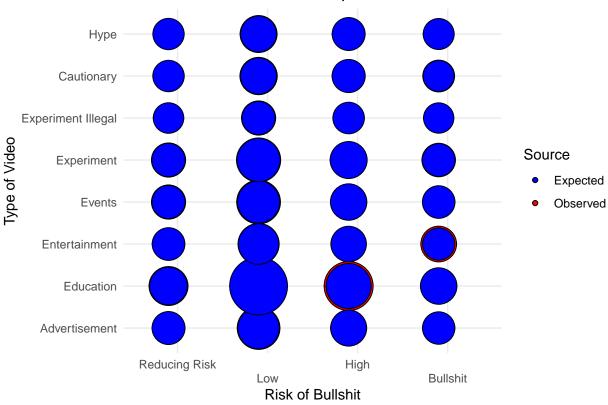
Fisher's MD

```
data <- readr::read_csv("Final_Main_Coder.csv")</pre>
## Rows: 394 Columns: 3-- Column specification ----
## Delimiter: ","
## chr (3): Video ID, Type of Video, Risk of Bullshit
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
contingency_table <- table(data$`Type of Video`, data$`Risk of Bullshit`)</pre>
fisher_result <- fisher.test(contingency_table, simulate.p.value = TRUE)</pre>
p value <- fisher result$p.value
cat("p-value:", p_value, "\n")
## p-value: 0.0004997501
# Extract observed values
observed_values <- contingency_table</pre>
print(observed_values)
##
##
                        Bullshit High Low Reducing Risk
##
     Advertisement
                               0
                                    7 31
##
     Cautionary
                                    0 11
                                                       0
                               1
##
     Education
                               6
                                   67 117
                                                      16
    Entertainment
##
                               7
                                    1 24
                                                       Λ
##
    Events
                                    2 36
                                                       4
##
     Experiment
                               3
                                    4 38
                                                       4
##
     Experiment Illegal
                               0
                                    0
                                       4
                                                       0
     Нуре
                               0
                                    1 10
##
# Perform chi-square test to calculate expected values
chi_result <- chisq.test(contingency_table)</pre>
## Warning in chisq.test(contingency_table): Chi-squared approximation may be
## incorrect
expected_values <- chi_result$expected</pre>
print(expected_values)
##
##
                         Bullshit
                                         High
                                                     Low Reducing Risk
##
     Advertisement
                        1.6395939 7.9086294 26.137056
                                                             2.3147208
     Cautionary
                        0.5177665 2.4974619
                                                             0.7309645
##
                                                8.253807
##
     Education
                        8.8883249 42.8730964 141.690355
                                                            12.5482234
##
    Entertainment
                        1.3807107 6.6598985 22.010152
                                                            1.9492386
##
    Events
                        1.8121827 8.7411168 28.888325
                                                             2.5583756
##
                        2.1142132 10.1979695 33.703046
     Experiment
                                                             2.9847716
     Experiment Illegal 0.1725888 0.8324873
##
                                                2.751269
                                                             0.2436548
##
                                              7.565990
                                                             0.6700508
     Нуре
                        0.4746193 2.2893401
```

```
# Convert observed_values to a data frame
observed_data <- as.data.frame.table(observed_values)</pre>
# Perform chi-square test to calculate expected values
chi result <- chisq.test(observed values)</pre>
## Warning in chisq.test(observed_values): Chi-squared approximation may be
## incorrect
expected_values <- chi_result$expected</pre>
# Convert expected_values to a data frame
expected_data <- as.data.frame.table(expected_values)</pre>
# Rename the columns
names(observed_data) <- c("Type_of_Video", "Risk_of_Bullshit", "Observed")</pre>
names(expected_data) <- c("Type_of_Video", "Risk_of_Bullshit", "Expected")</pre>
# Reorder the factor levels
observed_data$Type_of_Video <- factor(observed_data$Type_of_Video, levels = c("Advertisement", "Educati
observed_data$Risk_of_Bullshit <- factor(observed_data$Risk_of_Bullshit, levels = c("Reducing Risk", "L
expected_data$Type_of_Video <- factor(expected_data$Type_of_Video, levels = c("Advertisement", "Educati
expected_data$Risk_of_Bullshit <- factor(expected_data$Risk_of_Bullshit, levels = c("Reducing Risk", "L
library(ggplot2)
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
# Combine observed and expected data
combined_data <- bind_rows(</pre>
  observed_data %>% mutate(Source = "Observed"),
  expected_data %>% mutate(Source = "Expected")
)
# Reorder levels for better visualization
combined_data$Type_of_Video <- factor(</pre>
  combined_data$Type_of_Video,
  levels = c(
    "Advertisement", "Education", "Entertainment", "Events",
    "Experiment", "Experiment Illegal", "Cautionary", "Hype"
combined_data$Risk_of_Bullshit <- factor(</pre>
  combined_data$Risk_of_Bullshit,
  levels = c("Reducing Risk", "Low", "High", "Bullshit")
```

Warning: The `<scale>` argument of `guides()` cannot be `FALSE`. Use "none"
instead as of ggplot2 3.3.4.

Combined Observed and Expected Values Balloon Plot



```
expand = expansion(mult = c(0.05, 0.05))
) +
guides(size = FALSE, fill = guide_legend(title = "Source")) +
scale_fill_manual(values = c("blue", "red")) +
coord_cartesian(clip = "off")
```

Combined Observed and Expected Values Balloon Plot

