## heatmap

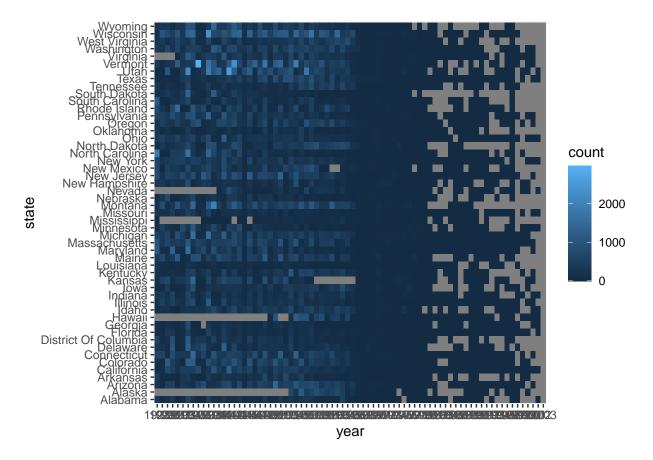
#### Hanna Rodrigues Ferreira

13 junho,2021

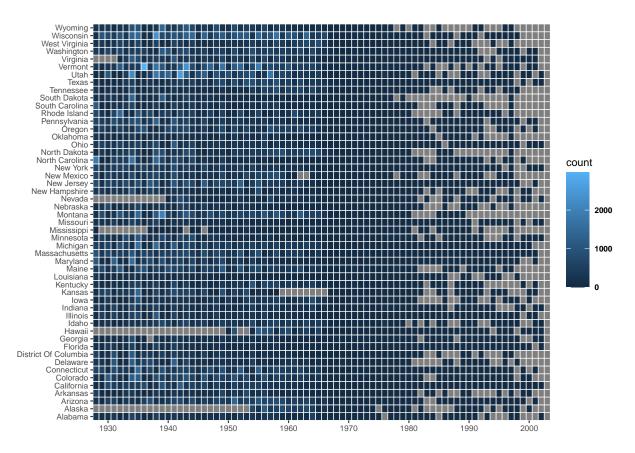
```
knitr::opts_chunk$set(echo = TRUE)
library(ggplot2)
library(dplyr)
library(tidyr)
library(stringr)
m <- read.csv("measles_lev1.csv",header=T,stringsAsFactors=F,skip=2)</pre>
m2 <- m \%>\%
  # convert data to long format
  gather (key="state",
         value="value",
         -YEAR,-WEEK) %>%
  # rename columns
  setNames(c("year",
              "week",
              "state",
             "value")) %>%
  mutate(year=factor(year)) %>%
  mutate(week=factor(week)) %>%
  mutate(value=as.numeric(value))
## Warning in mask$eval_all_mutate(quo): NAs introduzidos por coerção
# removes . and change states to title case using custom function
fn_tc <- function(x) paste(str_to_title(unlist(strsplit(x,"[.]"))),collapse=" ")</pre>
m2$state <- sapply(m2$state,fn_tc)</pre>
# custom sum function returns NA when all values in set are NA,
# in a set mixed with NAs, NAs are removed and remaining summed.
na_sum <- function(x)</pre>
  if(all(is.na(x))) val <- sum(x,na.rm=F)</pre>
  if(!all(is.na(x))) val <- sum(x,na.rm=T)</pre>
  return(val)
}
# sum incidences for all weeks into one year
m3 <- m2 %>%
```

```
group_by(year,state) %>%
summarise(count=na_sum(value)) %>%
as.data.frame()
```

## 'summarise()' has grouped output by 'year'. You can override using the '.groups' argument.

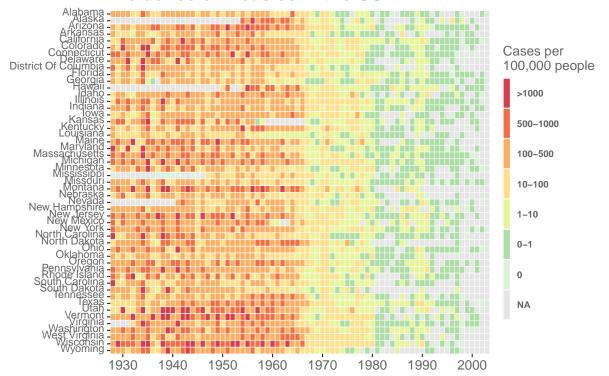


```
theme_grey(base_size=8)+
#theme options
theme(
    #bold font for legend text
    legend.text=element_text(face="bold"),
    #set thickness of axis ticks
    axis.ticks=element_line(size=0.4),
    #remove plot background
    plot.background=element_blank(),
    #remove plot border
    panel.border=element_blank())
```



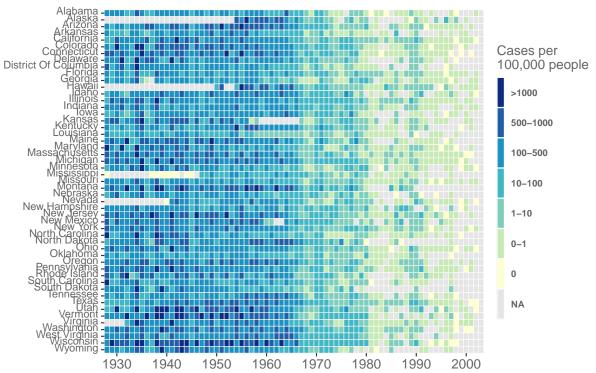
```
# assign text colour
textcol <- "grey40"
# further modified ggplot
p <- ggplot(m4,aes(x=year,y=state,fill=countfactor))+</pre>
  geom_tile(colour="white",size=0.2)+
  guides(fill=guide_legend(title="Cases per\n100,000 people"))+
  labs(x="",y="",title="Incidence of Measles in the US")+
  scale_y_discrete(expand=c(0,0))+
  scale_x_discrete(expand=c(0,0),breaks=c("1930","1940","1950","1960","1970","1980","1990","2000"))+
  scale_fill_manual(values=c("#d53e4f","#f46d43","#fdae61","#fee08b","#e6f598","#abdda4","#ddf1da"),na.
  #coord_fixed()+
  theme_grey(base_size=10)+
  theme(legend.position="right",legend.direction="vertical",
        legend.title=element_text(colour=textcol),
        legend.margin=margin(grid::unit(0,"cm")),
        legend.text=element_text(colour=textcol,size=7,face="bold"),
        legend.key.height=grid::unit(0.8, "cm"),
        legend.key.width=grid::unit(0.2,"cm"),
        axis.text.x=element_text(size=10,colour=textcol),
        axis.text.y=element_text(vjust=0.2,colour=textcol),
        axis.ticks=element_line(size=0.4),
        plot.background=element_blank(),
        panel.border=element_blank(),
        plot.margin=margin(0.7,0.4,0.1,0.2,"cm"),
        plot.title=element_text(colour=textcol,hjust=0,size=14,face="bold"))
```

#### Incidence of Measles in the US



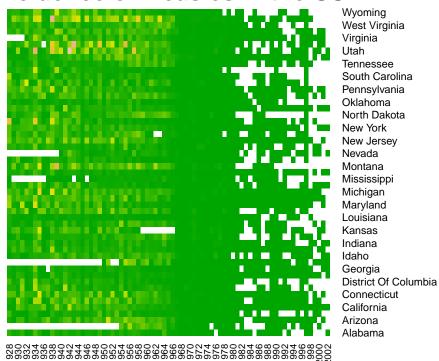
```
library(RColorBrewer)
# further modified ggplot
p <- ggplot(m4,aes(x=year,y=state,fill=countfactor))+</pre>
  geom_tile(colour="white",size=0.2)+
  guides(fill=guide_legend(title="Cases per\n100,000 people"))+
  labs(x="",y="",title="Incidence of Measles in the US")+
  scale y discrete(expand=c(0,0))+
  scale x discrete(expand=c(0,0),breaks=c("1930","1940","1950","1960","1970","1980","1980","1990","2000"))+
  scale_fill_manual(values=rev(brewer.pal(7,"YlGnBu")),na.value="grey90")+
  theme grey(base size=10)+
  theme(legend.position="right",legend.direction="vertical",
        legend.title=element text(colour=textcol),
        legend.margin=margin(grid::unit(0,"cm")),
        legend.text=element_text(colour=textcol,size=7,face="bold"),
        legend.key.height=grid::unit(0.8,"cm"),
        legend.key.width=grid::unit(0.2, "cm"),
        axis.text.x=element_text(size=10,colour=textcol),
        axis.text.y=element_text(vjust=0.2,colour=textcol),
        axis.ticks=element_line(size=0.4),
        plot.background=element_blank(),
        panel.border=element_blank(),
        plot.margin=margin(0.7,0.4,0.1,0.2,"cm"),
        plot.title=element_text(colour=textcol,hjust=0,size=14,face="bold"))
р
```





```
# load package
library(gplots) # heatmap.2() function
## Warning: package 'gplots' was built under R version 4.0.5
##
## Attaching package: 'gplots'
## The following object is masked from 'package:stats':
##
##
       lowess
library(plotrix) # gradient.rect() function
##
## Attaching package: 'plotrix'
## The following object is masked from 'package:gplots':
##
##
       plotCI
```

### Incidence of Measles in the US



# **Incidence of Measles in the US**

