

Visualization solutions

Introduction

This is a (partial) key for the exercises in the handout *Visualization*

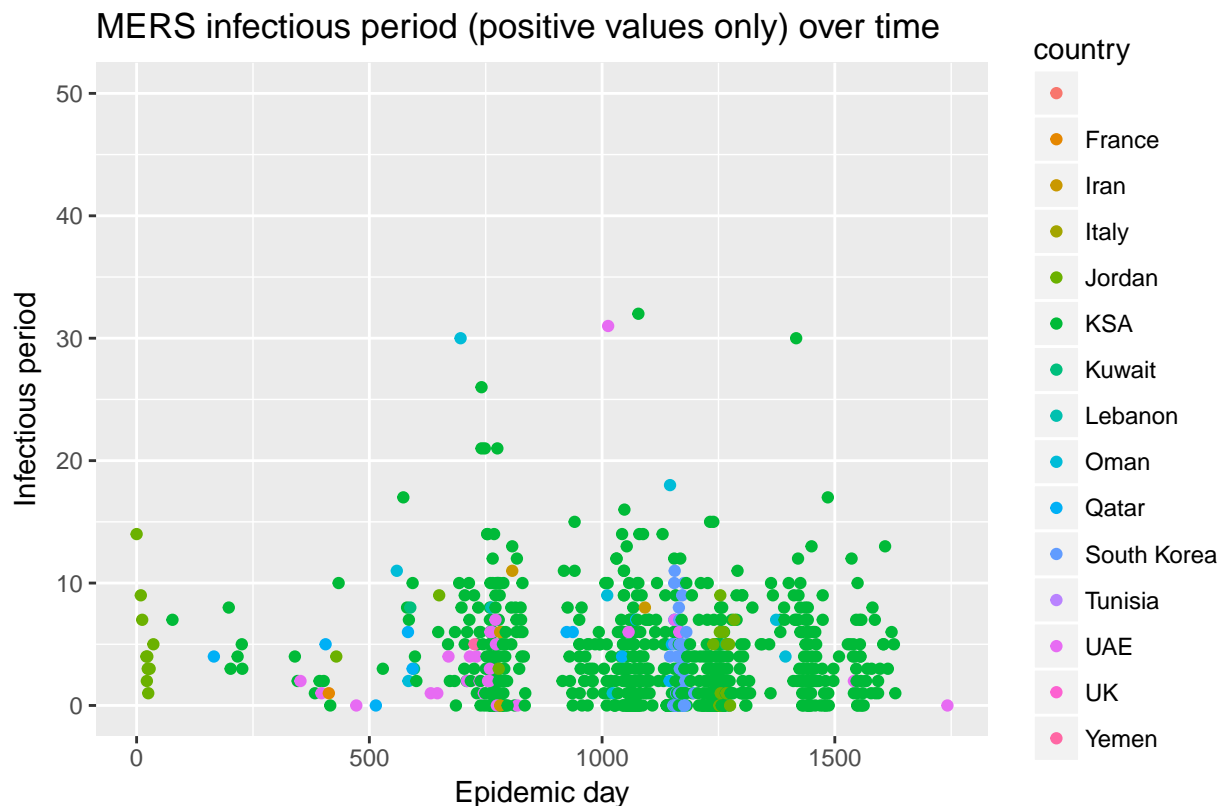
Exercise. Use our corrected infectious period variable (`infectious.period2`) to study the change in the infectious period over the course of the MERS epidemic.

```
## [1] "Date"
```

```
## [1] "difftime"
```

```
mers$infectious.period2 <- ifelse(mers$infectious.period<0,0,mers$infectious.period)
```

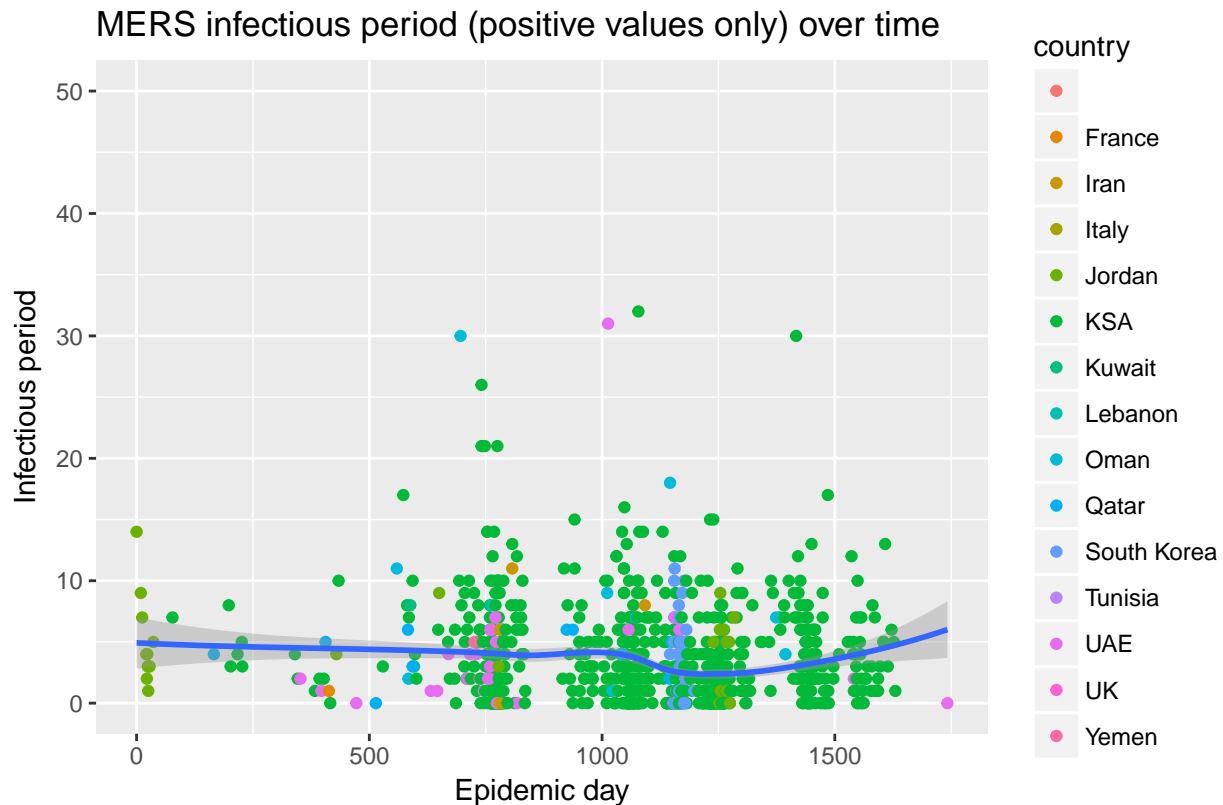
```
ggplot(data=mers, mapping=aes(x=epi.day, y=infectious.period2)) +  
  geom_point(mapping = aes(color=country)) +  
  scale_y_continuous(limits = c(0, 50)) +  
  labs(x='Epidemic day', y='Infectious period',  
       title='MERS infectious period (positive values only) over time', caption="Data from: https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv")
```



Data from: <https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv>

Exercise. In data from many outbreaks it can be seen that there is a kind of *societal learning*. When the infection first emerges it is not quickly recognized, public health resources have not been mobilized, it is not known what symptoms are diagnostic, how to treat, etc. But, quickly, this information is collected and the outbreak is contained. Is there evidence of this kind of societal learning in the mers data. Add a curve fit using `geom_smooth` to explore this question. Hint: I solved using the `loess` method because the default smoother (`gam`) failed.

```
ggplot(data=mers, mapping=aes(x=epi.day, y=infectious.period2)) +
  geom_point(mapping = aes(color=country)) +
  geom_smooth(method='loess') +
  scale_y_continuous(limits = c(0, 50)) +
  labs(x='Epidemic day', y='Infectious period',
        title='MERS infectious period (positive values only) over time', caption="Data from: https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv")
```

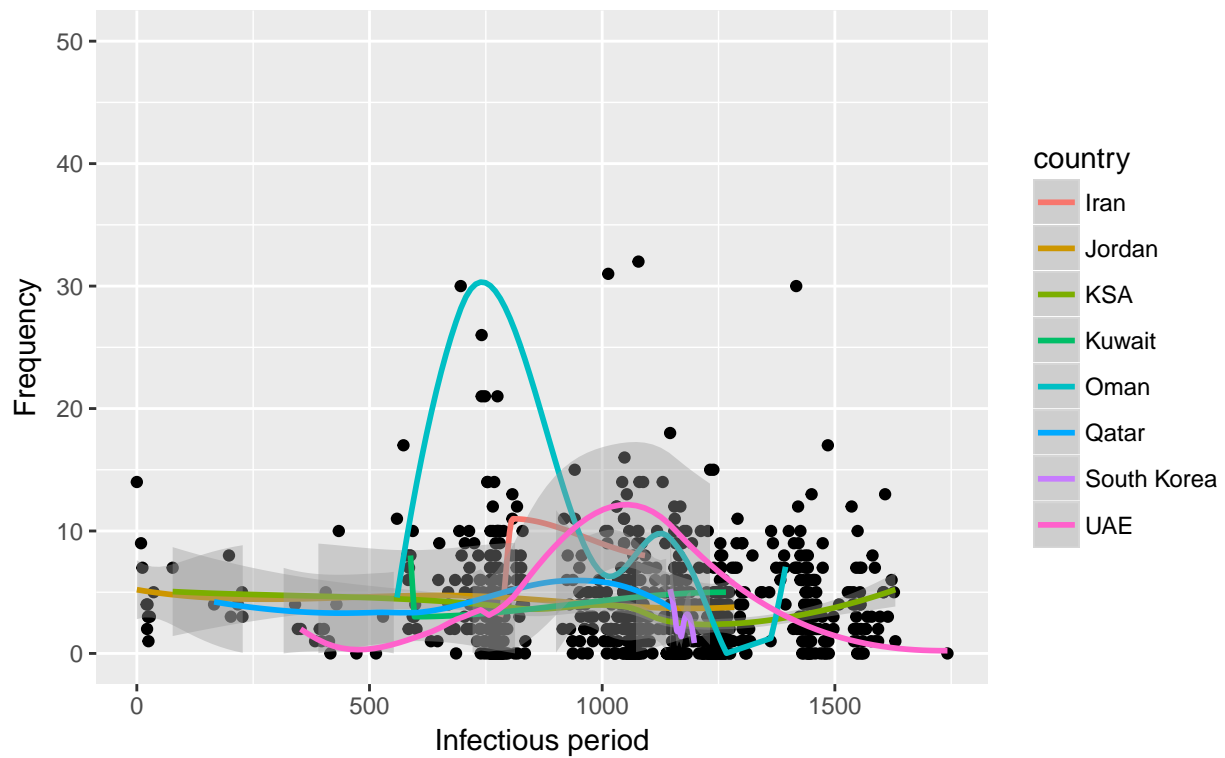


Data from: <https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv>

Exercise. Plot `infectious.period2` against time, as before, but this time add a separate smooth fit for each country.

```
ggplot(data=mers, mapping=aes(x=epi.day, y=infectious.period2)) +
  geom_point() +
  geom_smooth(method='loess', mapping = aes(color=country)) +
  scale_y_continuous(limits = c(0, 50)) +
  labs(x='Infectious period', y='Frequency',
        title='MERS infectious period (positive values only) over time', caption="Data from: https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv")
```

MERS infectious period (positive values only) over time



Data from: <https://github.com/rambaut/MERS-Cases/blob/gh-pages/data/cases.csv>