking further we see that AR Ads are most effective at Low and High Complexities
- At Medium Complexity, each Ad Type performs similarly

Verifying from our Tukeys HSD test: > * Low Complexity: Moderate effect ($F = 3.593$, $p = 0.0293$) - Ad types do differ significantly > * Medium Complexity: No significant effect ($F = 0.274$, $p = 0.7603$) - Ad types perform similarly > * High Complexity: Strongest effect ($F = 6.928$, $p = 0.0012$) - Ad types show very significant differences

Relationship between Visual Complexity and User Movement

```

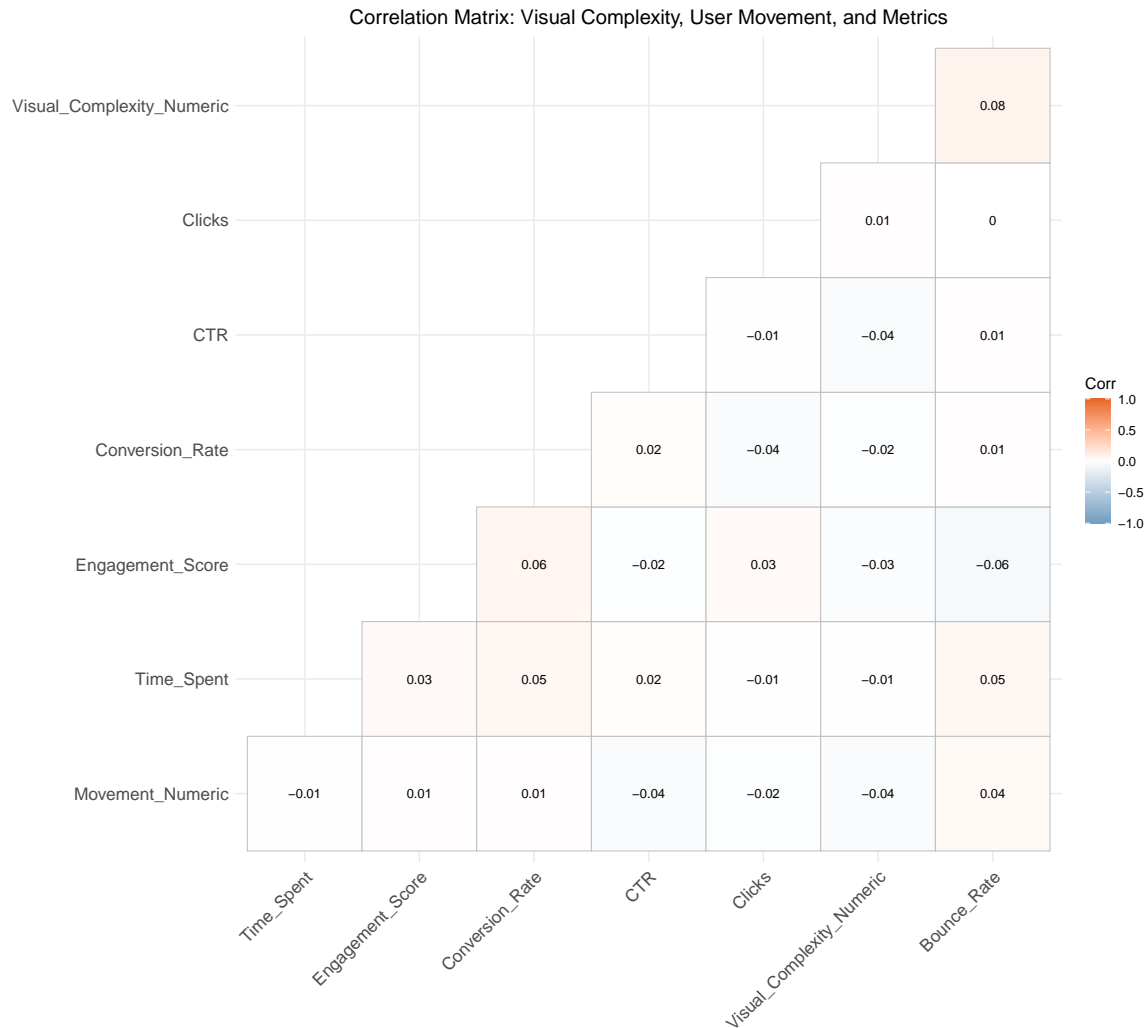
# Select relevant columns for correlation
correlation_cols <- c("Visual_Complexity_Numeric", "Movement_Numeric",
                      "Clicks", "Time_Spent", "Engagement_Score",
                      "Conversion_Rate", "Bounce_Rate", "CTR")

correlation_data <- data %>%
  dplyr::select(dplyr::all_of(correlation_cols))

# Calculate correlation matrix
cor_matrix <- cor(correlation_data)

# Create correlation plot
ggcorrplot(cor_matrix,
            hc.order = TRUE,
            type = "lower",
            lab = TRUE,
            lab_size = 3,
            colors = c("#6D9EC1", "white", "#E46726"),
            title = "Correlation Matrix: Visual Complexity, User Movement, and Metrics",
            ggtheme = theme_minimal()) +
  theme(
    plot.title = element_text(hjust = 0.5, size = 14),
    axis.text.x = element_text(angle = 45, hjust = 1),
    axis.text.y = element_text(hjust = 1)
  )

```



```
# Add statistical significance tests
correlation_tests <- data.frame(
  Variable1 = character(),
  Variable2 = character(),
  Correlation = numeric(),
  P_Value = numeric()
)

variables <- colnames(correlation_data)
for(i in 1:(length(variables)-1)) {
  for(j in (i+1):length(variables)) {
    test <- cor.test(correlation_data[[variables[i]]],
                     correlation_data[[variables[j]]])
    correlation_tests <- rbind(correlation_tests,
                              data.frame(Variable1 = variables[i],
                                         Variable2 = variables[j],
                                         Correlation = test$estimate,
                                         P_Value = test$p.value))
  }
}
```

```

# Display significant correlations (p < 0.05)
cat("\nStatistically Significant Correlations (p < 0.05):\n\n")

##
## Statistically Significant Correlations (p < 0.05):

significant_cors <- correlation_tests %>%
  filter(P_Value < 0.05) %>%
  arrange(P_Value) %>%
  mutate(
    Correlation = round(Correlation, 3),
    P_Value = round(P_Value, 4)
  )

# Print each correlation clearly
for(i in 1:nrow(significant_cors)) {
  cat(sprintf("%s and %s:\n",
              significant_cors$Variable1[i],
              significant_cors$Variable2[i]))
  cat(sprintf("  Correlation: %.3f\n", significant_cors$Correlation[i]))
  cat(sprintf("  P-value: %.4f\n\n", significant_cors$P_Value[i]))
}

## Visual_Complexity_Numeric and Bounce_Rate:
##   Correlation: 0.082
##   P-value: 0.0093
##
## Engagement_Score and Bounce_Rate:
##   Correlation: -0.065
##   P-value: 0.0402

```

Looking at the correlation plot, we don't see any strong correlations between Visual Complexity and User Movement or any of the other metrics

Visual Complexity and Bounce Rate > * very weak positive correlation > * it is statistically significant ($p < 0.01$), however the practical effect is unnoticeable

Engagement Score and Bounce Rate > * very weak negative correlation > * it is statistically significant ($p < 0.05$), however again, the practical effect is unnoticeable

Ad Characteristics Analysis

Now here's the interesting part, let's see which characteristics have the greatest impact on our metrics

Analysis of Ad Type Performance

```

# Create summary statistics for Ad Type
ad_type_summary <- data %>%
  group_by(Ad_Type) %>%

```

```

summarise(across(c(Clicks, Time_Spent, Engagement_Score,
                  Conversion_Rate, Bounce_Rate, CTR),
              list(
                mean = ~mean(.x, na.rm = TRUE),
                sd = ~sd(.x, na.rm = TRUE)
              )),
          n = n()) %>%
ungroup()

# Print summary statistics
cat("Summary Statistics by Ad Type:\n")

```

Summary Statistics by Ad Type:

```
print(ad_type_summary)
```

```

## # A tibble: 3 x 14
##   Ad_Type Clicks_mean Clicks_sd Time_Spent_mean Time_Spent_sd
##   <chr>      <dbl>    <dbl>         <dbl>         <dbl>
## 1 2D          159.      83.6           65.1           30.7
## 2 3D          152.      84.2           63.4           32.4
## 3 AR          156.      80.4           68.8           31.8
## # i 9 more variables: Engagement_Score_mean <dbl>, Engagement_Score_sd <dbl>,
## #   Conversion_Rate_mean <dbl>, Conversion_Rate_sd <dbl>,
## #   Bounce_Rate_mean <dbl>, Bounce_Rate_sd <dbl>, CTR_mean <dbl>, CTR_sd <dbl>,
## #   n <int>

```

```

# Function to create a single metric plot
plot_metric <- function(data, metric_name) {
  ggplot(data, aes(x = Ad_Type, y = !!sym(paste0(metric_name, "_mean")))) +
    geom_bar(stat = "identity", fill = "#4169E1", alpha = 0.7) +
    geom_errorbar(aes(ymin = !!sym(paste0(metric_name, "_mean")) - !!sym(paste0(metric_name, "_sd")),
                     ymax = !!sym(paste0(metric_name, "_mean")) + !!sym(paste0(metric_name, "_sd")),
                     width = 0.2) +
    theme_minimal() +
    labs(title = paste("Average", gsub("_", " ", metric_name), "by Ad Type"),
         x = "Ad Type",
         y = paste("Average", gsub("_", " ", metric_name))) +
    theme(axis.text.x = element_text(angle = 45, hjust = 1),
          plot.title = element_text(hjust = 0.5, size = 12))
}

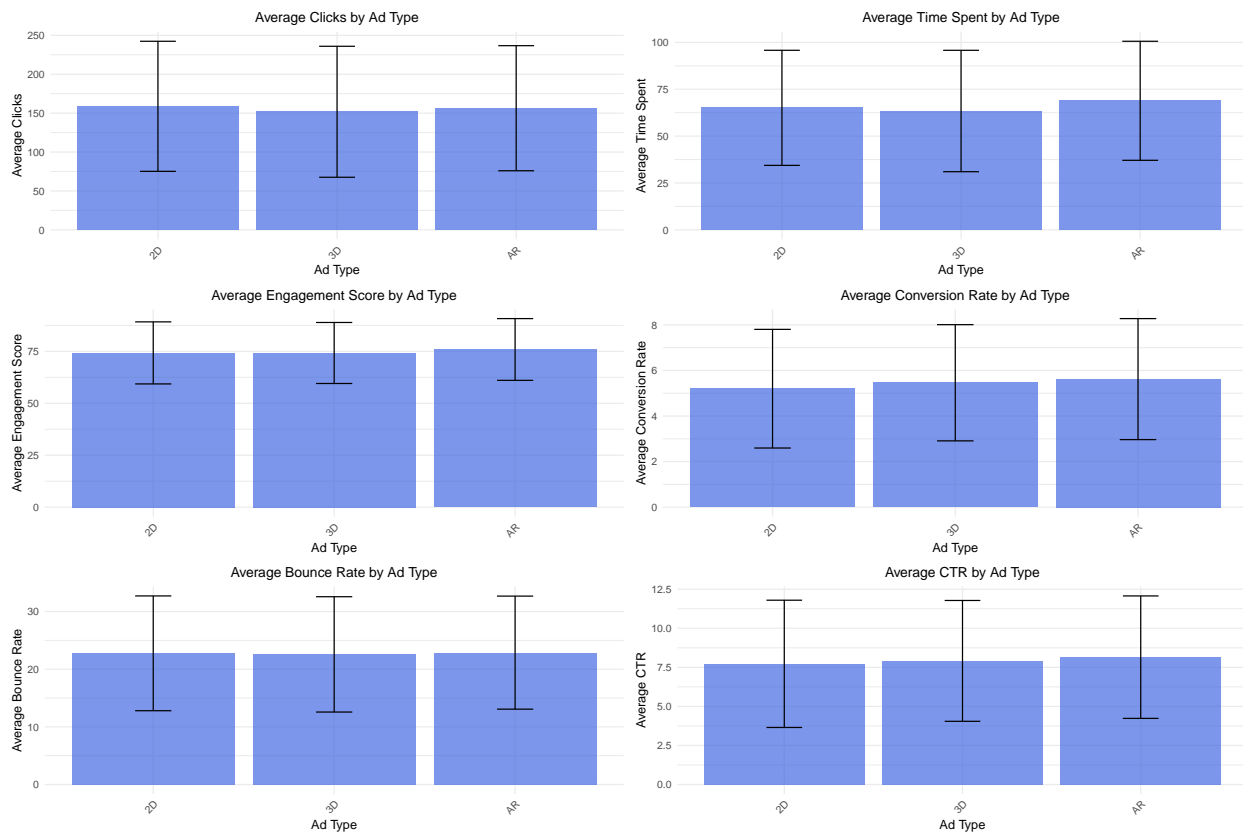
# Create plots for each metric
metrics <- c("Clicks", "Time_Spent", "Engagement_Score",
            "Conversion_Rate", "Bounce_Rate", "CTR")

plots <- lapply(metrics, function(metric) plot_metric(ad_type_summary, metric))

# Display plots in a grid with proper spacing
grid.arrange(grobs = plots,
             ncol = 2,
             widths = c(1, 1),

```

```
heights = c(1, 1, 1),
padding = unit(2, "line"))
```



```
# Perform ANOVA tests for each metric
cat("\nStatistical Analysis for Ad Type:\n")
```

```
##
## Statistical Analysis for Ad Type:
```

```
for (metric in metrics) {
  cat(paste("\n", gsub("_", " ", metric), "ANOVA Results:\n"))
  model <- aov(as.formula(paste(metric, "~ Ad_Type")), data = data)
  print(summary(model))

  # If ANOVA is significant, perform Tukey's test
  if (summary(model)[[1]]$"Pr(>F)"[1] < 0.05) {
    cat("\nTukey's HSD Test Results:\n")
    print(TukeyHSD(model))
  }
}
```

```
##
## Clicks ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type    2   7932   3966   0.581   0.56
```



```
## Residuals    997 6807515    6828
##
## Time Spent ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type     2   6033    3017   2.981 0.0512 .
## Residuals   997 1009009    1012
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Engagement Score ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type     2    663    331.7   1.514 0.221
## Residuals   997 218463    219.1
##
## Conversion Rate ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type     2     23   11.704   1.729 0.178
## Residuals   997   6747    6.767
##
## Bounce Rate ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type     2     18    9.20   0.093 0.911
## Residuals   997  98149   98.44
##
## CTR ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Ad_Type     2     26   12.99   0.841 0.431
## Residuals   997  15393   15.44
```

```
# Create a summary table of best performing ad types
best_performers <- data.frame(
  Metric = character(),
  Best_Ad_Type = character(),
  Mean_Value = numeric(),
  Significant = character()
)

for (metric in metrics) {
  # Get best performing ad type
  best_idx <- which.max(ad_type_summary[[paste0(metric, "_mean")]])

  # Check significance
  model <- aov(as.formula(paste(metric, "~ Ad_Type")), data = data)
  is_significant <- summary(model)[[1]]$"Pr(>F)"[1] < 0.05

  best_performers <- rbind(best_performers, data.frame(
    Metric = metric,
    Best_Ad_Type = ad_type_summary$Ad_Type[best_idx],
    Mean_Value = round(ad_type_summary[[paste0(metric, "_mean")]][best_idx], 2),
    Significant = ifelse(is_significant, "Yes", "No")
  ))
}

cat("\nBest Performing Ad Types for Each Metric:\n")
```

```
##
## Best Performing Ad Types for Each Metric:
```

```
print(best_performers)
```

```
##           Metric Best_Ad_Type Mean_Value Significant
## 1           Clicks           2D    158.78           No
## 2        Time_Spent           AR     68.84           No
## 3 Engagement_Score           AR     75.91           No
## 4 Conversion_Rate           AR      5.62           No
## 5      Bounce_Rate           AR     22.88           No
## 6              CTR           AR      8.15           No
```

Analysis of Age Group Performance

```
# Create summary statistics for Ad Type
age_group_summary <- data %>%
  group_by(Age_Group) %>%
  summarise(across(c(Clicks, Time_Spent, Engagement_Score,
                     Conversion_Rate, Bounce_Rate, CTR),
                  list(
                    mean = ~mean(.x, na.rm = TRUE),
                    sd = ~sd(.x, na.rm = TRUE)
                  )),
            n = n()) %>%
  ungroup()

# Print summary statistics
cat("Summary Statistics by Age Group:\n")
```

```
## Summary Statistics by Age Group:
```

```
print(age_group_summary)
```

```
## # A tibble: 5 x 14
##   Age_Group Clicks_mean Clicks_sd Time_Spent_mean Time_Spent_sd
##   <chr>      <dbl>      <dbl>          <dbl>          <dbl>
## 1 18-24      159.       83.5           65.1           31.3
## 2 25-34      154.       83.2           67.7           32.6
## 3 35-44      153.       82.6           65.6           31.2
## 4 45-54      139.       81.2           59.9           32.8
## 5 55+        163.       67.5           59.2           30.0
## # i 9 more variables: Engagement_Score_mean <dbl>, Engagement_Score_sd <dbl>,
## #   Conversion_Rate_mean <dbl>, Conversion_Rate_sd <dbl>,
## #   Bounce_Rate_mean <dbl>, Bounce_Rate_sd <dbl>, CTR_mean <dbl>, CTR_sd <dbl>,
## #   n <int>
```

```
# Function to create a single metric plot
plot_metric <- function(data, metric_name) {
  ggplot(data, aes(x = Age_Group, y = !!sym(paste0(metric_name, "_mean")))) +
```

```

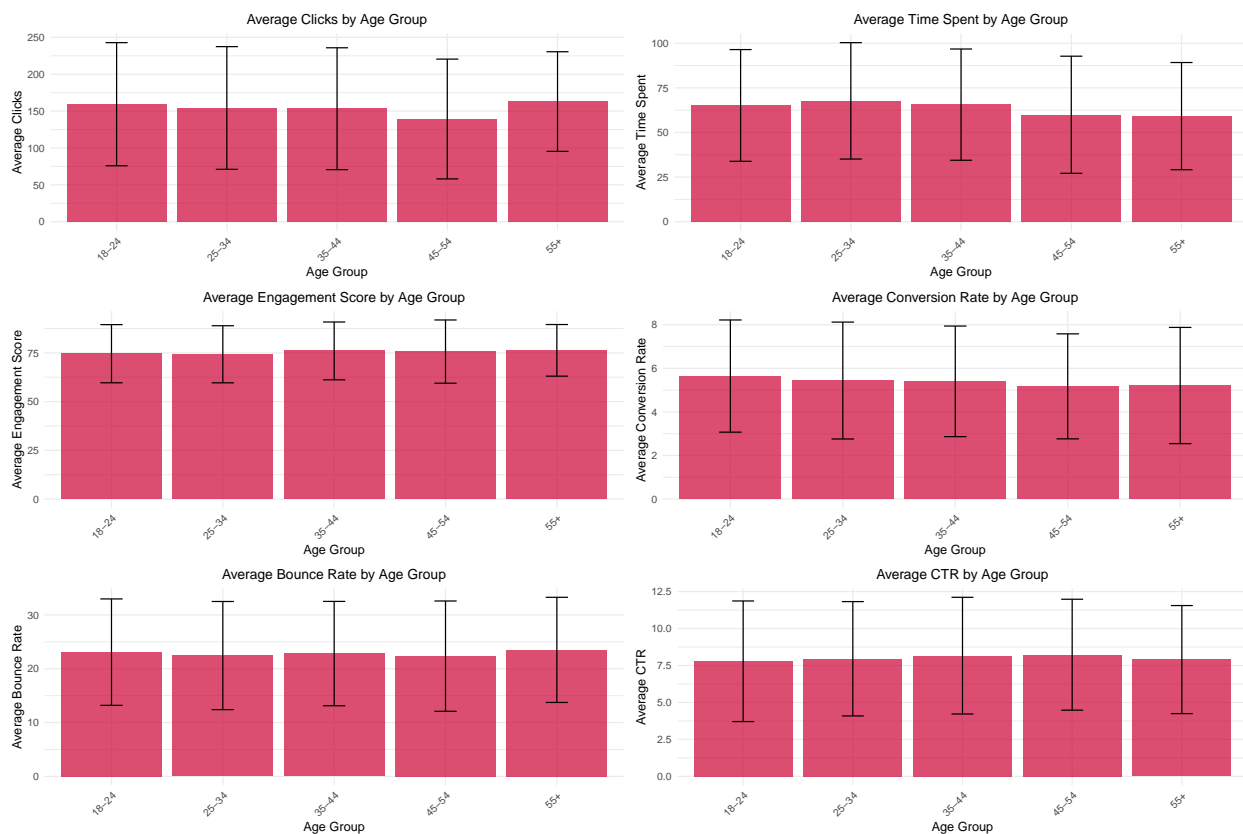
geom_bar(stat = "identity", fill = "#CC0033", alpha = 0.7) +
geom_errorbar(aes(ymin = !!sym(paste0(metric_name, "_mean")) - !!sym(paste0(metric_name, "_sd")),
  ymax = !!sym(paste0(metric_name, "_mean")) + !!sym(paste0(metric_name, "_sd")),
  width = 0.2) +
theme_minimal() +
labs(title = paste("Average", gsub("_", " ", metric_name), "by Age Group"),
  x = "Age Group",
  y = paste("Average", gsub("_", " ", metric_name))) +
theme(axis.text.x = element_text(angle = 45, hjust = 1),
  plot.title = element_text(hjust = 0.5, size = 12))
}

# Create plots for each metric
metrics <- c("Clicks", "Time_Spent", "Engagement_Score",
  "Conversion_Rate", "Bounce_Rate", "CTR")

plots <- lapply(metrics, function(metric) plot_metric(age_group_summary, metric))

# Display plots in a grid with proper spacing
grid.arrange(grobs = plots,
  ncol = 2,
  widths = c(1, 1),
  heights = c(1, 1, 1),
  padding = unit(2, "line"))

```



```
# Perform ANOVA tests for each metric
cat("\nStatistical Analysis for Age Group:\n")
```

```
##
```

```
## Statistical Analysis for Age Group:
```

```
for (metric in metrics) {
  cat(paste("\n", gsub("_", " ", metric), "ANOVA Results:\n"))
  model <- aov(as.formula(paste(metric, "~ Age_Group")), data = data)
  print(summary(model))

  # If ANOVA is significant, perform Tukey's test
  if (summary(model)[[1]]$"Pr(>F)"[1] < 0.05) {
    cat("\nTukey's HSD Test Results:\n")
    print(TukeyHSD(model))
  }
}
```

```
##
```

```
## Clicks ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4   22808     5702   0.835  0.503
## Residuals  995 6792638     6827
```

```
##
```

```
## Time Spent ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4   4994     1249   1.23  0.296
## Residuals  995 1010048     1015
```

```
##
```

```
## Engagement Score ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4    537    134.3   0.611  0.655
## Residuals  995 218590    219.7
```

```
##
```

```
## Conversion Rate ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4     17    4.368   0.644  0.631
## Residuals  995   6753    6.787
```

```
##
```

```
## Bounce Rate ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4     98    24.51   0.249  0.911
## Residuals  995  98069    98.56
```

```
##
```

```
## CTR ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Age_Group    4     22    5.541   0.358  0.838
## Residuals  995  15397   15.474
```

```
# Create a summary table of best performing ad types
```

```
best_performers <- data.frame(
  Metric = character(),
```

```

Best_Age_Group = character(),
Mean_Value = numeric(),
Significant = character()
)

for (metric in metrics) {
  # Get best performing ad type
  best_idx <- which.max(age_group_summary[[paste0(metric, "_mean")]])

  # Check significance
  model <- aov(as.formula(paste(metric, "~ Age_Group")), data = data)
  is_significant <- summary(model)[[1]]$"Pr(>F)"[1] < 0.05

  best_performers <- rbind(best_performers, data.frame(
    Metric = metric,
    Best_Age_Group = age_group_summary$Age_Group[best_idx],
    Mean_Value = round(age_group_summary[[paste0(metric, "_mean")]][best_idx], 2),
    Significant = ifelse(is_significant, "Yes", "No")
  ))
}

cat("\nMost Impacted Age Groups for Each Metric:\n")

```

```

##
## Most Impacted Age Groups for Each Metric:

```

```
print(best_performers)
```

```

##           Metric Best_Age_Group Mean_Value Significant
## 1           Clicks           55+      162.97          No
## 2        Time_Spent        25-34       67.72          No
## 3 Engagement_Score           55+       76.30          No
## 4  Conversion_Rate        18-24        5.64          No
## 5      Bounce_Rate           55+       23.50          No
## 6              CTR        45-54        8.23          No

```

Analysis of Device Type Performance

```

# Create summary statistics for Device Type
device_type_summary <- data %>%
  group_by(Device_Type) %>%
  summarise(across(c(Clicks, Time_Spent, Engagement_Score,
                     Conversion_Rate, Bounce_Rate, CTR),
    list(
      mean = ~mean(.x, na.rm = TRUE),
      sd = ~sd(.x, na.rm = TRUE)
    )),
    n = n()) %>%
  ungroup()

```

```

# Print summary statistics
cat("Summary Statistics by Device Type:\n")

## Summary Statistics by Device Type:

print(device_type_summary)

## # A tibble: 3 x 14
##   Device_Type Clicks_mean Clicks_sd Time_Spent_mean Time_Spent_sd
##   <chr>         <dbl>     <dbl>         <dbl>         <dbl>
## 1 Desktop      158.       79.5          64.8          32.3
## 2 Mobile       153.       83.6          65.7          31.5
## 3 Tablet       154.       86.2          69.7          32.8
## # i 9 more variables: Engagement_Score_mean <dbl>, Engagement_Score_sd <dbl>,
## #   Conversion_Rate_mean <dbl>, Conversion_Rate_sd <dbl>,
## #   Bounce_Rate_mean <dbl>, Bounce_Rate_sd <dbl>, CTR_mean <dbl>, CTR_sd <dbl>,
## #   n <int>

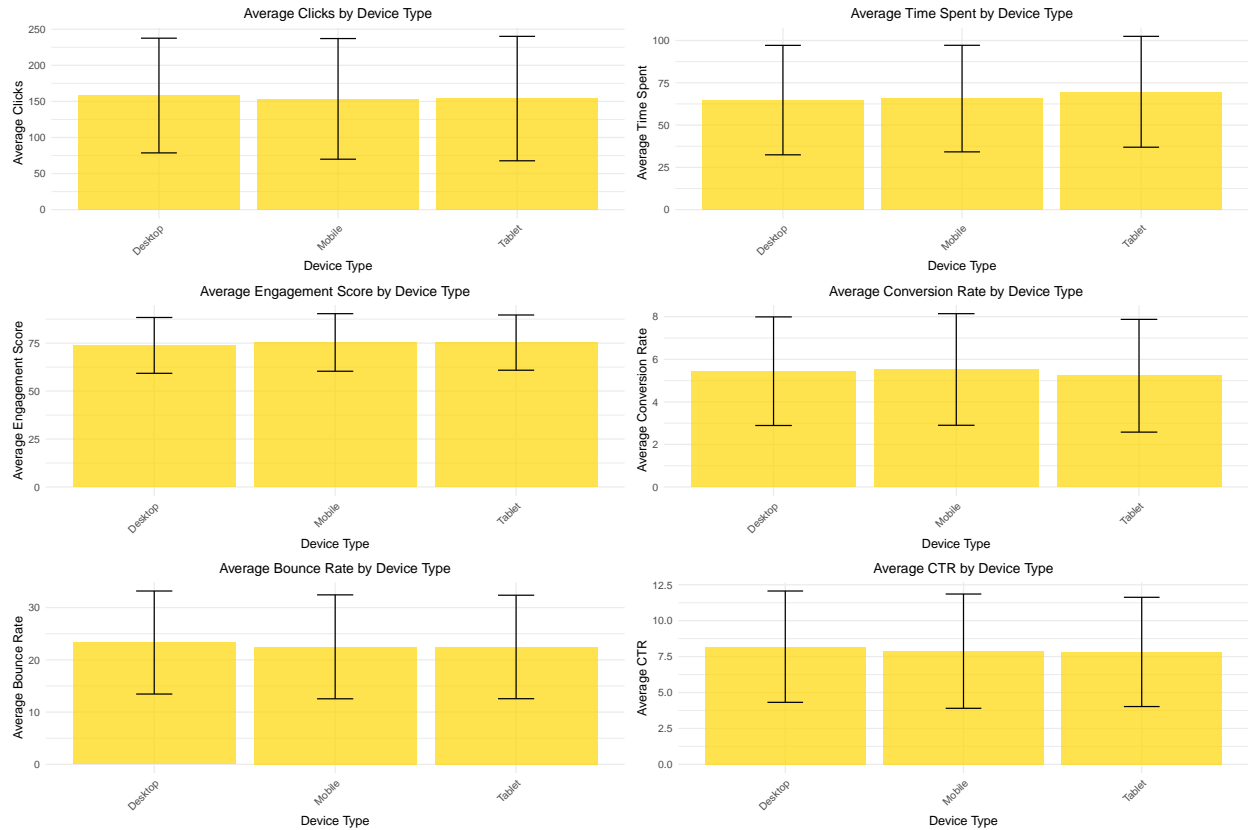
# Function to create a single metric plot
plot_metric <- function(data, metric_name) {
  ggplot(data, aes(x = Device_Type, y = !!sym(paste0(metric_name, "_mean")))) +
    geom_bar(stat = "identity", fill = "#FFD700", alpha = 0.7) + # Changed to yellow
    geom_errorbar(aes(ymin = !!sym(paste0(metric_name, "_mean")) - !!sym(paste0(metric_name, "_sd")),
                      ymax = !!sym(paste0(metric_name, "_mean")) + !!sym(paste0(metric_name, "_sd")),
                      width = 0.2) +
    theme_minimal() +
    labs(title = paste("Average", gsub("_", " ", metric_name), "by Device Type"),
         x = "Device Type",
         y = paste("Average", gsub("_", " ", metric_name))) +
    theme(axis.text.x = element_text(angle = 45, hjust = 1),
          plot.title = element_text(hjust = 0.5, size = 12))
}

# Create plots for each metric
metrics <- c("Clicks", "Time_Spent", "Engagement_Score",
            "Conversion_Rate", "Bounce_Rate", "CTR")

plots <- lapply(metrics, function(metric) plot_metric(device_type_summary, metric))

# Display plots in a grid with proper spacing
grid.arrange(grobs = plots,
             ncol = 2,
             widths = c(1, 1),
             heights = c(1, 1, 1),
             padding = unit(2, "line"))

```



```
# Perform ANOVA tests for each metric
cat("\nStatistical Analysis for Device Type:\n")
```

```
##
## Statistical Analysis for Device Type:
```

```
for (metric in metrics) {
  cat(paste("\n", gsub("_", " ", metric), "ANOVA Results:\n"))
  model <- aov(as.formula(paste(metric, "~ Device_Type")), data = data)
  print(summary(model))

  # If ANOVA is significant, perform Tukey's test
  if (summary(model)[[1]]$"Pr(>F)"[1] < 0.05) {
    cat("\nTukey's HSD Test Results:\n")
    print(TukeyHSD(model))
  }
}
```

```
##
## Clicks ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Device_Type  2    4362    2181   0.319  0.727
## Residuals  997 6811085    6832
##
## Time Spent ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
```

```
## Device_Type 2 1878 938.9 0.924 0.397
## Residuals 997 1013164 1016.2
##
## Engagement Score ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## Device_Type 2 481 240.4 1.096 0.334
## Residuals 997 218646 219.3
##
## Conversion Rate ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## Device_Type 2 8 3.957 0.583 0.558
## Residuals 997 6763 6.783
##
## Bounce Rate ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## Device_Type 2 143 71.30 0.725 0.484
## Residuals 997 98025 98.32
##
## CTR ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## Device_Type 2 22 10.85 0.703 0.495
## Residuals 997 15397 15.44
```

```
# Create a summary table of best performing device types
best_performers <- data.frame(
  Metric = character(),
  Best_Device_Type = character(),
  Mean_Value = numeric(),
  Significant = character()
)

for (metric in metrics) {
  # Get best performing device type
  best_idx <- which.max(device_type_summary[[paste0(metric, "_mean")]])

  # Check significance
  model <- aov(as.formula(paste(metric, "~ Device_Type")), data = data)
  is_significant <- summary(model)[[1]]$"Pr(>F)"[1] < 0.05

  best_performers <- rbind(best_performers, data.frame(
    Metric = metric,
    Best_Device_Type = device_type_summary$Device_Type[best_idx],
    Mean_Value = round(device_type_summary[[paste0(metric, "_mean")]][best_idx], 2),
    Significant = ifelse(is_significant, "Yes", "No")
  ))
}

cat("\nBest Performing Device Types for Each Metric:\n")
```

```
##
## Best Performing Device Types for Each Metric:
```



```
print(best_performers)
```

```
##           Metric Best_Device_Type Mean_Value Significant
## 1      Clicks      Desktop      158.13          No
## 2    Time_Spent      Tablet      69.68          No
## 3 Engagement_Score      Mobile      75.33          No
## 4 Conversion_Rate      Mobile      5.52          No
## 5    Bounce_Rate      Desktop      23.32          No
## 6          CTR      Desktop      8.19          No
```

Analysis of Visual Complexity Performance

```
# Create summary statistics for Visual Complexity
visual_complexity_summary <- data %>%
  group_by(Visual_Complexity) %>%
  summarise(across(c(Clicks, Time_Spent, Engagement_Score,
                    Conversion_Rate, Bounce_Rate, CTR),
    list(
      mean = ~mean(.x, na.rm = TRUE),
      sd = ~sd(.x, na.rm = TRUE)
    ),
    n = n()) %>%
  ungroup()

# Print summary statistics
cat("Summary Statistics by Visual Complexity:\n")
```

```
## Summary Statistics by Visual Complexity:
```

```
print(visual_complexity_summary)
```

```
## # A tibble: 3 x 14
##   Visual_Complexity Clicks_mean Clicks_sd Time_Spent_mean Time_Spent_sd
##   <fct>             <dbl>     <dbl>         <dbl>         <dbl>
## 1 Low              152.      85.4           66.2           30.6
## 2 Medium           156.      80.8           65.8           32.7
## 3 High            155.      83.9           65.5           31.4
## # i 9 more variables: Engagement_Score_mean <dbl>, Engagement_Score_sd <dbl>,
## #   Conversion_Rate_mean <dbl>, Conversion_Rate_sd <dbl>,
## #   Bounce_Rate_mean <dbl>, Bounce_Rate_sd <dbl>, CTR_mean <dbl>, CTR_sd <dbl>,
## #   n <int>
```

```
# Function to create a single metric plot
plot_metric <- function(data, metric_name) {
  ggplot(data, aes(x = Visual_Complexity, y = !!sym(paste0(metric_name, "_mean")))) +
    geom_bar(stat = "identity", fill = "#228B22", alpha = 0.7) + # Changed to forest green
    geom_errorbar(aes(ymin = !!sym(paste0(metric_name, "_mean")) - !!sym(paste0(metric_name, "_sd")),
                     ymax = !!sym(paste0(metric_name, "_mean")) + !!sym(paste0(metric_name, "_sd")),
                     width = 0.2) +
    theme_minimal() +
```

```

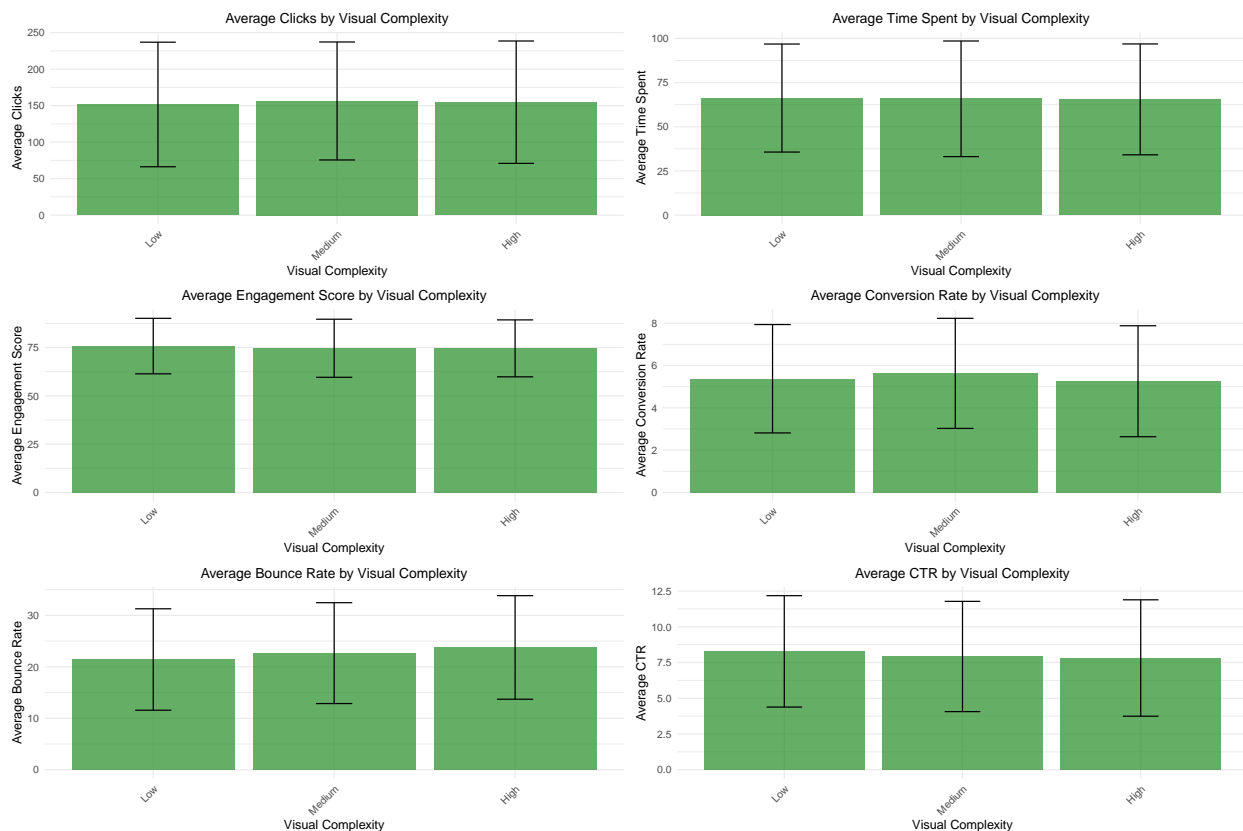
labs(title = paste("Average", gsub("_", " ", metric_name), "by Visual Complexity"),
     x = "Visual Complexity",
     y = paste("Average", gsub("_", " ", metric_name))) +
theme(axis.text.x = element_text(angle = 45, hjust = 1),
      plot.title = element_text(hjust = 0.5, size = 12))
}

# Create plots for each metric
metrics <- c("Clicks", "Time_Spent", "Engagement_Score",
            "Conversion_Rate", "Bounce_Rate", "CTR")

plots <- lapply(metrics, function(metric) plot_metric(visual_complexity_summary, metric))

# Display plots in a grid with proper spacing
grid.arrange(grobs = plots,
             ncol = 2,
             widths = c(1, 1),
             heights = c(1, 1, 1),
             padding = unit(2, "line"))

```



```

# Perform ANOVA tests for each metric
cat("\nStatistical Analysis for Visual Complexity:\n")

```

```

##
## Statistical Analysis for Visual Complexity:

```

```

for (metric in metrics) {
  cat(paste("\n", gsub("_", " ", metric), "ANOVA Results:\n"))
  model <- aov(as.formula(paste(metric, "~ Visual_Complexity")), data = data)
  print(summary(model))

  # If ANOVA is significant, perform Tukey's test
  if (summary(model)[[1]]$"Pr(>F)"[1] < 0.05) {
    cat("\nTukey's HSD Test Results:\n")
    print(TukeyHSD(model))
  }
}

##
## Clicks ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2     3349    1674   0.245  0.783
## Residuals       997 6812098    6833
##
## Time Spent ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2        68    33.9   0.033  0.967
## Residuals       997 1014974   1018.0
##
## Engagement Score ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2     216    108.1   0.492  0.611
## Residuals       997 218911    219.6
##
## Conversion Rate ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2      28   13.893   2.054  0.129
## Residuals       997   6743    6.763
##
## Bounce Rate ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2     664   331.9   3.394  0.034 *
## Residuals       997  97503    97.8
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Tukey's HSD Test Results:
##   Tukey multiple comparisons of means
##     95% family-wise confidence level
##
## Fit: aov(formula = as.formula(paste(metric, "~ Visual_Complexity")), data = data)
##
## $Visual_Complexity
##           diff          lwr          upr          p adj
## Medium-Low  1.236859 -0.6927145  3.166432  0.2891785
## High-Low    2.343086  0.2233881  4.462783  0.0259893
## High-Medium 1.106227 -0.5990504  2.811504  0.2806779
##
##

```

```
## CTR ANOVA Results:
##           Df Sum Sq Mean Sq F value Pr(>F)
## Visual_Complexity  2      28   13.80   0.894  0.409
## Residuals        997  15391   15.44

# Create a summary table of best performing visual complexity levels
best_performers <- data.frame(
  Metric = character(),
  Best_Visual_Complexity = character(),
  Mean_Value = numeric(),
  Significant = character()
)

for (metric in metrics) {
  # Get best performing visual complexity level
  best_idx <- which.max(visual_complexity_summary[[paste0(metric, "_mean")]])

  # Check significance
  model <- aov(as.formula(paste(metric, "~ Visual_Complexity")), data = data)
  is_significant <- summary(model)[[1]]$"Pr(>F)"[1] < 0.05

  best_performers <- rbind(best_performers, data.frame(
    Metric = metric,
    Best_Visual_Complexity = visual_complexity_summary$Visual_Complexity[best_idx],
    Mean_Value = round(visual_complexity_summary[[paste0(metric, "_mean")]][best_idx], 2),
    Significant = ifelse(is_significant, "Yes", "No")
  ))
}

cat("\nBest Performing Visual Complexity Levels for Each Metric:\n")
```

```
##
## Best Performing Visual Complexity Levels for Each Metric:
```

```
print(best_performers)
```

```
##           Metric Best_Visual_Complexity Mean_Value Significant
## 1           Clicks           Medium      156.38           No
## 2        Time_Spent           Low       66.25           No
## 3 Engagement_Score           Low       75.79           No
## 4   Conversion_Rate           Medium      5.63           No
## 5      Bounce_Rate           High      23.77           Yes
## 6             CTR           Low       8.28           No
```

Analysis of User Movement Performance

```
# Create summary statistics for User Movement
user_movement_summary <- data %>%
  group_by(User_Movement_Data) %>%
  summarise(across(c(Clicks, Time_Spent, Engagement_Score,
                     Conversion_Rate, Bounce_Rate, CTR),
```

```

        list(
          mean = ~mean(.x, na.rm = TRUE),
          sd = ~sd(.x, na.rm = TRUE)
        ),
        n = n()) %>%
  ungroup()

# Print summary statistics
cat("Summary Statistics by User Movement:\n")

```

```
## Summary Statistics by User Movement:
```

```
print(user_movement_summary)
```

```
## # A tibble: 3 x 14
##   User_Movement_Data Clicks_mean Clicks_sd Time_Spent_mean Time_Spent_sd
##   <chr>              <dbl>      <dbl>          <dbl>          <dbl>
## 1 gaze, movement      151.      83.5            64.9            32.0
## 2 movement only      159.      82.1            66.7            32.2
## 3 no movement        153.      81.9            65.8            31.1
## # i 9 more variables: Engagement_Score_mean <dbl>, Engagement_Score_sd <dbl>,
## #   Conversion_Rate_mean <dbl>, Conversion_Rate_sd <dbl>,
## #   Bounce_Rate_mean <dbl>, Bounce_Rate_sd <dbl>, CTR_mean <dbl>, CTR_sd <dbl>,
## #   n <int>
```

```

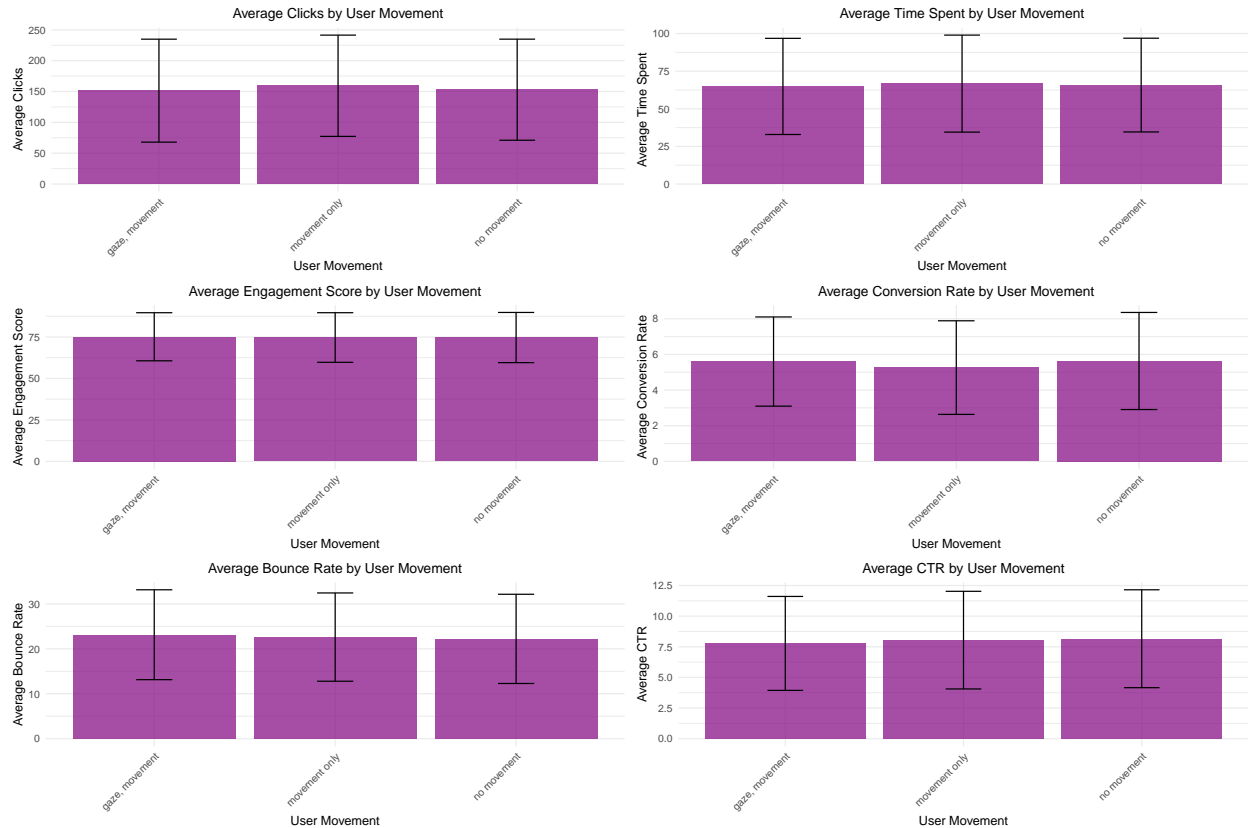
# Function to create a single metric plot
plot_metric <- function(data, metric_name) {
  ggplot(data, aes(x = User_Movement_Data, y = !!sym(paste0(metric_name, "_mean")))) +
    geom_bar(stat = "identity", fill = "#800080", alpha = 0.7) + # Changed to purple
    geom_errorbar(aes(ymin = !!sym(paste0(metric_name, "_mean")) - !!sym(paste0(metric_name, "_sd")),
                      ymax = !!sym(paste0(metric_name, "_mean")) + !!sym(paste0(metric_name, "_sd")),
                      width = 0.2) +
    theme_minimal() +
    labs(title = paste("Average", gsub("_", " ", metric_name), "by User Movement"),
         x = "User Movement",
         y = paste("Average", gsub("_", " ", metric_name))) +
    theme(axis.text.x = element_text(angle = 45, hjust = 1),
          plot.title = element_text(hjust = 0.5, size = 12))
}

# Create plots for each metric
metrics <- c("Clicks", "Time_Spent", "Engagement_Score",
            "Conversion_Rate", "Bounce_Rate", "CTR")

plots <- lapply(metrics, function(metric) plot_metric(user_movement_summary, metric))

# Display plots in a grid with proper spacing
grid.arrange(grobs = plots,
             ncol = 2,
             widths = c(1, 1),
             heights = c(1, 1, 1),
             padding = unit(2, "line"))

```



```
# Perform ANOVA tests for each metric
cat("\nStatistical Analysis for User Movement:\n")
```

```
##
```

```
## Statistical Analysis for User Movement:
```

```
for (metric in metrics) {
  cat(paste("\n", gsub("_", " ", metric), "ANOVA Results:\n"))
  model <- aov(as.formula(paste(metric, "~ User_Movement_Data")), data = data)
  print(summary(model))

  # If ANOVA is significant, perform Tukey's test
  if (summary(model)[[1]]$"Pr(>F)"[1] < 0.05) {
    cat("\nTukey's HSD Test Results:\n")
    print(TukeyHSD(model))
  }
}
```

```
##
```

```
## Clicks ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## User_Movement_Data  2   13464    6732  0.987  0.373
## Residuals       997 6801982    6822
```

```
##
```

```
## Time Spent ANOVA Results:
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
```

```
## User_Movement_Data 2 684 342.2 0.336 0.714
## Residuals 997 1014358 1017.4
##
## Engagement Score ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## User_Movement_Data 2 49 24.58 0.112 0.894
## Residuals 997 219078 219.74
##
## Conversion Rate ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## User_Movement_Data 2 30 14.767 2.184 0.113
## Residuals 997 6741 6.761
##
## Bounce Rate ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## User_Movement_Data 2 127 63.32 0.644 0.525
## Residuals 997 98041 98.34
##
## CTR ANOVA Results:
## Df Sum Sq Mean Sq F value Pr(>F)
## User_Movement_Data 2 24 12.05 0.78 0.459
## Residuals 997 15395 15.44
```

```
# Create a summary table of best performing user movement types
best_performers <- data.frame(
  Metric = character(),
  Best_User_Movement = character(),
  Mean_Value = numeric(),
  Significant = character()
)

for (metric in metrics) {
  # Get best performing user movement type
  best_idx <- which.max(user_movement_summary[[paste0(metric, "_mean")]])

  # Check significance
  model <- aov(as.formula(paste(metric, "~ User_Movement_Data")), data = data)
  is_significant <- summary(model)[[1]]$"Pr(>F)"[1] < 0.05

  best_performers <- rbind(best_performers, data.frame(
    Metric = metric,
    Best_User_Movement = user_movement_summary$User_Movement_Data[best_idx],
    Mean_Value = round(user_movement_summary[[paste0(metric, "_mean")]][best_idx], 2),
    Significant = ifelse(is_significant, "Yes", "No")
  ))
}

cat("\nBest Performing User Movement Types for Each Metric:\n")
```

```
##
## Best Performing User Movement Types for Each Metric:
```

```
print(best_performers)
```

##	Metric	Best_User_Movement	Mean_Value	Significant
## 1	Clicks	movement only	159.33	No
## 2	Time_Spent	movement only	66.73	No
## 3	Engagement_Score	gaze, movement	75.15	No
## 4	Conversion_Rate	no movement	5.63	No
## 5	Bounce_Rate	gaze, movement	23.14	No
## 6	CTR	no movement	8.16	No