

Heatmaps Phi Bounds and Cole's C vs. Yule's Q vs. Phi

Function: Compare coefficient1 with coefficient2 by using a heatmap

- coefficient1: Coefficient that gets evaluated, i.e. whose values get displayed in the heatmap
- coefficient2.inv: Coefficient that gets fixed and used to create contingency tables for varying marginal distributions
- level: Level at which coefficient2 gets fixed
- xlim, ylim: range of PA and PB as a vector. Recommendation: use c(0.01, 0.99) but don't include 0 or 1. This will produce errors.
- xlab, ylab: axis description with strings
- PA: Marginal probability for event A (row 1 in contingency table)
- PB: Marginal probability for event B (column 1 in contingency table)

```
plot_contour_binary <- function(coefficient1, coefficient2.inv,
  level, xlim, ylim, grid, xlab, ylab, bins) {
  # function yielding the value for coefficient1 given
  # the contingency table which is determined by
  # coefficient2.inv, PA and PB
  f <- function(coefficient1, coefficient2.inv, PA, PB) {
    as.numeric(coefficient1(coefficient2.inv(level, c(PA,
      1 - PA, PB, 1 - PB)), alpha = FALSE))
  }

  PA <- seq(xlim[1], xlim[2], length.out = grid)
  PB <- seq(ylim[1], ylim[2], length.out = grid)
  data <- expand_grid(x = PA, y = PB)
  z <- rep(NA, grid^2)
  for (i in 1:grid) {
    for (j in 1:grid) {
      z[(i - 1) * grid + j] <- f(coefficient1, coefficient2.inv,
        PA[i], PB[j])
    }
  }
  data <- mutate(data, z)

  labels <- rep("", bins + 1)

  addlabels <- seq(from = -1, to = 1, length.out = 11)

  addlabelsat <- seq(from = 1, to = bins + 1, length.out = 11)

  labels[addlabelsat] <- addlabels

  g <- ggplot(data, aes(x = x, y = y, fill = z)) + geom_tile() +
    scale_fill_stepsn(n.breaks = bins, colors = hcl.colors(20,
      "Spectral"), limits = c(-1, 1), labels = labels) +
```

```

    guides(fill = guide_colourbar(title = NULL, ticks = FALSE,
      barheight = 10)) + xlab(xlab) + ylab(ylab)

    # 'Green-Brown' or 'Blue-Red 3' instead of 'Spectral'
    # are colorblind-safe, but for smaller ranges of values
    # do not distinguish as well

    g
  }
}

plot_contour_trivial <- function(level, xlim, ylim, grid, xlab,
  ylab, bins) {
  # function yielding a trivial heatmap with value
  # 'level', i.e. the whole square in one color
  f <- function(PA, PB) {
    level
  }

  PA <- seq(xlim[1], xlim[2], length.out = grid)
  PB <- seq(ylim[1], ylim[2], length.out = grid)
  data <- expand_grid(x = PA, y = PB)
  z <- rep(NA, grid^2)
  for (i in 1:grid) {
    for (j in 1:grid) {
      z[(i - 1) * grid + j] <- f(PA[i], PB[j])
    }
  }
  data <- mutate(data, z)

  labels <- rep("", bins + 1)

  addlabels <- seq(from = -1, to = 1, length.out = 11)

  addlabelsat <- seq(from = 1, to = bins + 1, length.out = 11)

  labels[addlabelsat] <- addlabels

  g <- ggplot(data, aes(x = x, y = y, fill = z)) + geom_tile() +
    scale_fill_stepsn(n.breaks = bins, colors = hcl.colors(20,
      "Spectral"), limits = c(-1, 1), labels = labels) +
    guides(fill = guide_colourbar(title = NULL, ticks = FALSE,
      barheight = 10)) + xlab(xlab) + ylab(ylab)

  # 'Green-Brown' or 'Blue-Red 3' instead of 'Spectral'
  # are colorblind-safe, but for smaller ranges of values
  # do not distinguish as well

  g
}

```

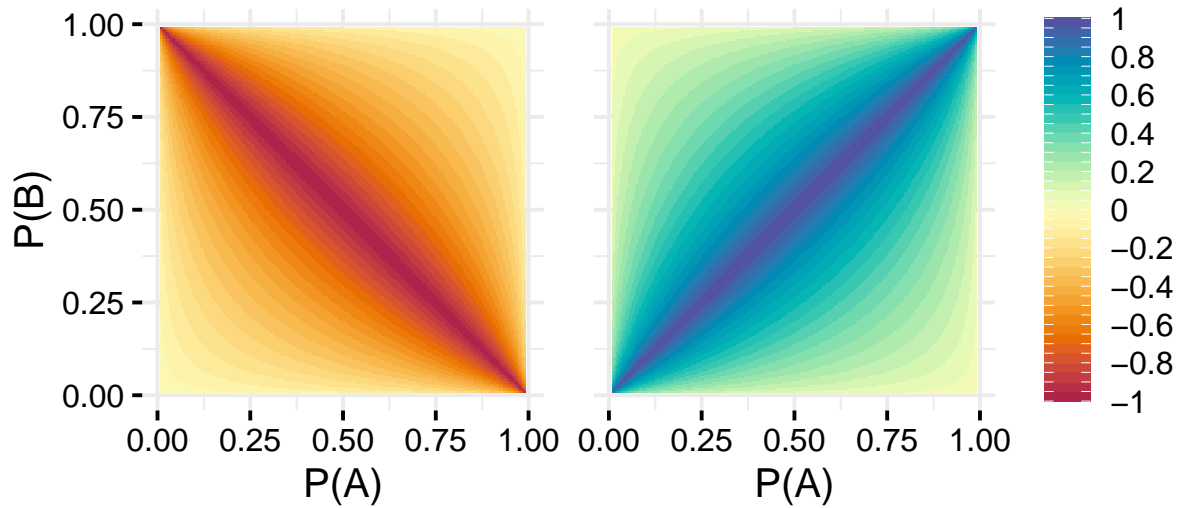
Including Plots

Bounds for Phi

```

gridsize <- 250
plot_contour_binary(Phi, Cole.inv, -1, c(0.01, 0.99), c(0.01,
  0.99), gridsize, "P(A)", "P(B)", 40) + coord_fixed() + theme_minimal(base_size = 15) +
  theme(axis.text.x = element_text(color = "black"), axis.text.y = element_text(color = "black"),
    axis.ticks = element_line(color = "black")) + plot_contour_binary(Phi,
  Cole.inv, 1, c(0.01, 0.99), c(0.01, 0.99), gridsize, "P(A)",
  "P(B)", 40) + coord_fixed() + theme_minimal(base_size = 15) +
  theme(axis.text.x = element_text(color = "black"), axis.ticks.x = element_line(color = "black"),
    axis.text.y = element_blank(), axis.ticks.y = element_blank(),
    axis.title.y = element_blank()) + plot_layout(guides = "collect",
  ncol = 2)

```



```

ggsave(filename = here("application/plots/phi_bounds.pdf"), width = 200,
  height = 90, device = "pdf", units = "mm")

```

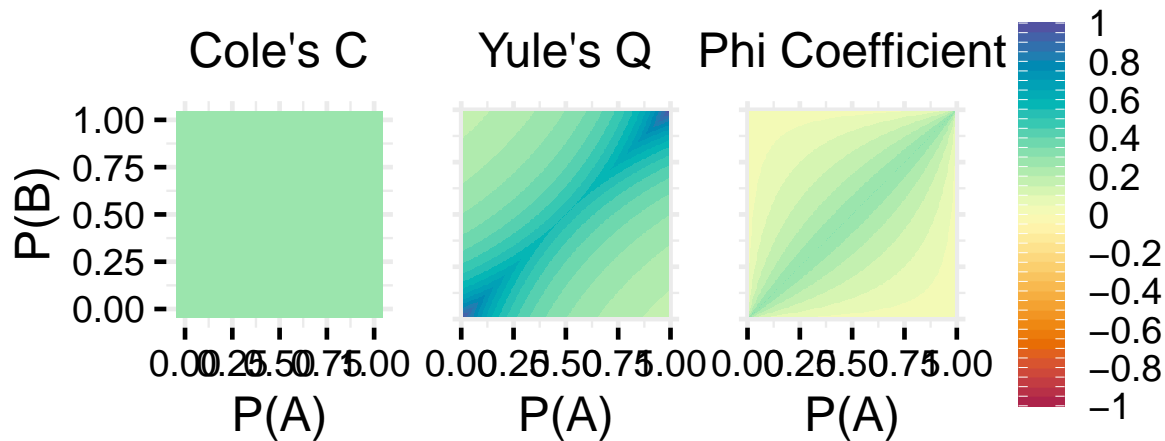
Heatmaps with Cole as the coefficient to be set and comparison with Cole, YuleQ and Phi

```

gridsize <- 250
adj_width <- 50
adj_height <- 95
base_size <- 18
plot_contour_trivial(0.3, c(0.01, 0.99), c(0.01, 0.99), 10, "P(A)",
  "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Cole's C") + theme(plot.title = element_text(hjust = 0.5,
  size = 18)) + theme(axis.text.x = element_text(color = "black"),
  axis.text.y = element_text(color = "black"), axis.ticks = element_line(color = "black")) +
  plot_contour_binary(YuleQ, Cole.inv, 0.3, c(0.01, 0.99),
  c(0.01, 0.99), gridsize, "P(A)", "P(B)", 40) + coord_fixed() +
  theme_minimal(base_size = base_size) + ggtitle("Yule's Q") +
  theme(plot.title = element_text(hjust = 0.5, size = 18)) +
  theme(axis.text.x = element_text(color = "black"), axis.ticks.x = element_line(color = "black"),
    axis.text.y = element_blank(), axis.ticks.y = element_blank(),
    axis.title.y = element_blank()) + plot_contour_binary(Phi,
  Cole.inv, 0.3, c(0.01, 0.99), c(0.01, 0.99), gridsize, "P(A)",
  "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Phi Coefficient") + theme(plot.title = element_text(hjust = 0.5,
  size = 18)) + theme(axis.text.x = element_text(color = "black"),
  axis.ticks.x = element_line(color = "black"), axis.text.y = element_blank(),

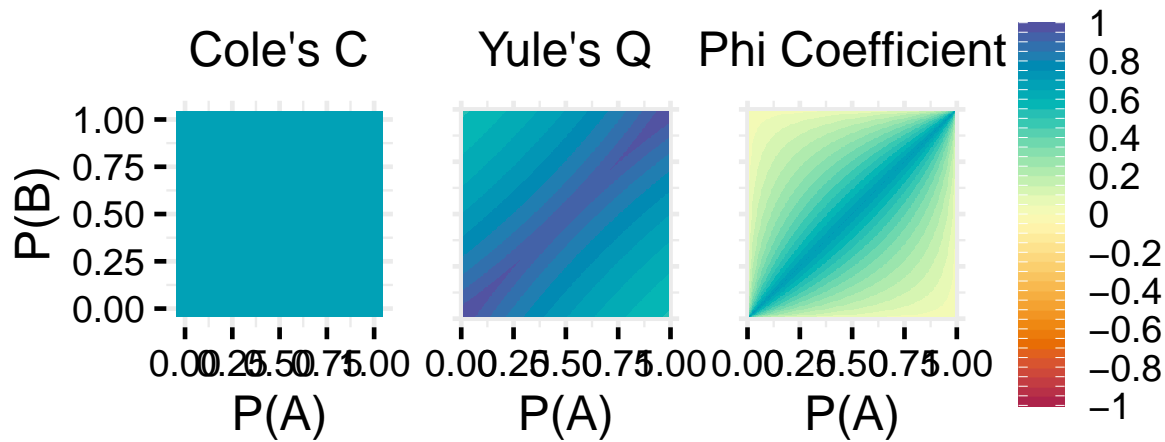
```

```
axis.ticks.y = element_blank(), axis.title.y = element_blank()) +
plot_layout(guides = "collect", ncol = 3)
```



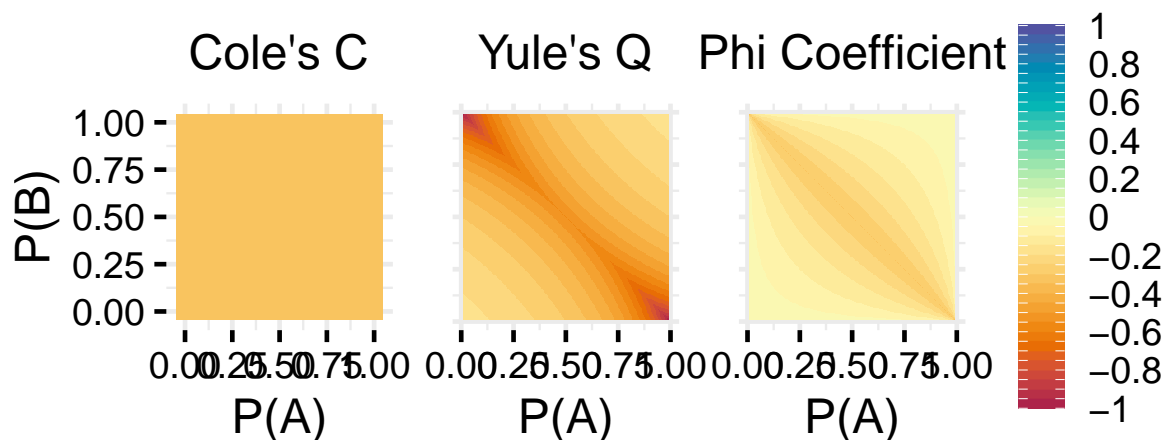
```
ggsave(filename = here("application/plots/cole_comparison_03.pdf"),
height = 200 - adj_height, width = (1 + sqrt(5))/2 * 200 -
adj_width, device = "pdf", units = "mm")

plot_contour_trivial(0.7, c(0.01, 0.99), c(0.01, 0.99), 10, "P(A)",
"P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
ggtitle("Cole's C") + theme(plot.title = element_text(hjust = 0.5,
size = 18)) + theme(axis.text.x = element_text(color = "black"),
axis.text.y = element_text(color = "black"), axis.ticks = element_line(color = "black")) +
plot_contour_binary(YuleQ, Cole.inv, 0.7, c(0.01, 0.99),
c(0.01, 0.99), gridsize, "P(A)", "P(B)", 40) + coord_fixed() +
theme_minimal(base_size = base_size) + ggtitle("Yule's Q") +
theme(plot.title = element_text(hjust = 0.5, size = 18)) +
theme(axis.text.x = element_text(color = "black"), axis.ticks.x = element_line(color = "black"),
axis.text.y = element_blank(), axis.ticks.y = element_blank(),
axis.title.y = element_blank()) + plot_contour_binary(Phi,
Cole.inv, 0.7, c(0.01, 0.99), c(0.01, 0.99), gridsize, "P(A)",
"P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
ggtitle("Phi Coefficient") + theme(plot.title = element_text(hjust = 0.5,
size = 18)) + theme(axis.text.x = element_text(color = "black"),
axis.ticks.x = element_line(color = "black"), axis.text.y = element_blank(),
axis.ticks.y = element_blank(), axis.title.y = element_blank()) +
plot_layout(guides = "collect", ncol = 3)
```



```
ggsave(filename = here("application/plots/cole_comparison_07.pdf"),
        height = 200 - adj_height, width = (1 + sqrt(5))/2 * 200 -
        adj_width, device = "pdf", units = "mm")

plot_contour_trivial(-0.3, c(0.01, 0.99), c(0.01, 0.99), 10,
  "P(A)", "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Cole's C") + theme(plot.title = element_text(hjust = 0.5,
    size = 18)) + theme(axis.text.x = element_text(color = "black"),
  axis.text.y = element_text(color = "black"), axis.ticks = element_line(color = "black")) +
  plot_contour_binary(YuleQ, Cole.inv, -0.3, c(0.01, 0.99),
    c(0.01, 0.99), gridsize, "P(A)", "P(B)", 40) + coord_fixed() +
  theme_minimal(base_size = base_size) + ggtitle("Yule's Q") +
  theme(plot.title = element_text(hjust = 0.5, size = 18)) +
  theme(axis.text.x = element_text(color = "black"), axis.ticks.x = element_line(color = "black"),
    axis.text.y = element_blank(), axis.ticks.y = element_blank(),
    axis.title.y = element_blank()) + plot_contour_binary(Phi,
  Cole.inv, -0.3, c(0.01, 0.99), c(0.01, 0.99), gridsize, "P(A)",
  "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Phi Coefficient") + theme(plot.title = element_text(hjust = 0.5,
    size = 18)) + theme(axis.text.x = element_text(color = "black"),
  axis.ticks.x = element_line(color = "black"), axis.text.y = element_blank(),
  axis.ticks.y = element_blank(), axis.title.y = element_blank()) +
  plot_layout(guides = "collect", ncol = 3)
```



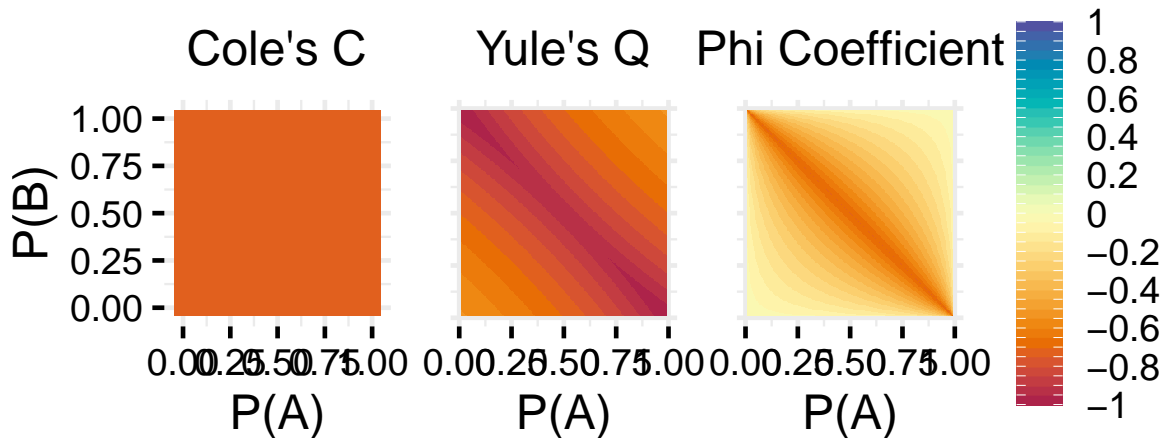
```
ggsave(filename = here("application/plots/cole_comparison_-03.pdf"),
        height = 200 - adj_height, width = (1 + sqrt(5))/2 * 200 -
```

```

adj_width, device = "pdf", units = "mm")

plot_contour_trivial(-0.7, c(0.01, 0.99), c(0.01, 0.99), 10,
  "P(A)", "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Cole's C") + theme(plot.title = element_text(hjust = 0.5,
    size = 18)) + theme(axis.text.x = element_text(color = "black"),
    axis.text.y = element_text(color = "black"), axis.ticks = element_line(color = "black")) +
  plot_contour_binary(YuleQ, Cole.inv, -0.7, c(0.01, 0.99),
    c(0.01, 0.99), gridsize, "P(A)", "P(B)", 40) + coord_fixed() +
  theme_minimal(base_size = base_size) + ggtitle("Yule's Q") +
  theme(plot.title = element_text(hjust = 0.5, size = 18)) +
  theme(axis.text.x = element_text(color = "black"), axis.ticks.x = element_line(color = "black"),
    axis.text.y = element_blank(), axis.ticks.y = element_blank(),
    axis.title.y = element_blank()) + plot_contour_binary(Phi,
  Cole.inv, -0.7, c(0.01, 0.99), c(0.01, 0.99), gridsize, "P(A)",
  "P(B)", 40) + coord_fixed() + theme_minimal(base_size = base_size) +
  ggtitle("Phi Coefficient") + theme(plot.title = element_text(hjust = 0.5,
    size = 18)) + theme(axis.text.x = element_text(color = "black"),
    axis.ticks.x = element_line(color = "black"), axis.text.y = element_blank(),
    axis.ticks.y = element_blank(), axis.title.y = element_blank()) +
  plot_layout(guides = "collect", ncol = 3)

```



```

ggsave(filename = here("application/plots/cole_comparison_-07.pdf"),
  height = 200 - adj_height, width = (1 + sqrt(5))/2 * 200 -
  adj_width, device = "pdf", units = "mm")

```