epiChart

Allison Collins 4/2/2020

Load data from the simulations

```
#load + concatenate the csvs
df <- list.files(path = "epi_csvs",full.names = TRUE) %>%
 lapply(read_csv) %>%
 bind rows
## Parsed with column specification:
## cols(
##
     `#dummy file` = col_character()
## )
## Parsed with column specification:
## cols(
##
    X1 = col_integer(),
    time = col_integer(),
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    S = col_double(),
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    E = col_double(),
    I = col_double(),
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    H = col_double(),
    C = col_double(),
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    R = col_double(),
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    hosp = col_double(),
     crits = col_double()
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```

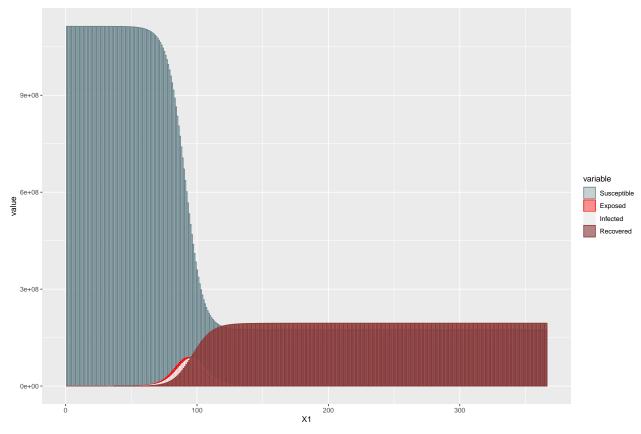
```
## Parsed with column specification:
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##
##
     crits = col_double()
## )
#create summary dataframe
new_df <- df %>%
  group_by(X1) %>%
  summarise(Susceptible = sum(S), Exposed = sum(E), Infected = sum(I), Recovered = sum(R))
```

Make chart



Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.