# Tile Map to depict mortality in the US

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This is a reproducible example to produce a tile map that shows mortality rates in the USA.

#### Libraries

```
library(tidyverse)
library(geofacet)
library(kani)

options(scipen = 99)
```

**Note:** I have used a library called kani which has some theme aesthetics for plotting. It can be installed by using devtools::install\_github("kanishkamisra/kani") in your R console.

#### Data import

We use data from IMHE's website that has gender-specific data about mortality rates due to different causes in the USA and its counties. We will only be dealing with mortality rates due to all causes in all states (present along with the counties) of the US(Including Hawaii and District of Columbia).

```
files <- list.files(pattern = '*.CSV', full.names = T, recursive = T)

mortality <- map_df(files, read_csv)

states <- data.frame(
    state_abb = c(state.abb, "DC"),
    location_name = c(state.name, "District of Columbia")
)

mortality <- mortality %>%
    filter(location_name %in% state.name | location_name == "District of Columbia") %>%
    filter(cause_name == "All causes" & sex == "Both") %>%
    inner_join(states)
```

### US Average for each year

```
usa_vs_state <- mortality %>%
  group_by(year_id) %>%
  mutate(usa_avg = mean(mx)) %>%
  ungroup() %>%
  select(
    state = location_name,
    state_abb,
    year = year_id,
    state_avg = mx,
```

```
usa_avg
) %>%
mutate(
    ribbon_color = case_when(
        state_avg > usa_avg ~ "#f8766d",
        usa_avg > state_avg ~ "#00bfc4"
),
    ribbon_value = case_when(
        state_avg > usa_avg ~ state_avg,
        usa_avg > state_avg ~ usa_avg,
        TRUE ~ state_avg
)
```

To produce the plot, we use the <code>geom\_ribbon</code> function to show more a more apparent difference in the state vs the usa average mortality due to all causes. The ribbon is essentially a color fill between the two line plots of the state mortality rates and the usa average. The color is determined by which rate is greater (worse in the case of mortality).

```
usa_state_plot <- usa_vs_state %>%
  gather(state_avg, usa_avg, key = "metric", value = "mortality_rate") %>%
  separate(metric, into = c("metric", "avg")) %>%
  mutate(metric = str_to_title(metric)) %>%
  select(-avg) %>%
  ggplot(aes(year, mortality_rate, color = metric)) +
  geom_line(size = 1) +
  geom_ribbon(aes(ymin = mortality_rate, ymax = ribbon_value,
                  linetype = NA, fill = ribbon color), alpha = 0.2, show.legend = F) +
  facet_geo(~state_abb, grid = "us_state_grid4") +
  scale fill identity() +
  theme_kani() +
  theme(
   legend.position = "top",
   legend.margin = margin(b = -1, unit = "cm"),
   plot.background = element_rect(fill = "white"),
   panel.background = element_rect(fill = "white"),
   legend.background = element_rect(fill = "white"),
   legend.key = element_rect(fill = "white"),
   strip.background = element_rect(fill = "white"),
   strip.text = element text(face = "bold"),
   legend.text = element_text(size = rel(1.2)),
   legend.title = element_text(size = rel(1.2))
  ) +
  labs(
   x = "Year",
   y = "Mortality Rate (per 100,000 people)",
   color = "Trend"
  )
ggsave("usa_state_mortality.png", usa_state_plot, height = 12, width = 15)
```

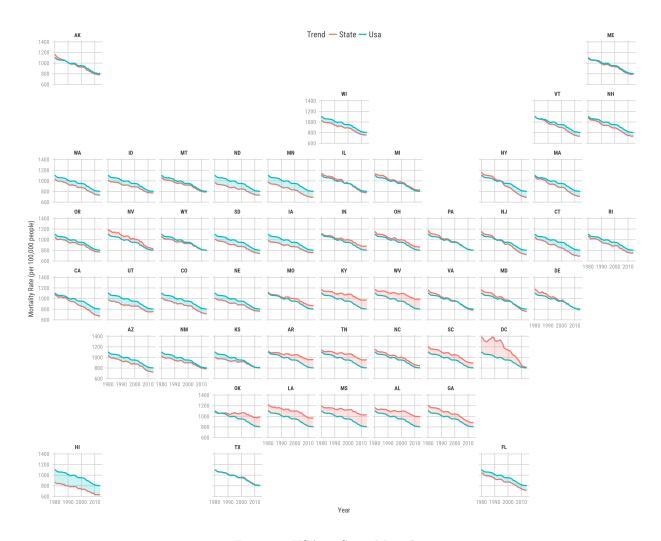


Figure 1: USA vs State Mortality

## Citation

Institute for Health Metrics and Evaluation (IHME). United States Mortality Rates by County 1980-2014. Seattle, United States: Institute for Health Metrics and Evaluation (IHME), 2016.