

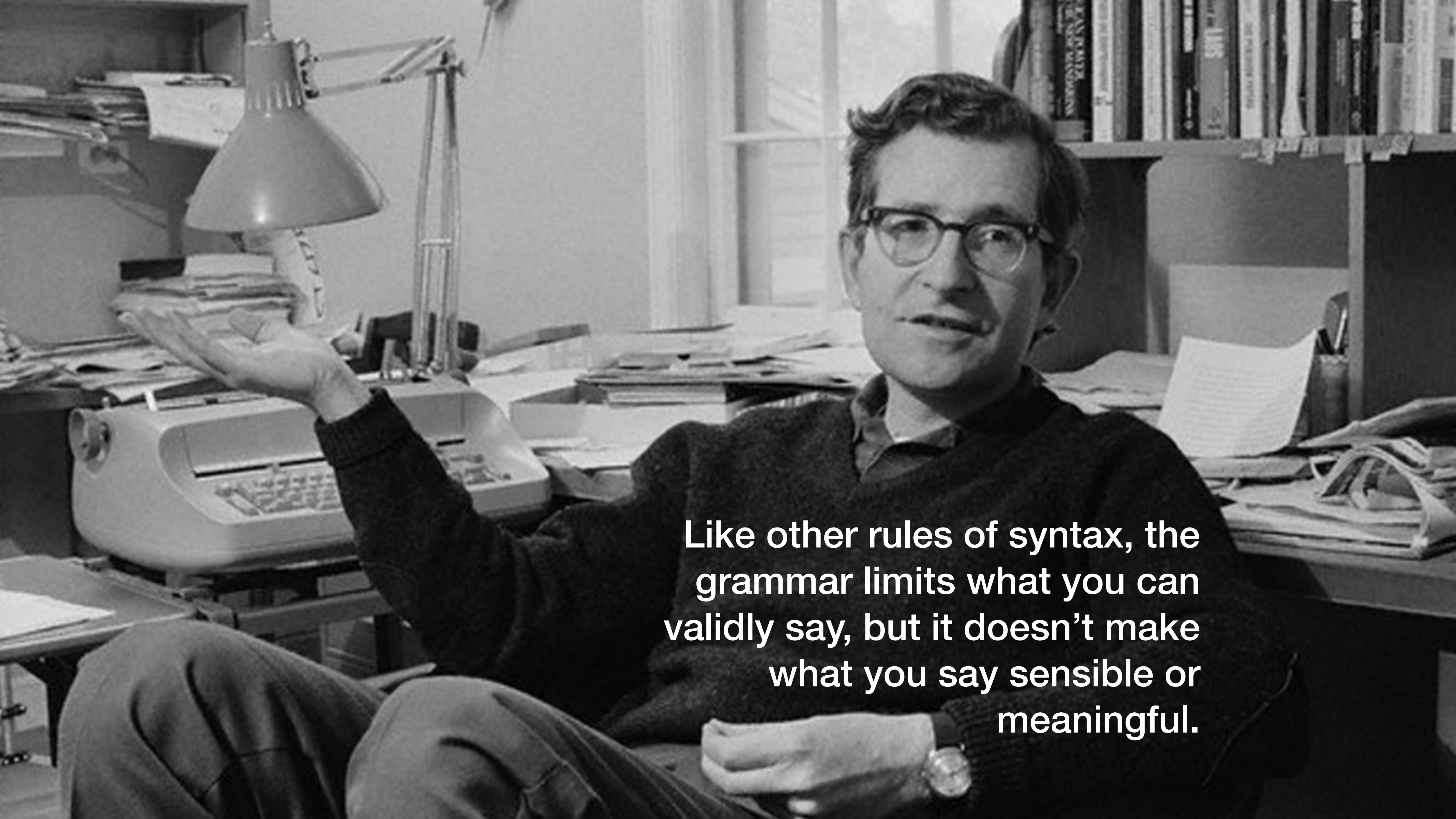
# Show the Right Numbers



ggplot  
IMPLEMENTS  
A GRAMMAR  
OF GRAPHICS

The grammar is a set of rules for how produce graphics from data, taking **pieces of data** and **mapping** them to **geometric objects** (like points and lines) **that have aesthetic attributes** (like position, color and size), together with further rules for **transforming the data if needed**, adjusting **scales**, or projecting the results onto a **coordinate system**.

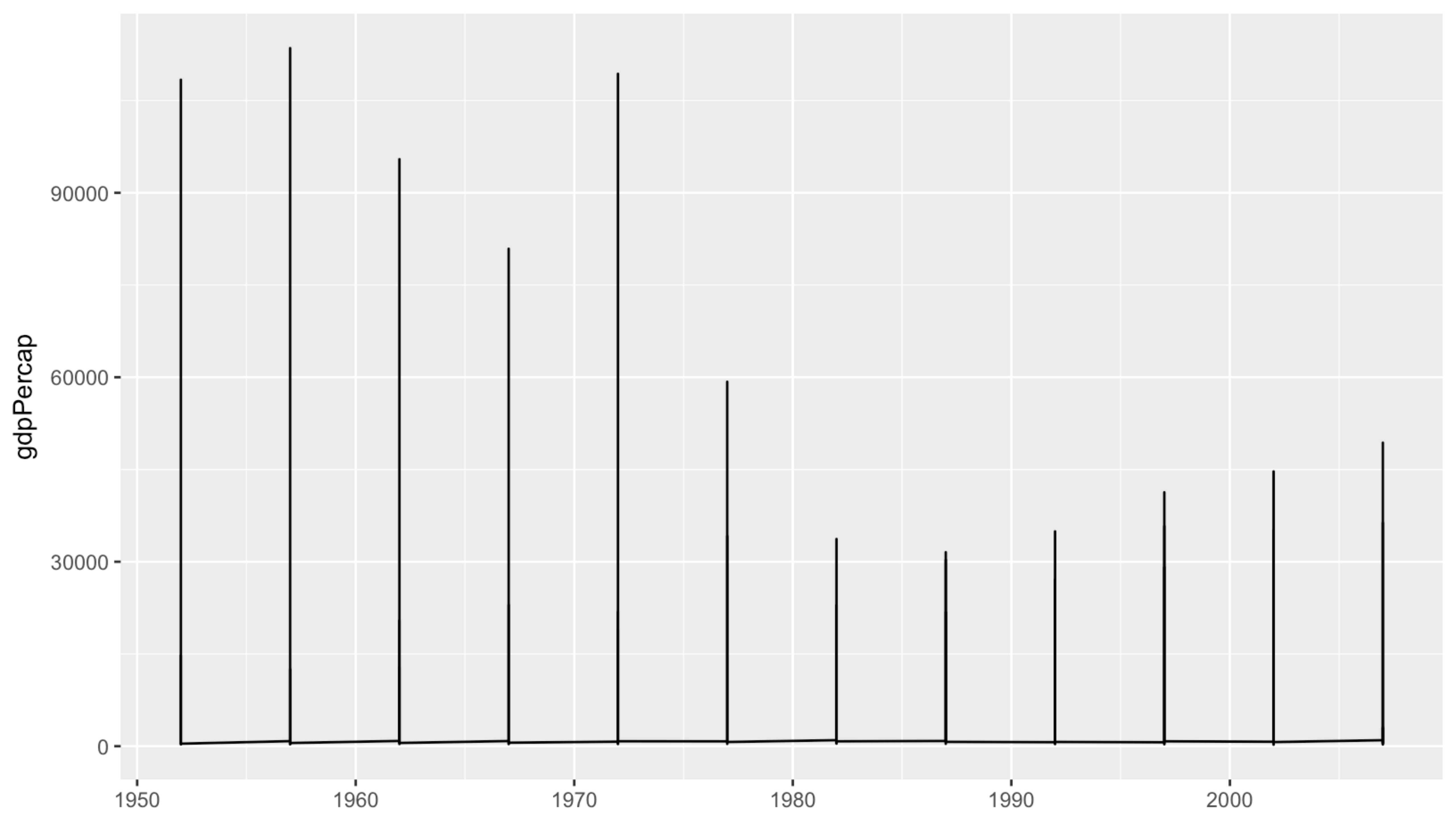


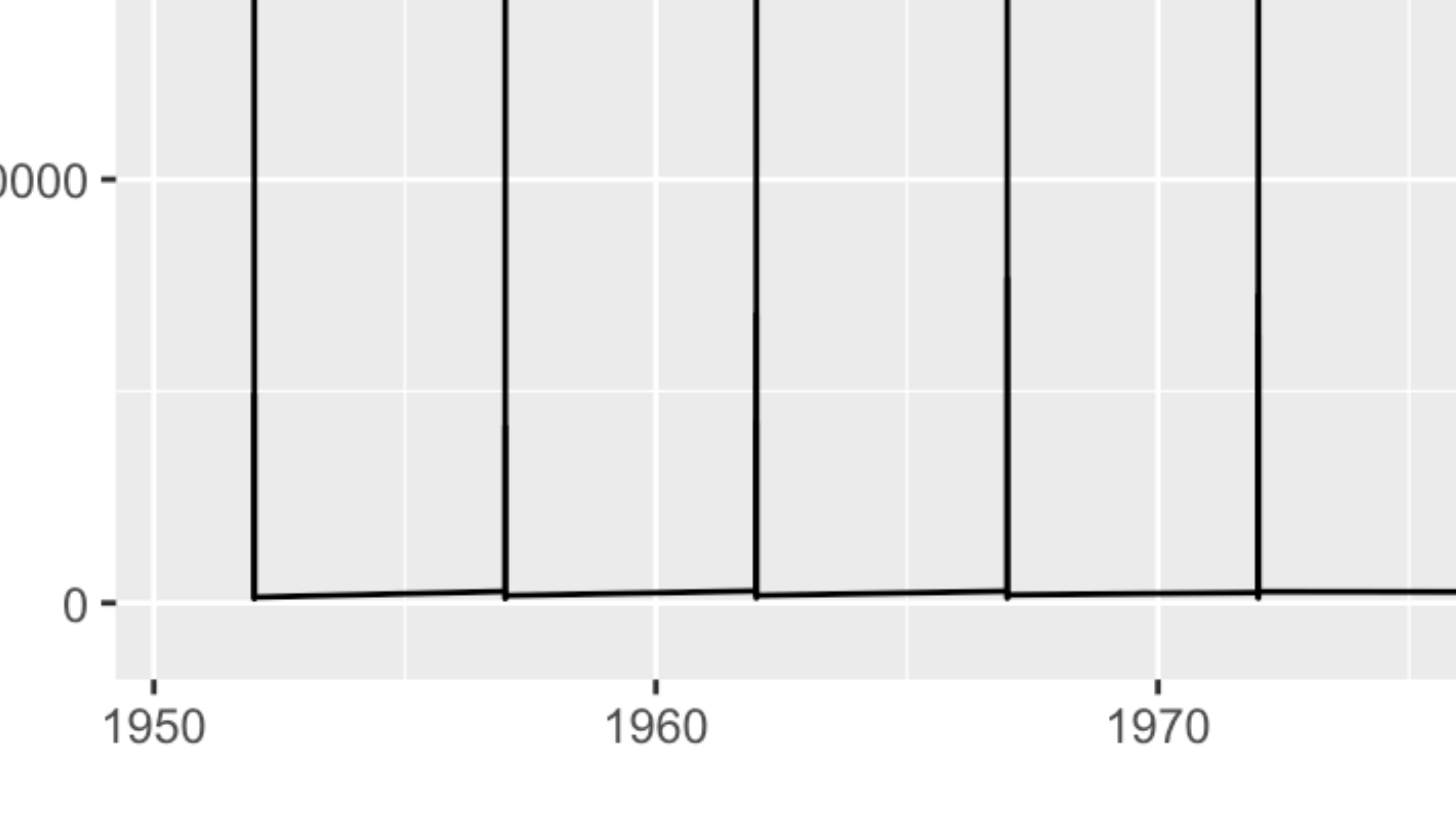


Like other rules of syntax, the  
grammar limits what you can  
validly say, but it doesn't make  
what you say sensible or  
meaningful.

# Grouped Data and the group aesthetic

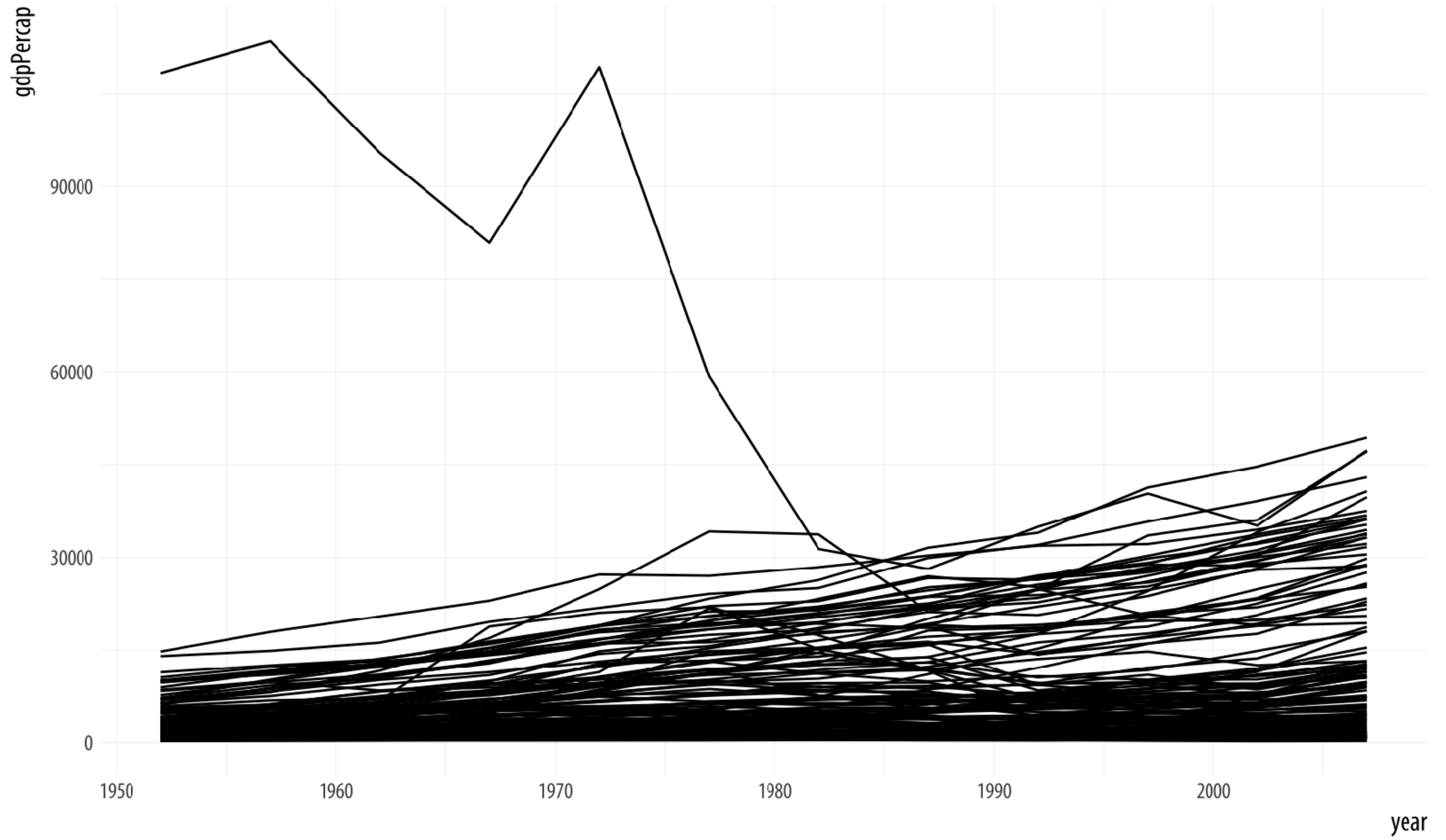
[illegible]





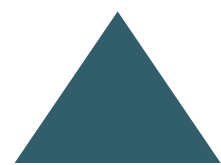


```
p <- ggplot(data = gapminder,  
            mapping = aes(x = year,  
                           y = gdpPerCap))  
p + geom_line(mapping = aes(group = country))
```

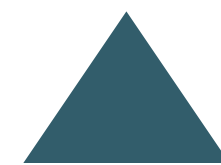


```
p <- ggplot(data = gapminder,  
            mapping = aes(x = year,  
                          y = gdpPercap))
```

```
p + geom_line(mapping =  
              aes(group = country)) +  
  facet_wrap(~ continent)
```

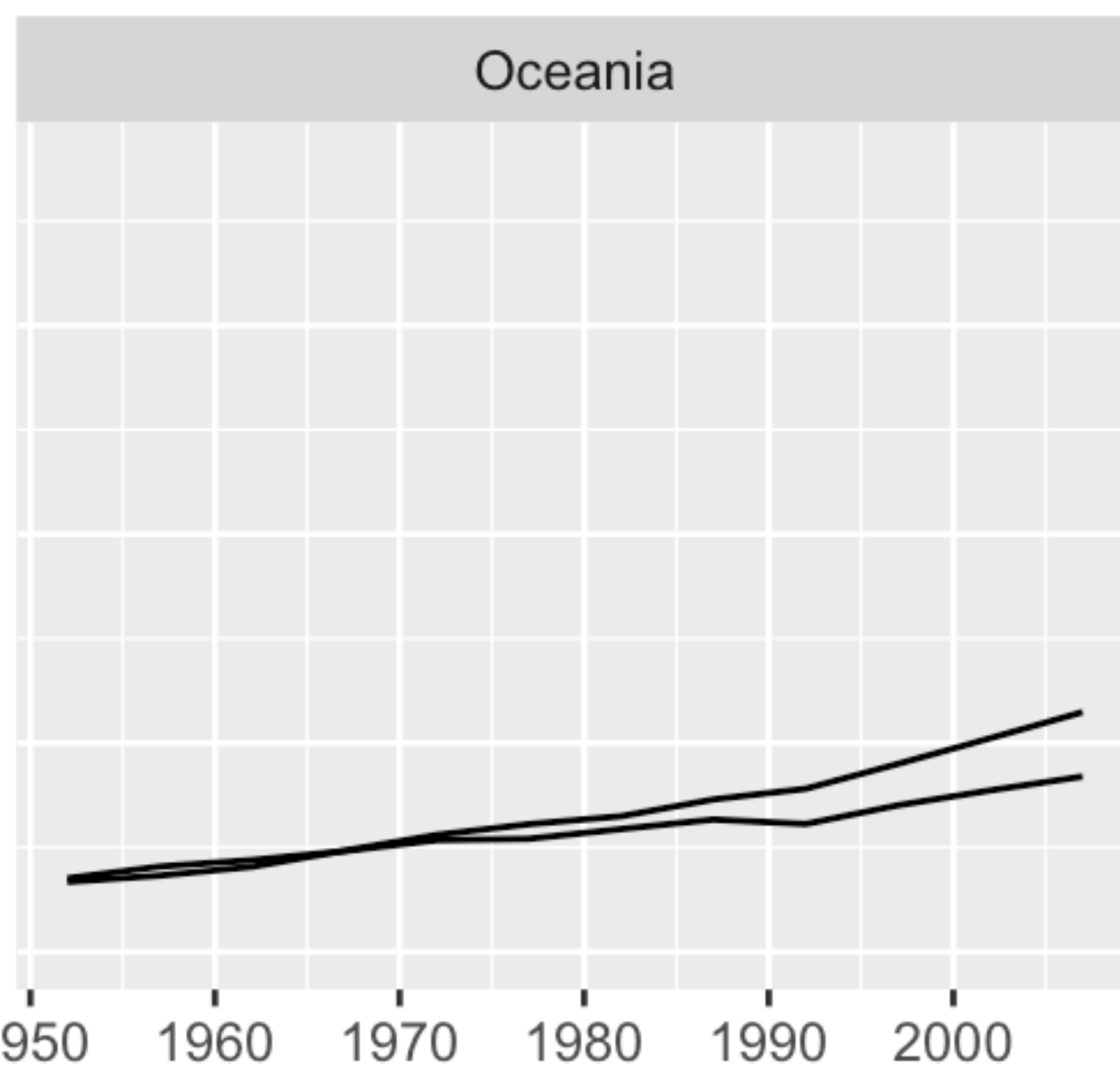
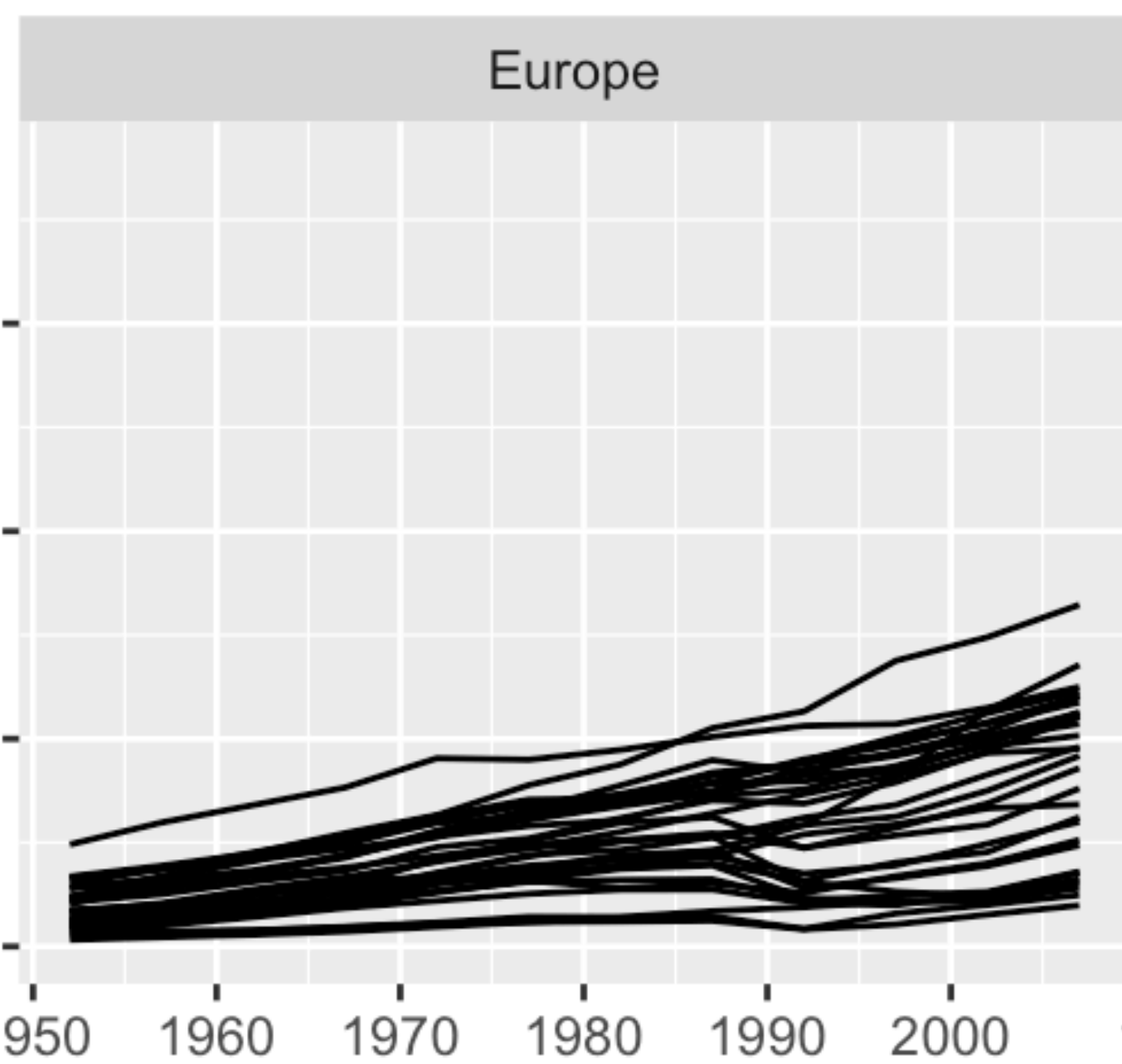
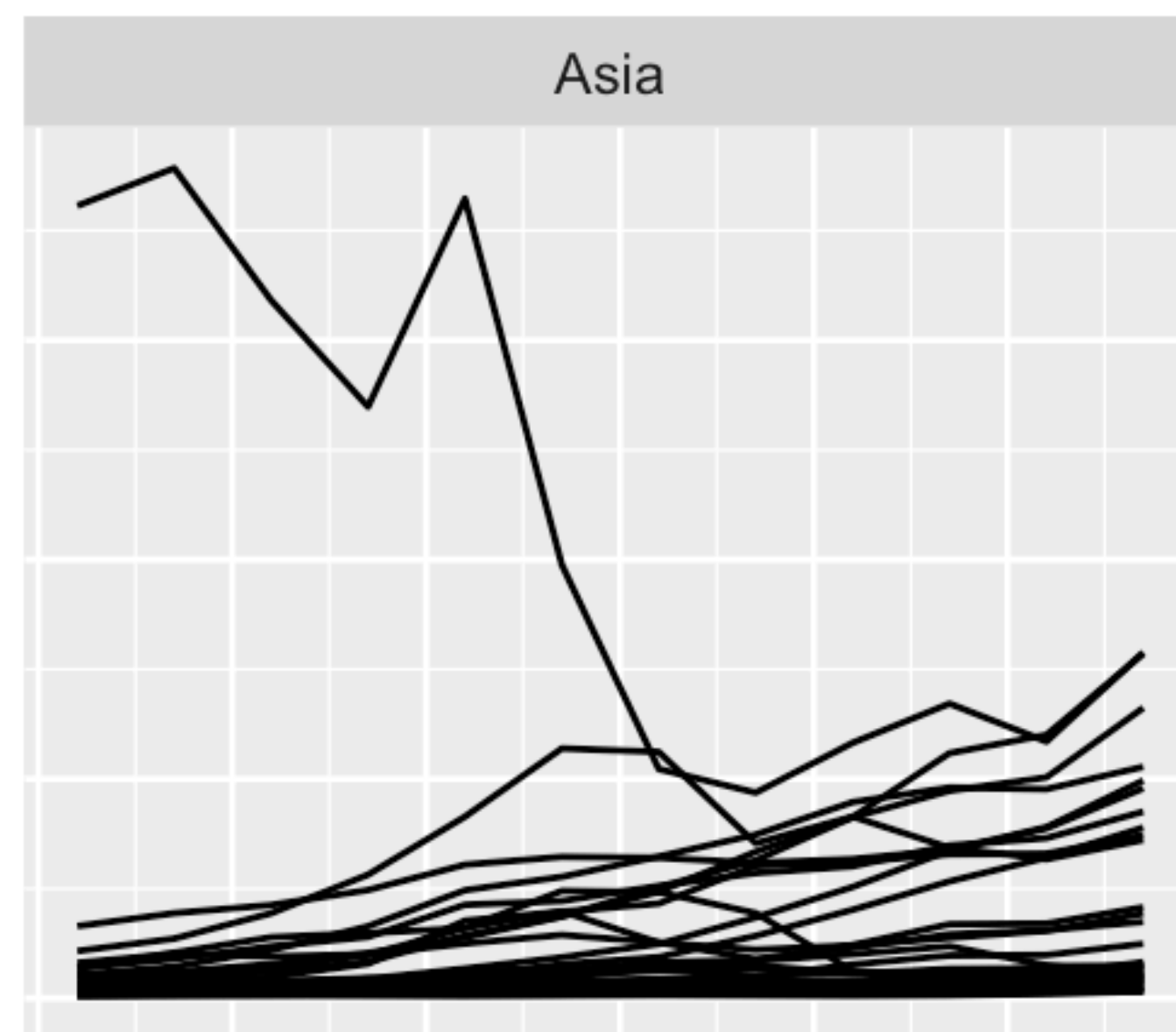
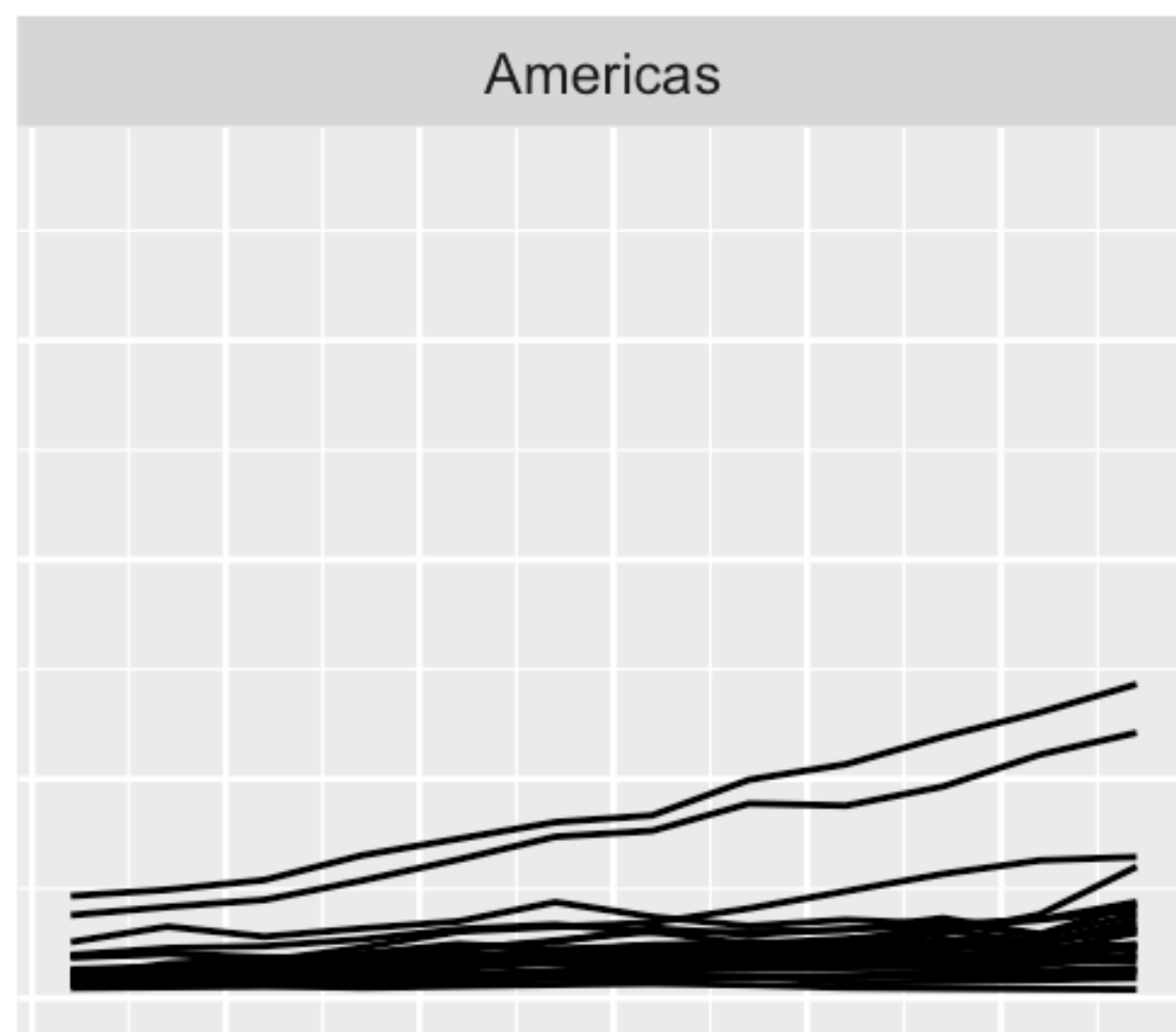
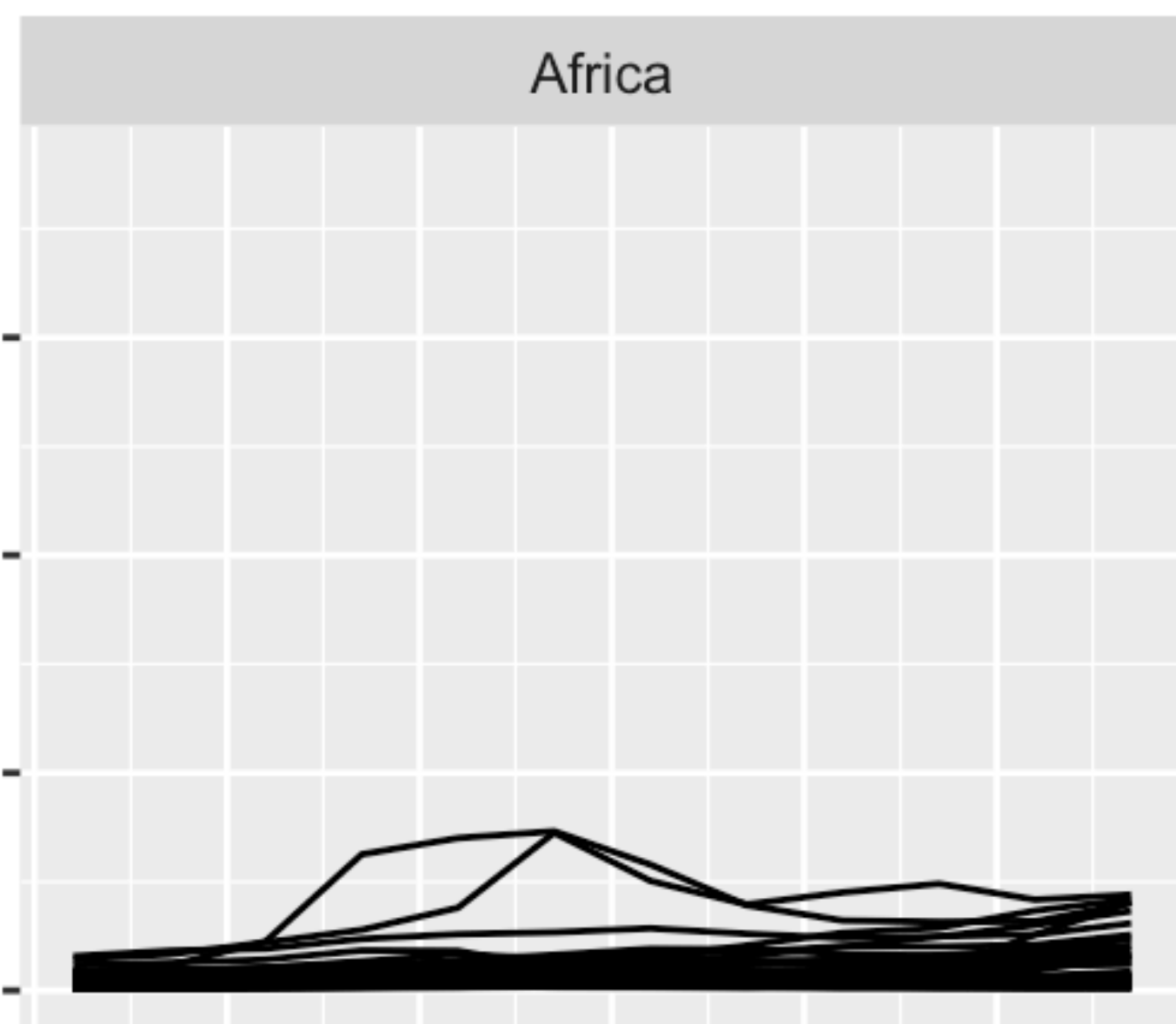


A facet is not a  
geom. It's a way  
of arranging geoms.



Facets use R's  
'formula' syntax. Read  
the ~ as "on" or "by".


gdpPercap



1950 1960 1970 1980 1990 2000

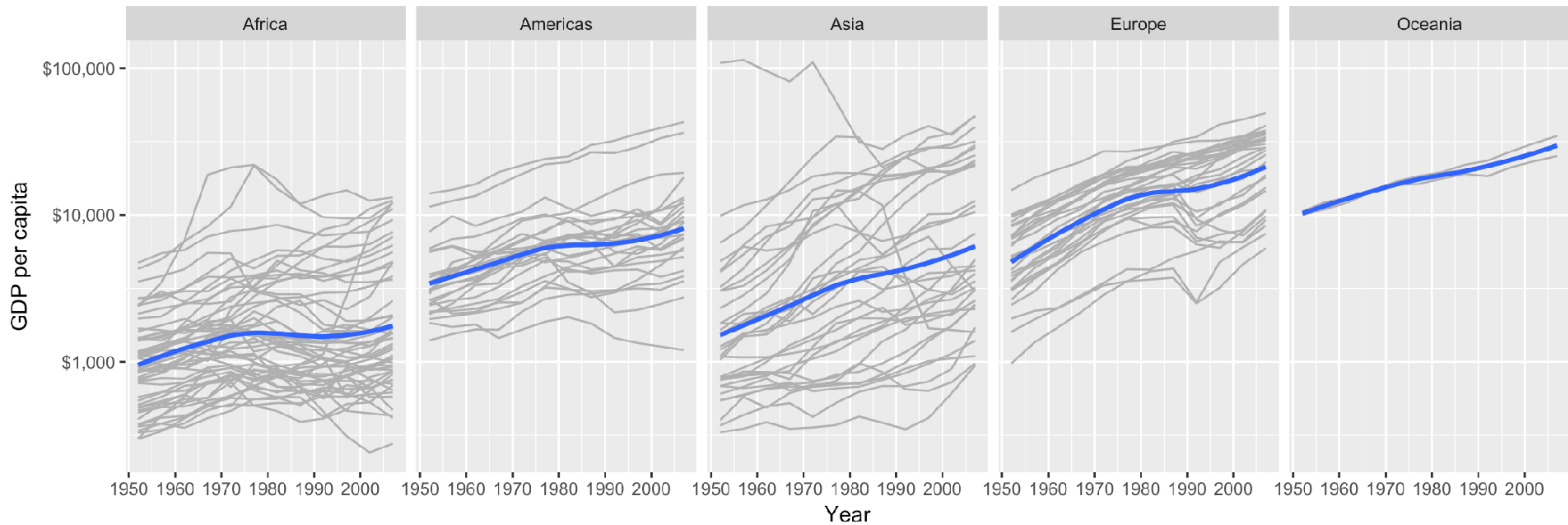
year

```
p + geom_line(color = "gray70",  
             mapping = aes(group = country)) +  
  geom_smooth(size = 1.1,  
             method = "loess",  
             se = FALSE) +  
  scale_y_log10(labels=scales::dollar) +  
  facet_wrap(~ continent, ncol = 5) +  
  labs(x = "Year",  
       y = "GDP per capita",  
       title = "GDP per capita on Five Continents")
```



The `labs()` function  
lets you name labels,  
title, subtitle, etc.

GDP per capita on Five Continents



**geoms CAN  
TRANSFORM  
DATA**



# gss\_sm

## A subset of General Social Survey Questions from 2016

```
> gss_sm
```

```
# A tibble: 2,867 x 32
```

	year	id	ballot	age	childs	sibs	degree	race	sex	region	income16	relig	marital	padeg	madeg
	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<fct>	<fct>	<fct>	<fct>	<fct>	<fct>	<fct>	<fct>	<fct>
1	2016	1	1	47	3	2	Bache...	White	Male	New E...	\$170000...	None	Married	Grad...	High...
2	2016	2	2	61	0	3	High ...	White	Male	New E...	\$50000 ...	None	Never ...	Lt H...	High...
3	2016	3	3	72	2	3	Bache...	White	Male	New E...	\$75000 ...	Cath...	Married	High...	Lt H...
4	2016	4	1	43	4	3	High ...	White	Fema...	New E...	\$170000...	Cath...	Married	NA	High...
5	2016	5	3	55	2	2	Gradu...	White	Fema...	New E...	\$170000...	None	Married	Bach...	High...
6	2016	6	2	53	2	2	Junio...	White	Fema...	New E...	\$60000 ...	None	Married	NA	High...
7	2016	7	1	50	2	2	High ...	White	Male	New E...	\$170000...	None	Married	High...	High...
8	2016	8	3	23	3	6	High ...	Other	Fema...	Middl...	\$30000 ...	Cath...	Married	Lt H...	Lt H...
9	2016	9	1	45	3	5	High ...	Black	Male	Middl...	\$60000 ...	Prot...	Married	Lt H...	Lt H...
10	2016	10	3	71	4	1	Junio...	White	Male	Middl...	\$60000 ...	None	Divorc...	High...	High...

```
# ... with 2,857 more rows, and 17 more variables: partyid <fct>, polviews <fct>, happy <fct>,  
# partners <fct>, grass <fct>, zodiac <fct>, pres12 <dbl>, wtssall <dbl>, income_rc <fct>, agegrp <fct>,  
# ageq <fct>, siblings <fct>, kids <fct>, religion <fct>, bigregion <fct>, partners_rc <fct>,  
# obama <dbl>
```

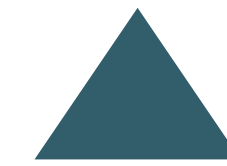
```
> |
```

```
with(gss_sm, table(religion))
```

```
##
```

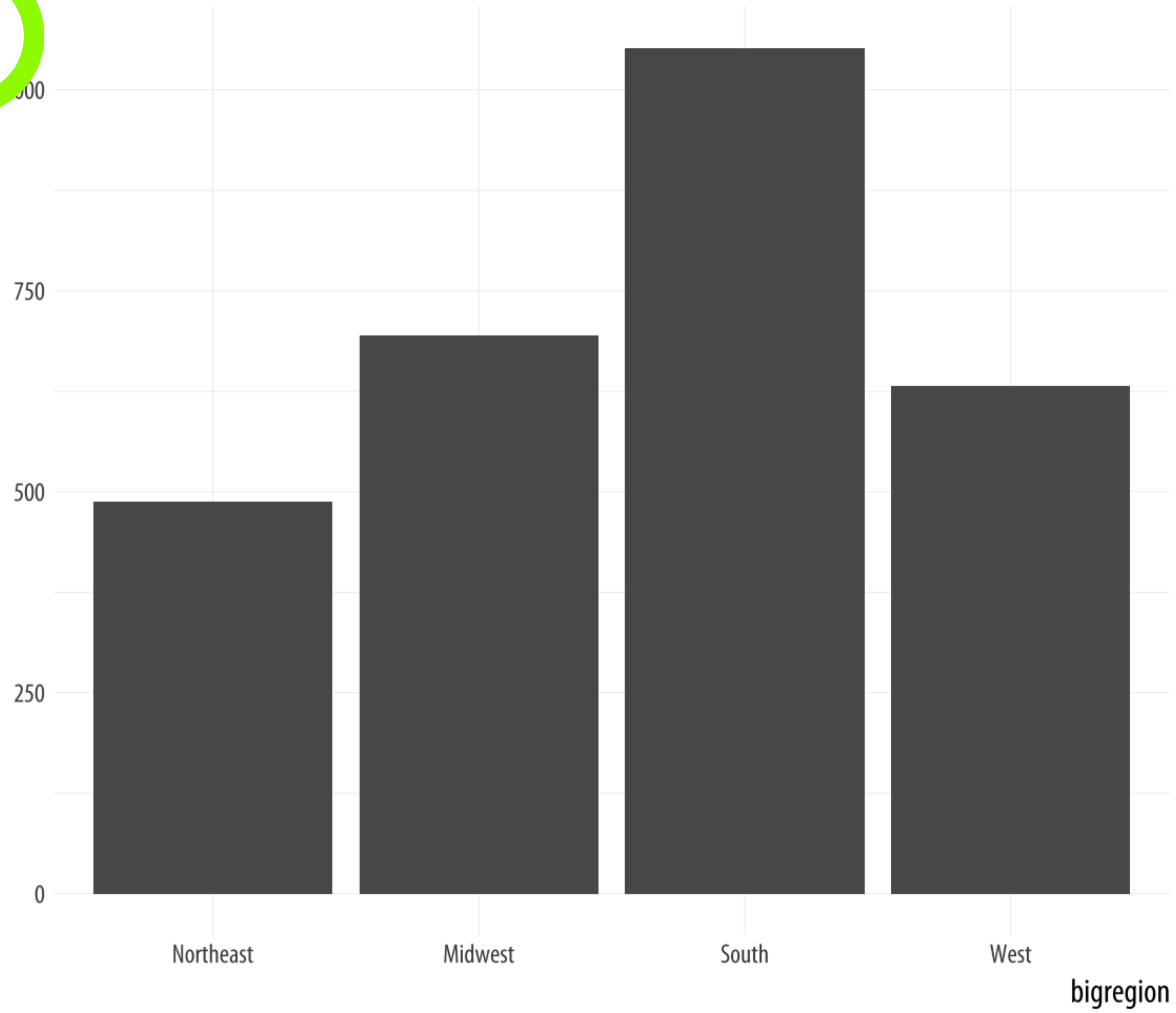
## Protestant	Catholic	Jewish	None	Other
## 1371	649	51	619	159

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion))  
p + geom_bar()
```



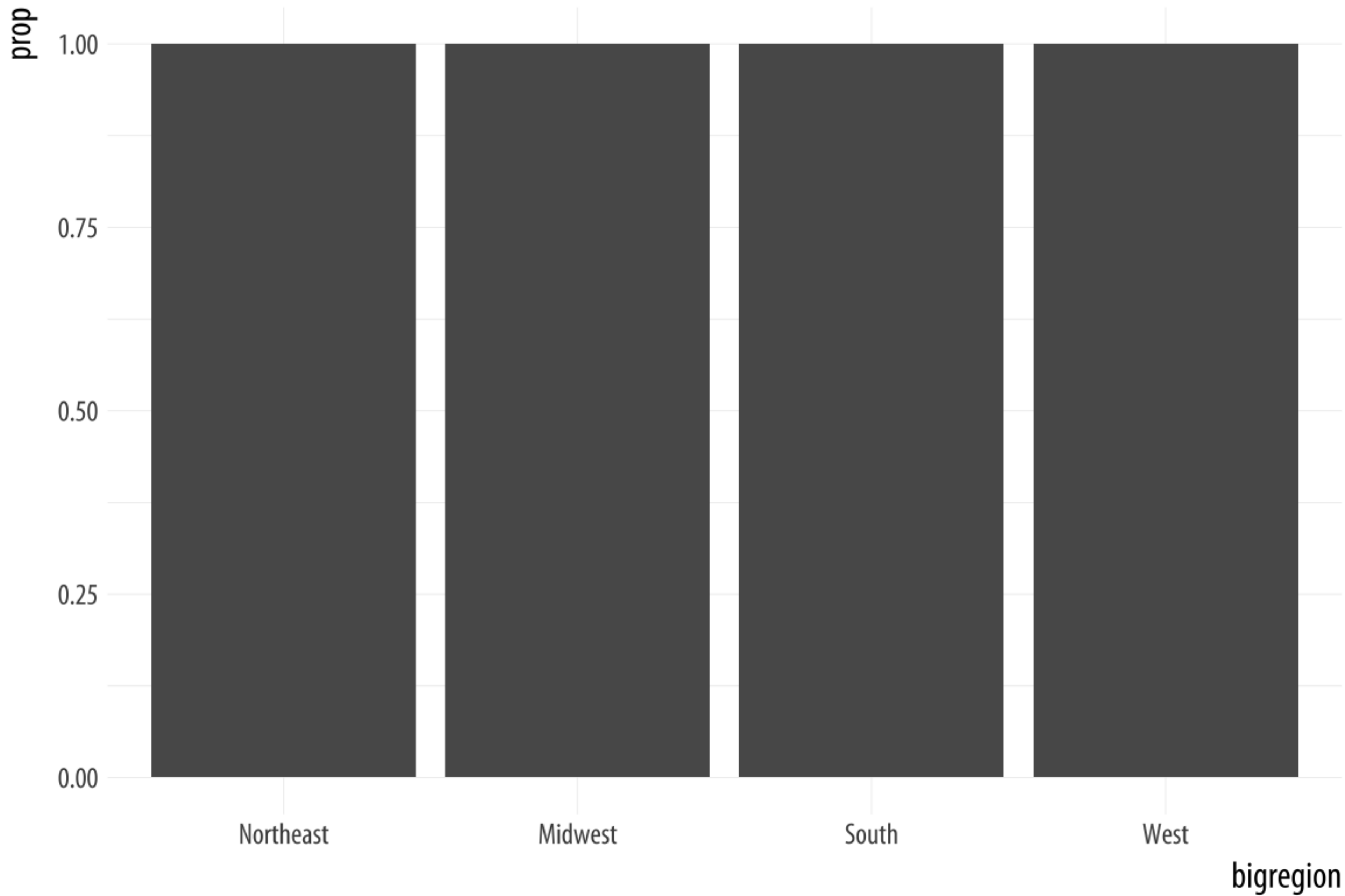
**Just the one aesthetic  
mapping, to x.**

count



The y-axis variable, `count`, is not in the data. Instead, ggplot has calculated it for us. It does this using the default `stat_` function associated with `geom_bar()`, `stat_count()`. This function can compute two new variables, `count`, and `prop` (short for **proportion**). The `count` statistic is the default one used.

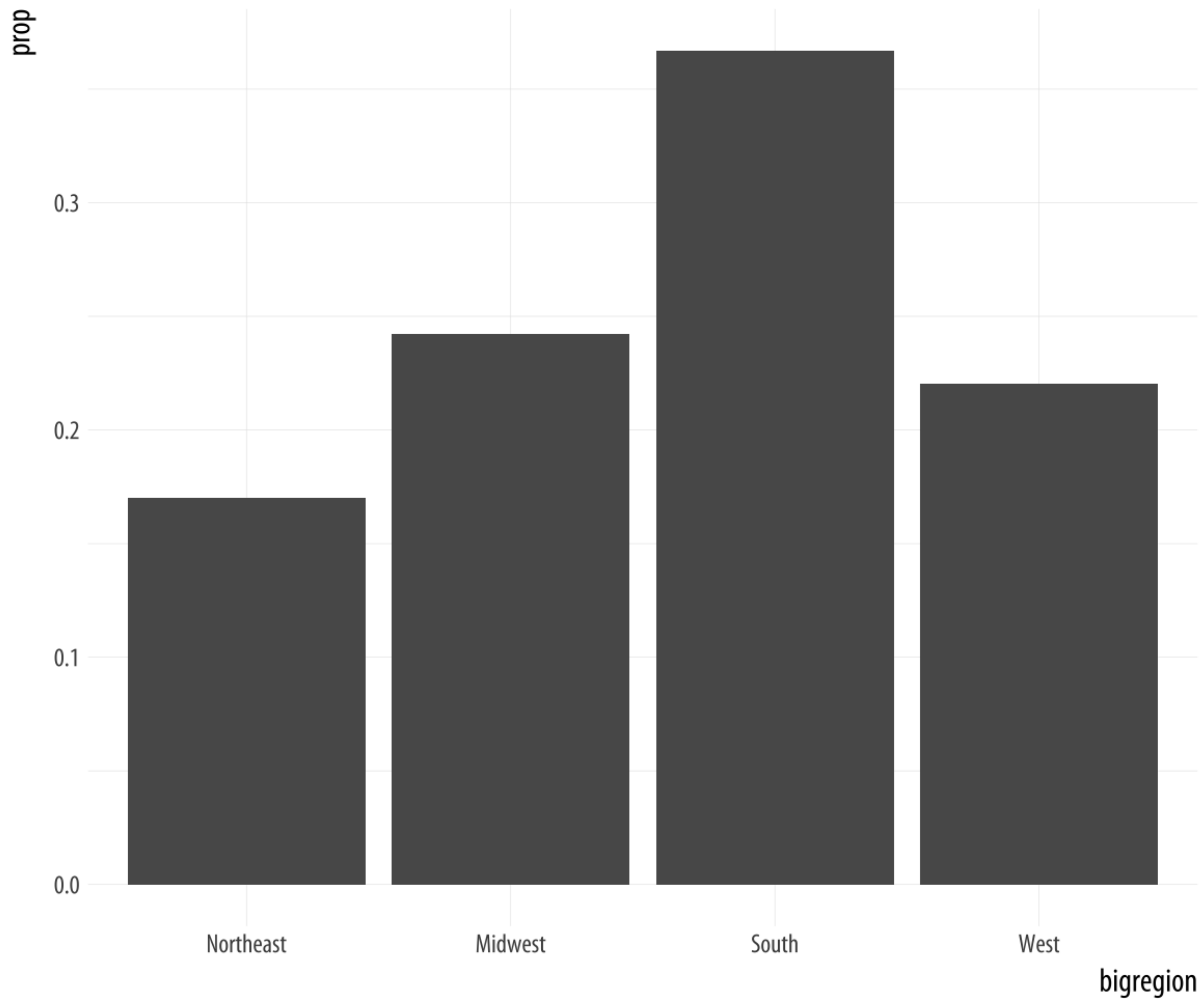
```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion))  
p + geom_bar(mapping = aes(y = ..prop..))
```





```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion))  
p + geom_bar(mapping = aes(y = ..prop.., group = 1))
```





```
p + geom_bar()
```

```
p + stat_count()
```

geom\_ functions call  
their default stat\_ functions  
behind the scenes. (And vice versa)

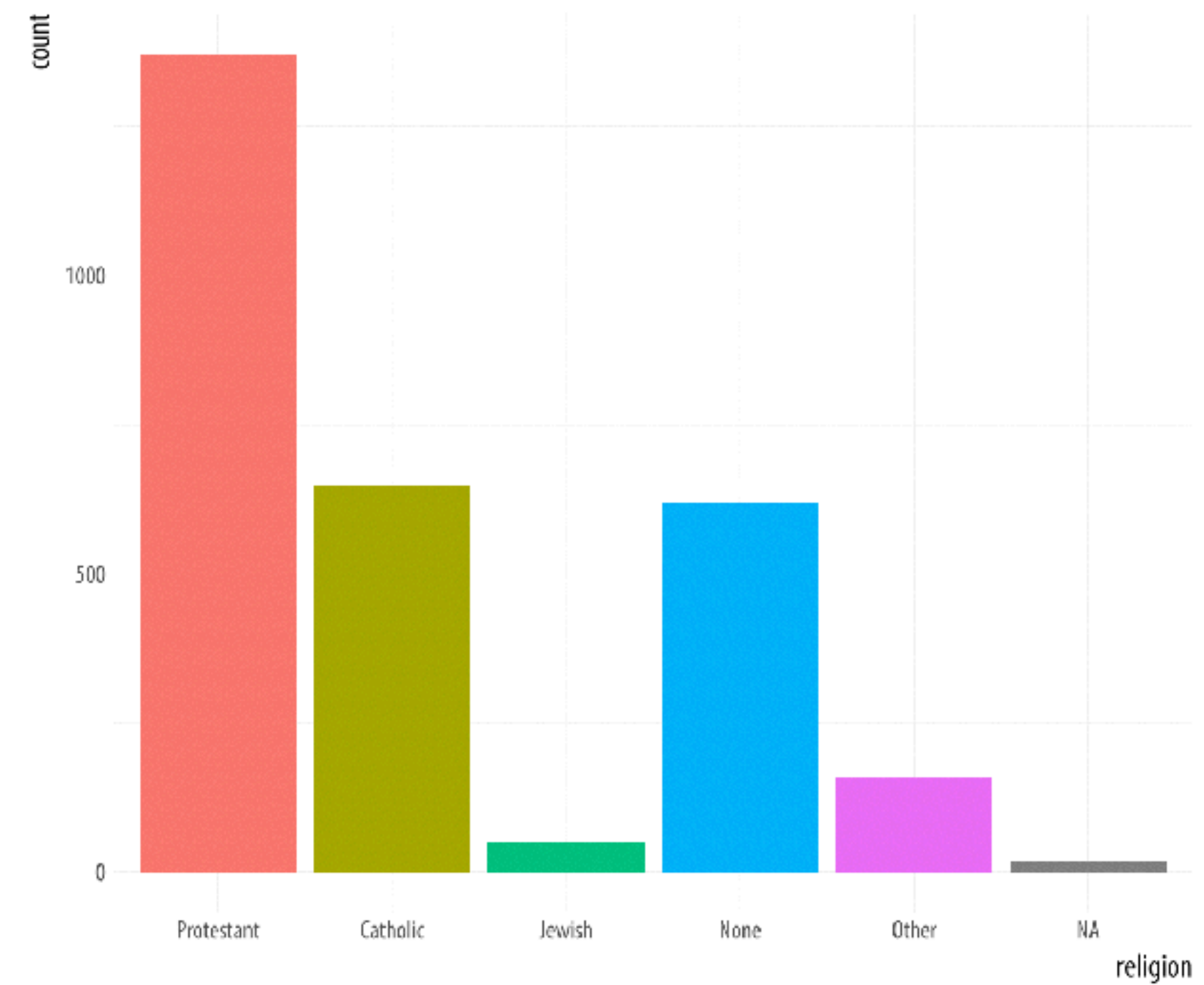
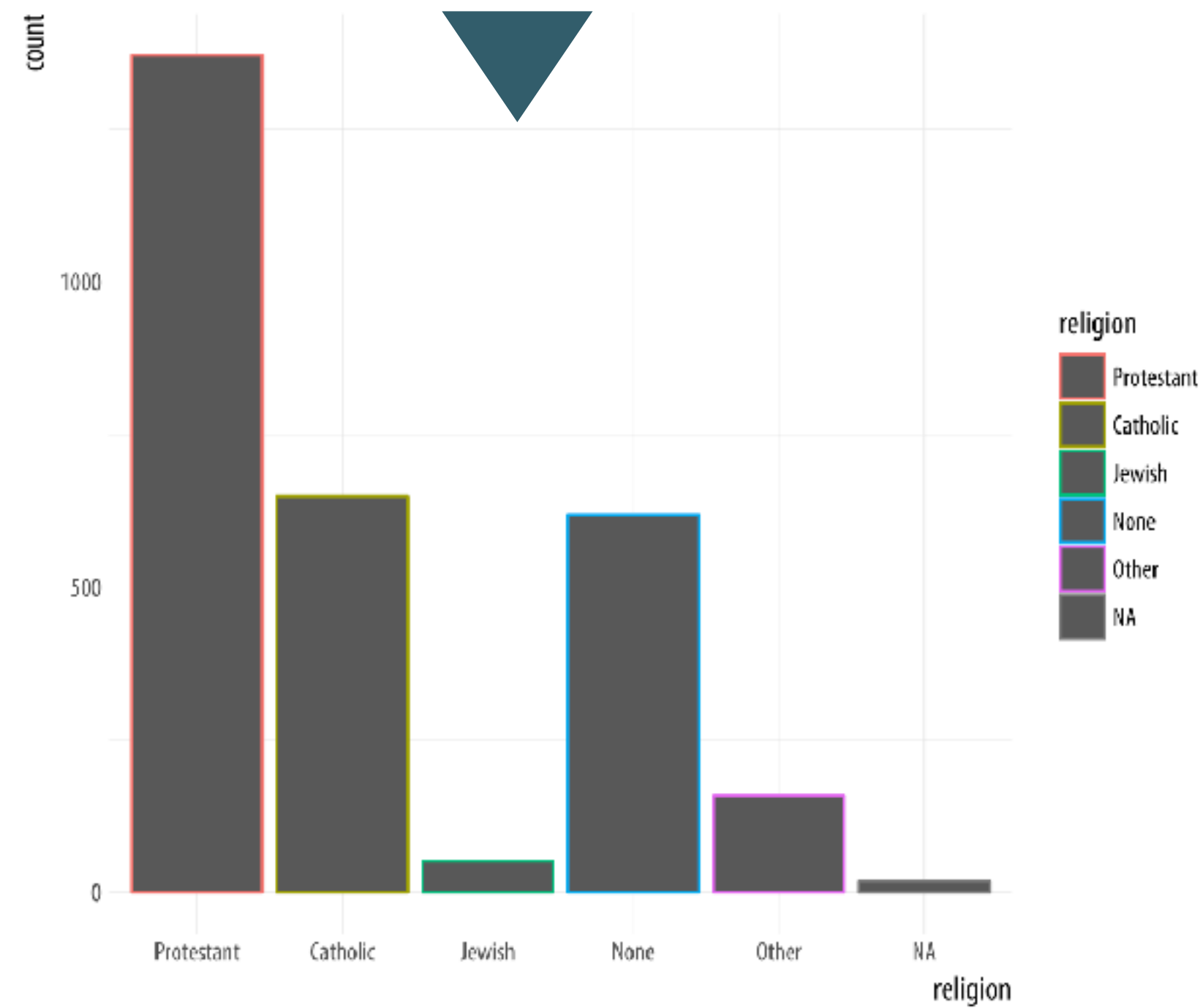
```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = religion))  
p + geom_bar()
```

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = religion, color = religion))  
p + geom_bar()
```

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = religion, fill = religion))  
p + geom_bar()
```

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = religion, fill = religion))  
p + geom_bar() + guides(fill = FALSE)
```

```
p <- ggplot(data = gss_sm,
            mapping = aes(x = religion, color = religion))
p + geom_bar()
```



```
p <- ggplot(data = gss_sm,
            mapping = aes(x = religion, fill = religion))
p + geom_bar() + guides(fill = FALSE)
```

# HISTOGRAMS & KERNEL DENSITIES

# midwest

## County-Level Census Data for Midwestern States

```
> midwest
# A tibble: 437 x 28
  PID county state area poptotal popdensity popwhite popblack popamerindian popasian popother percwhite
  <int> <chr> <chr> <dbl> <int> <dbl> <int> <int> <int> <int> <int> <dbl>
1  561 ADAMS IL 0.052 66090 1271. 63917 1702 98 249 124 96.7
2  562 ALEXA... IL 0.014 10626 759 7054 3496 19 48 9 66.4
3  563 BOND IL 0.022 14991 681. 14477 429 35 16 34 96.6
4  564 BOONE IL 0.017 30806 1812. 29344 127 46 150 1139 95.3
5  565 BROWN IL 0.018 5836 324. 5264 547 14 5 6 90.2
6  566 BUREAU IL 0.05 35688 714. 35157 50 65 195 221 98.5
7  567 CALHO... IL 0.017 5322 313. 5298 1 8 15 0 99.5
8  568 CARRO... IL 0.027 16805 622. 16519 111 30 61 84 98.3
9  569 CASS IL 0.024 13437 560. 13384 16 8 23 6 99.6
10 570 CHAMP... IL 0.058 173025 2983. 146506 16559 331 8033 1596 84.7
# ... with 427 more rows, and 16 more variables: percblack <dbl>, percamerindan <dbl>, percasian <dbl>,
# percother <dbl>, popadults <int>, perchsd <dbl>, percollege <dbl>, percprof <dbl>,
# poppovertyknown <int>, percpovertyknown <dbl>, percbelowpoverty <dbl>, percchildbelowpovert <dbl>,
# percadultpoverty <dbl>, percelderlypoverty <dbl>, inmetro <int>, category <chr>
> |
```

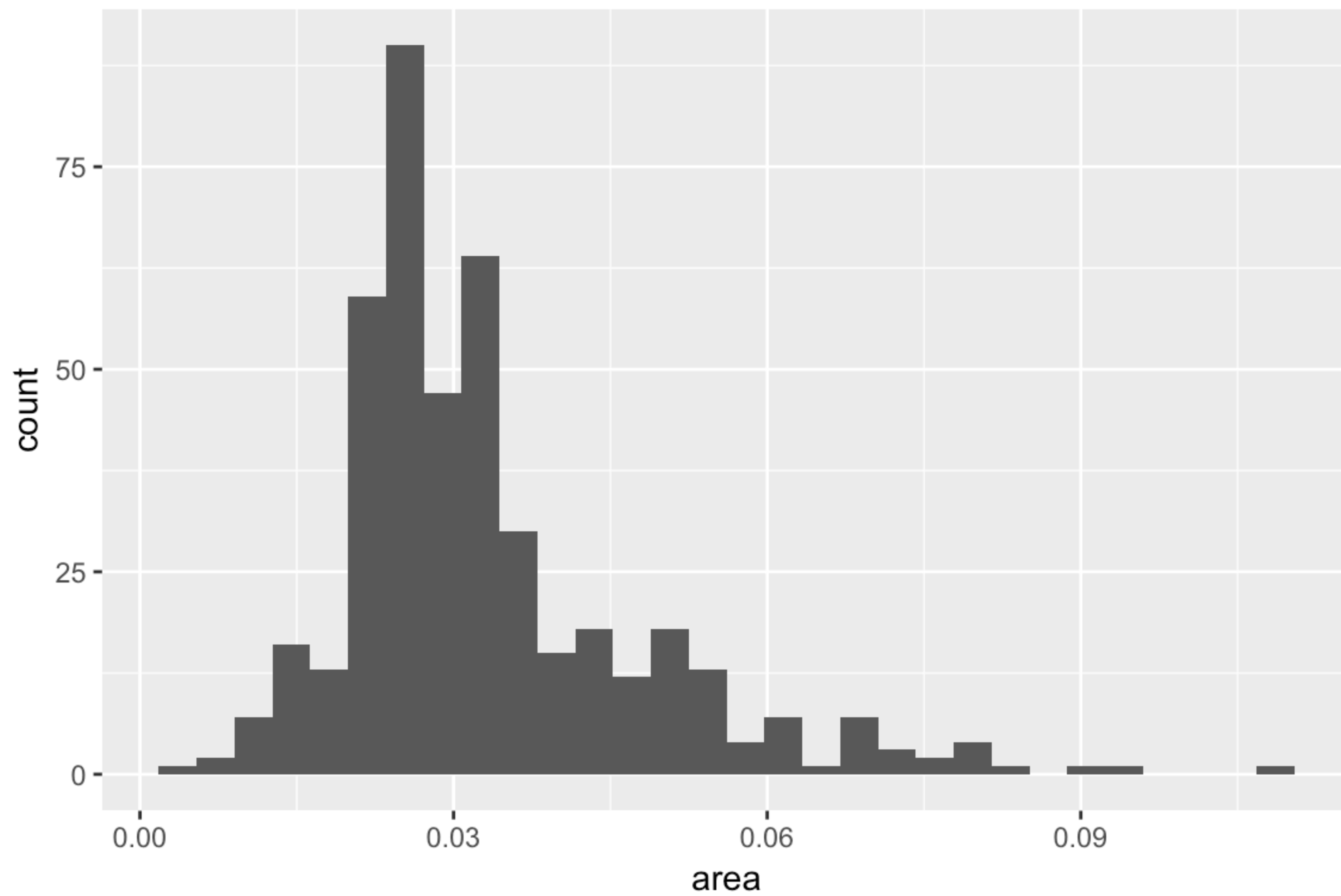


```
p <- ggplot(data = midwest,  
            mapping = aes(x = area))  
p + geom_histogram()
```

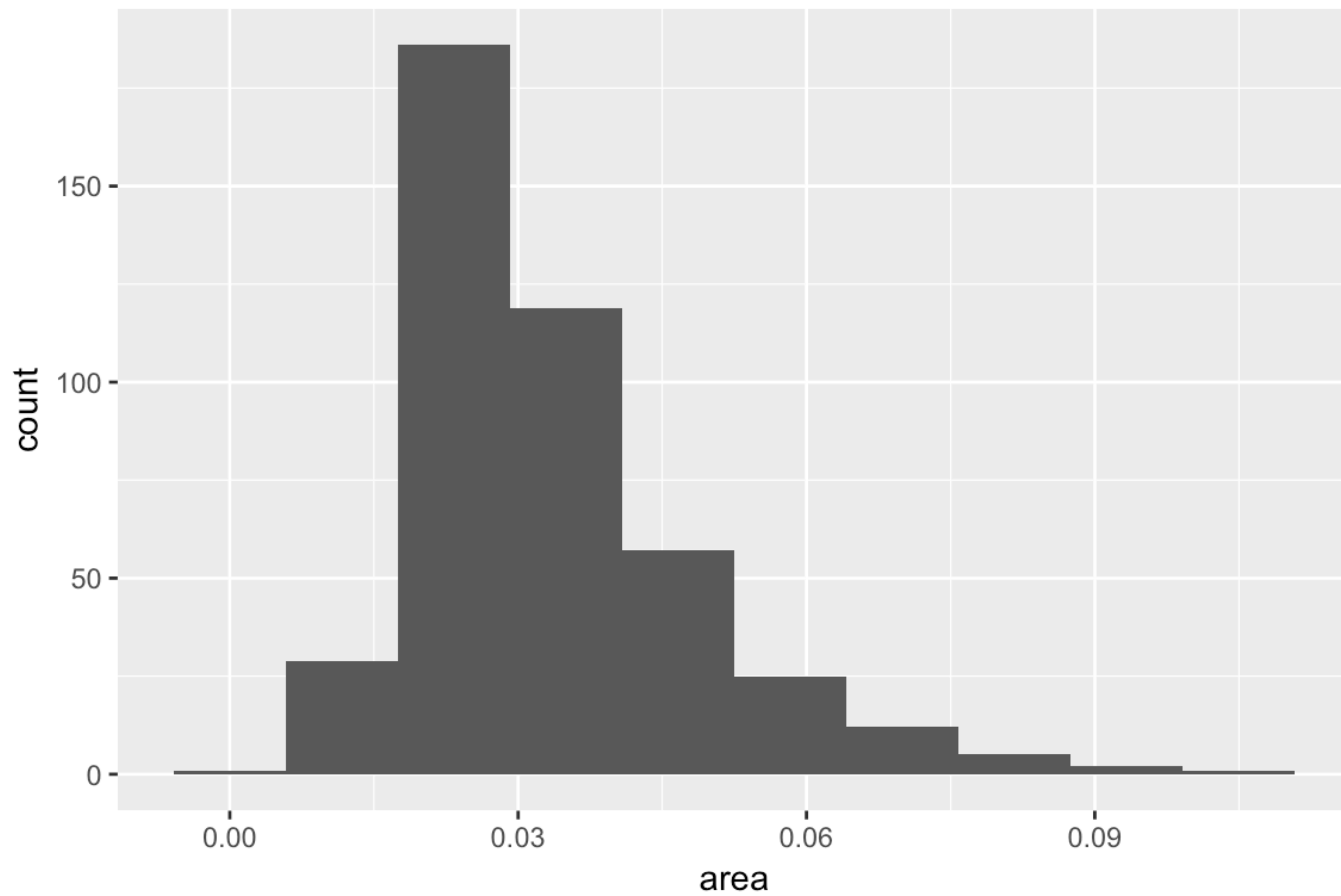
```
## `stat_bin()` using `bins = 30`.  
## Pick better value with `binwidth`.
```



The default stat for  
this geom has to make  
a choice, and is letting  
us know we might  
want to override it.



```
p <- ggplot(data = midwest,  
            mapping = aes(x = area))  
p + geom_histogram(bins = 10)
```



```
oh_wi <- c("OH", "WI")
```

subset our data  
on the fly ▼

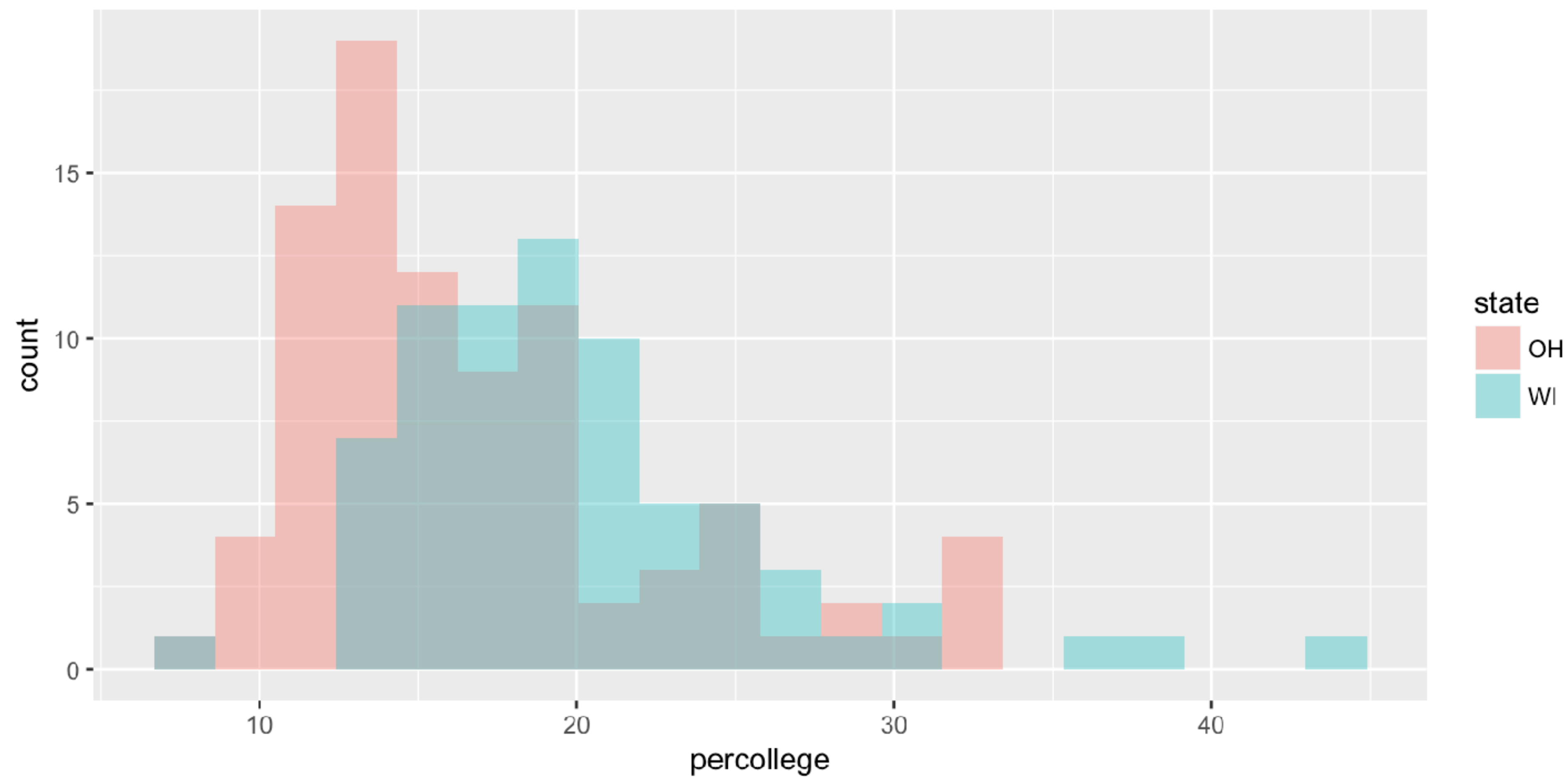
a convenient,  
built-in operator ▼

```
p <- ggplot(data = subset(midwest, state %in% oh_wi),  
            mapping = aes(x = percollege, fill = state))
```

```
p + geom_histogram(position = "identity",  
                   alpha = 0.4, bins = 20)
```



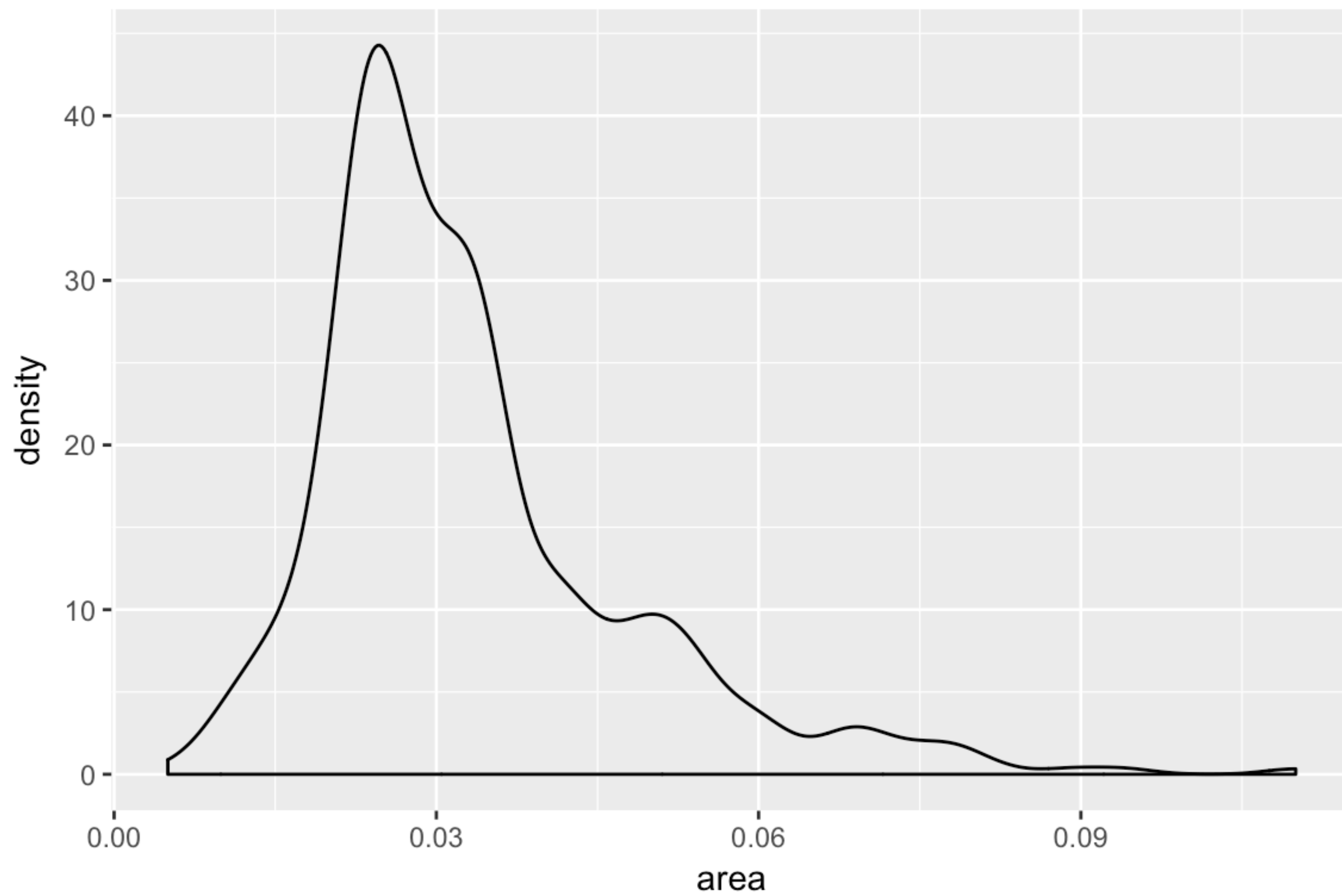
Just plot x by its  
values on the  
scale, don't stack  
or dodge



```
p <- ggplot(data = midwest,  
            mapping = aes(x = area))  
p + geom_density()
```

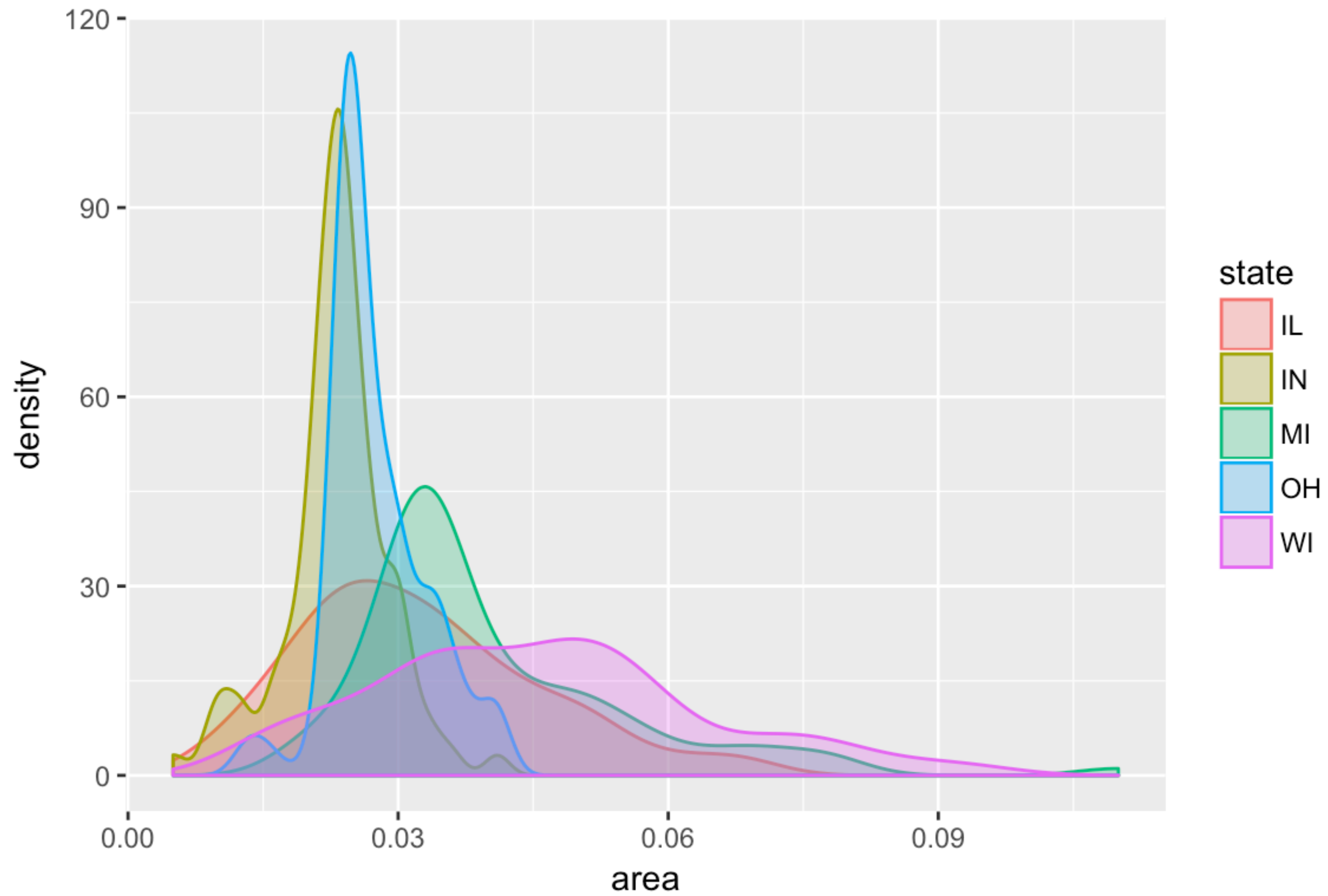
geom\_hist()'s continuous  
counterpart, geom\_density()





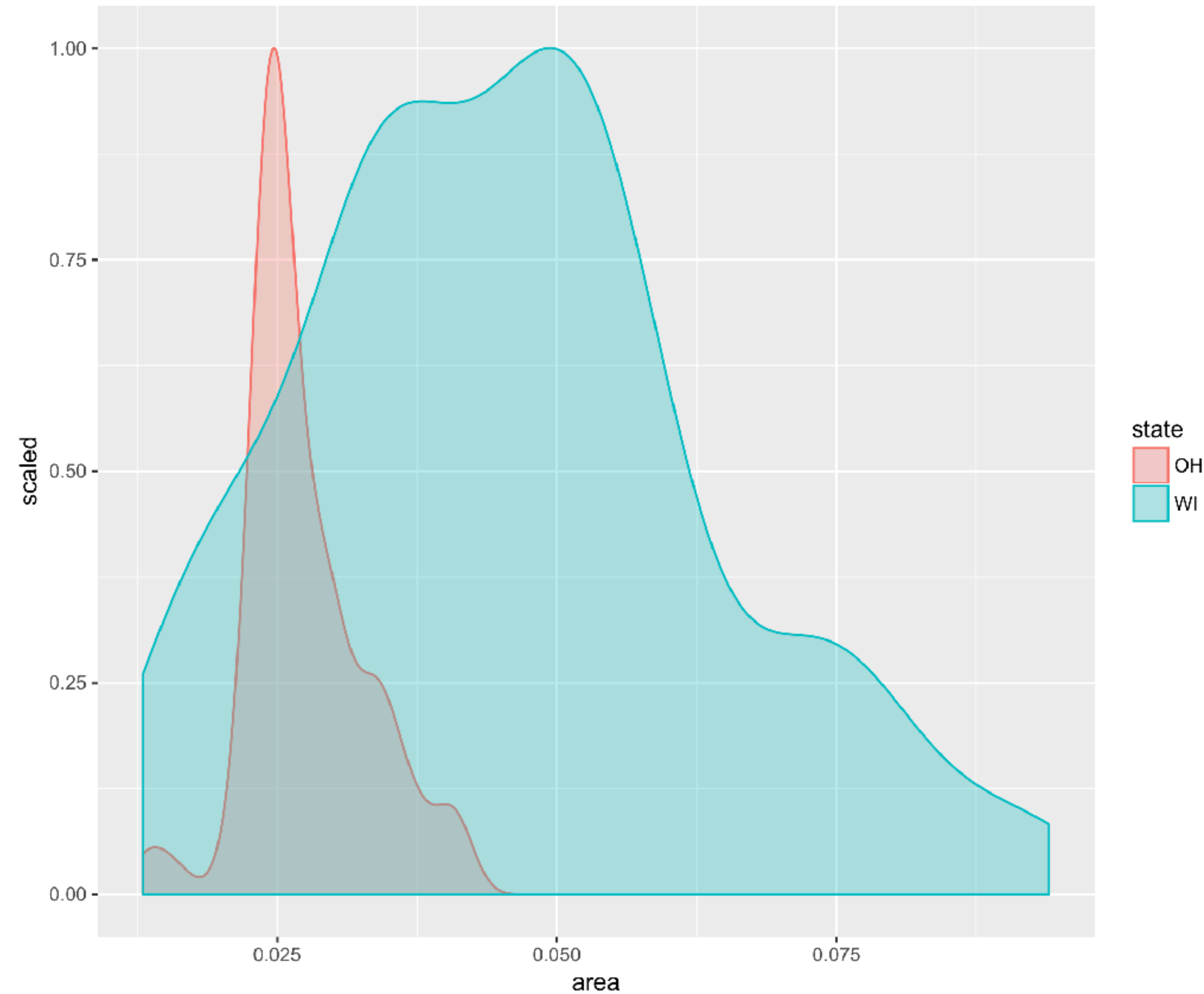
```
p <- ggplot(data = midwest,  
            mapping = aes(x = area,  
                          fill = state,  
                          color = state))
```

```
p + geom_density(alpha = 0.3)
```



```
p <- ggplot(data = subset(midwest, subset = state %in% OH_WI),  
            mapping = aes(x = area, fill = state, color = state))
```

```
p + geom_density(alpha = 0.3, mapping = (aes(y = ..scaled..)))
```



**AVOIDING  
TRANSFORMATIONS  
WHEN NECESSARY**

```
> titanic
```

```
##           fate gender      n percent
## 1 perished    male 1364      62.0
## 2 perished    female 126       5.7
## 3 survived    male  367      16.7
## 4 survived    female 344      15.6
```

**No counting up required?**  
**Then stat = identity**

```
p <- ggplot(data = titanic,  
            mapping = aes(x = fate,  
                           y = percent,  
                           fill = sex))  
p + geom_bar(stat = "identity",  
             position = "dodge") +  
  theme(legend.position = "top")
```

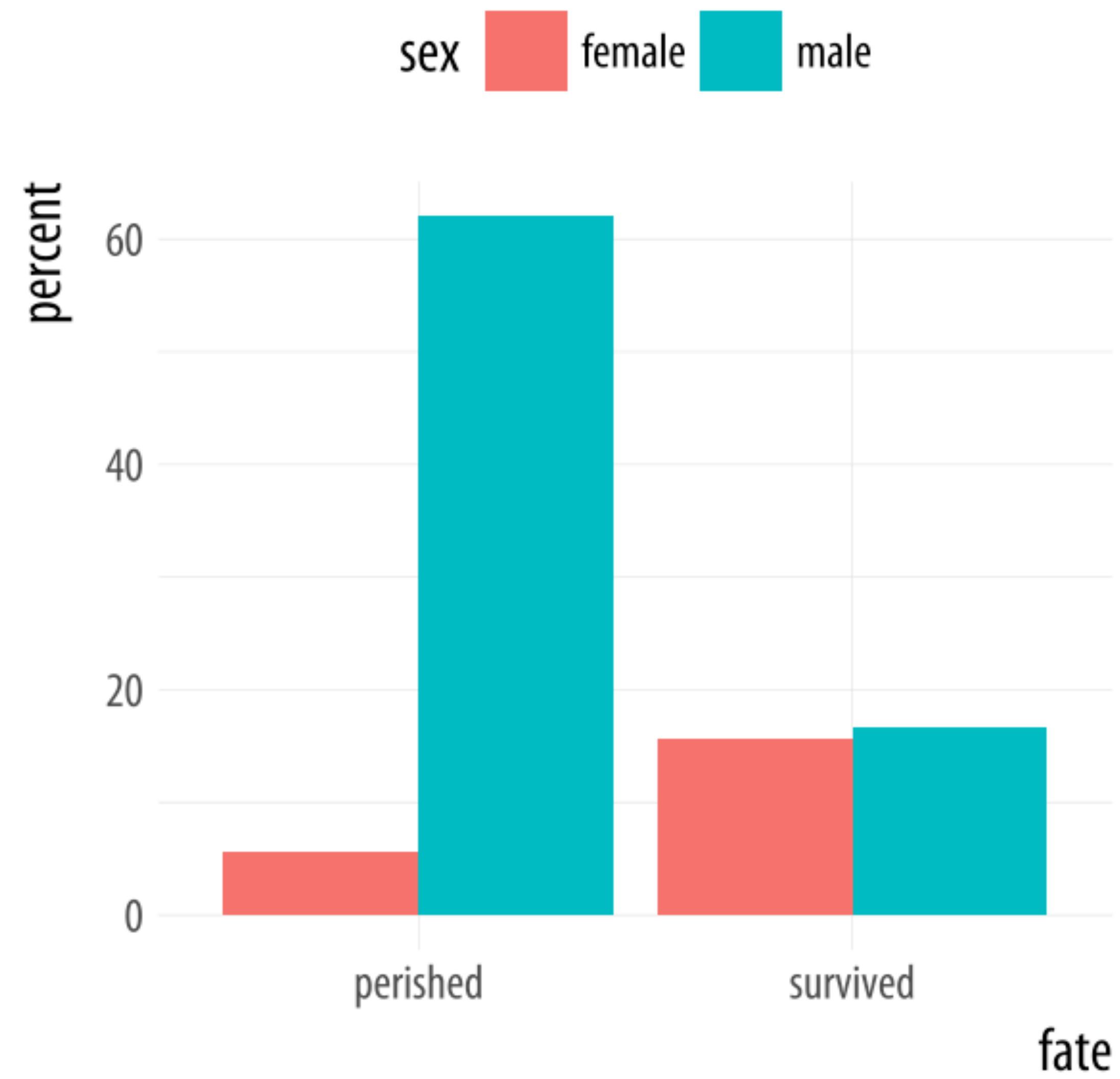


The `theme()` function  
controls parts of the  
plot that don't belong  
to its “grammatical”  
structure

```
p <- ggplot(data = titanic,  
            mapping = aes(x = fate,  
                          y = percent,  
                          fill = sex))  
  
p + geom_col(position = "dodge") +  
  theme(legend.position = "top")
```

Even better: for convenience when  
not counting up, just use `geom_col()`





```
oecd_sum
```

```
## # A tibble: 57 x 5
```

```
## # Groups:   year [57]
```

```
##      year other   usa  diff hi_lo
```

```
##      <int> <dbl> <dbl> <dbl> <chr>
```

```
##    1  1960  68.6  69.9  1.30  Below
```

```
##    2  1961  69.2  70.4  1.20  Below
```

```
##    3  1962  68.9  70.2  1.30  Below
```

```
##    4  1963  69.1  70.0  0.900  Below
```

```
##    5  1964  69.5  70.3  0.800  Below
```

```
##    6  1965  69.6  70.3  0.700  Below
```

```
##    7  1966  69.9  70.3  0.400  Below
```

```
##    8  1967  70.1  70.7  0.600  Below
```

```
##    9  1968  70.1  70.4  0.300  Below
```

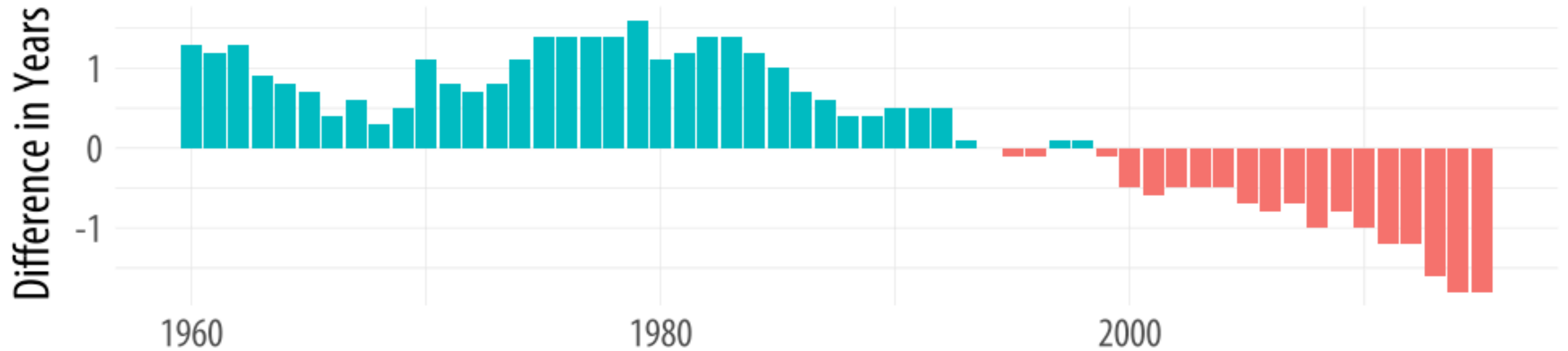
```
##   10  1969  70.1  70.6  0.500  Below
```

```
## # ... with 47 more rows
```



# The US Life Expectancy Gap

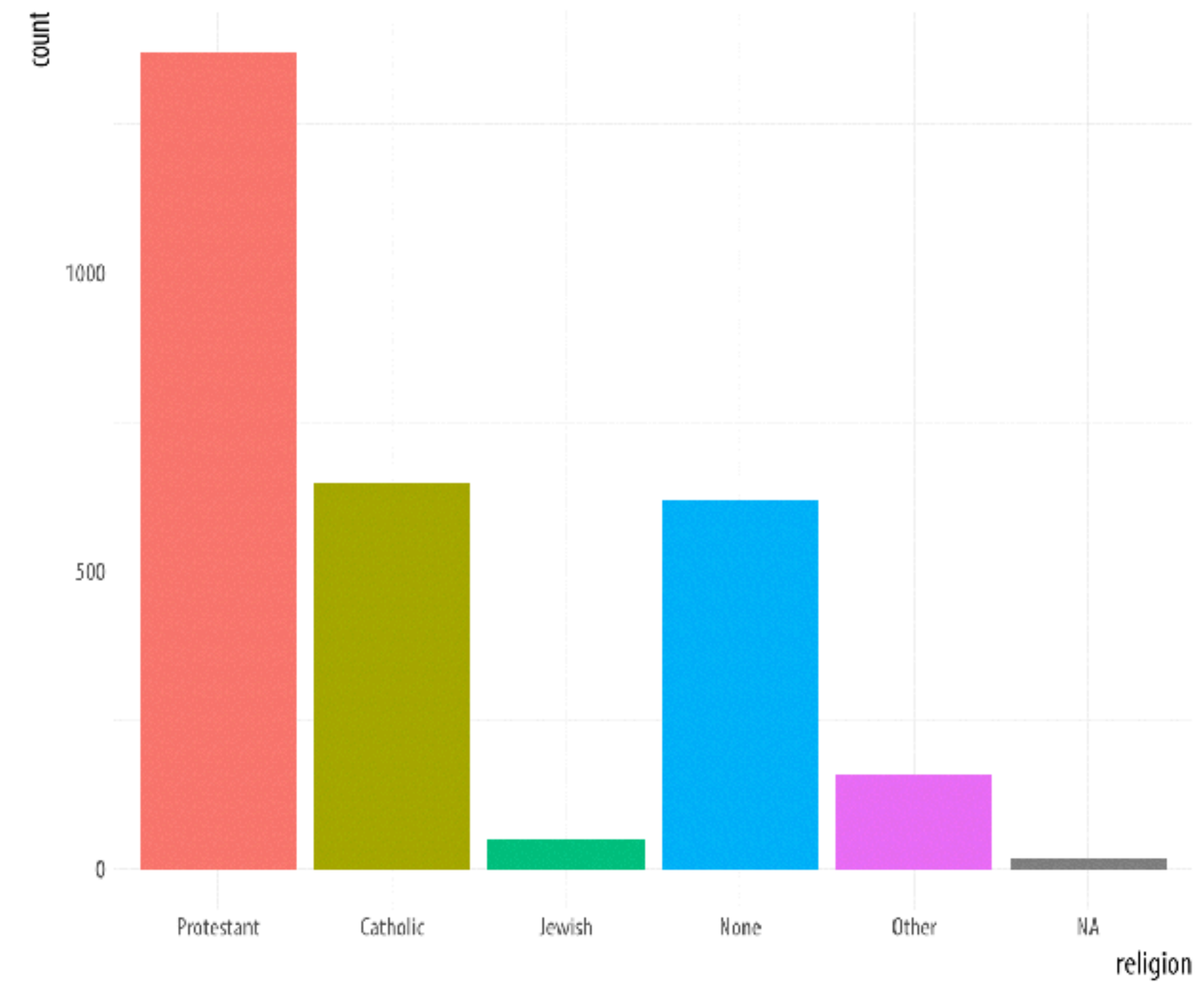
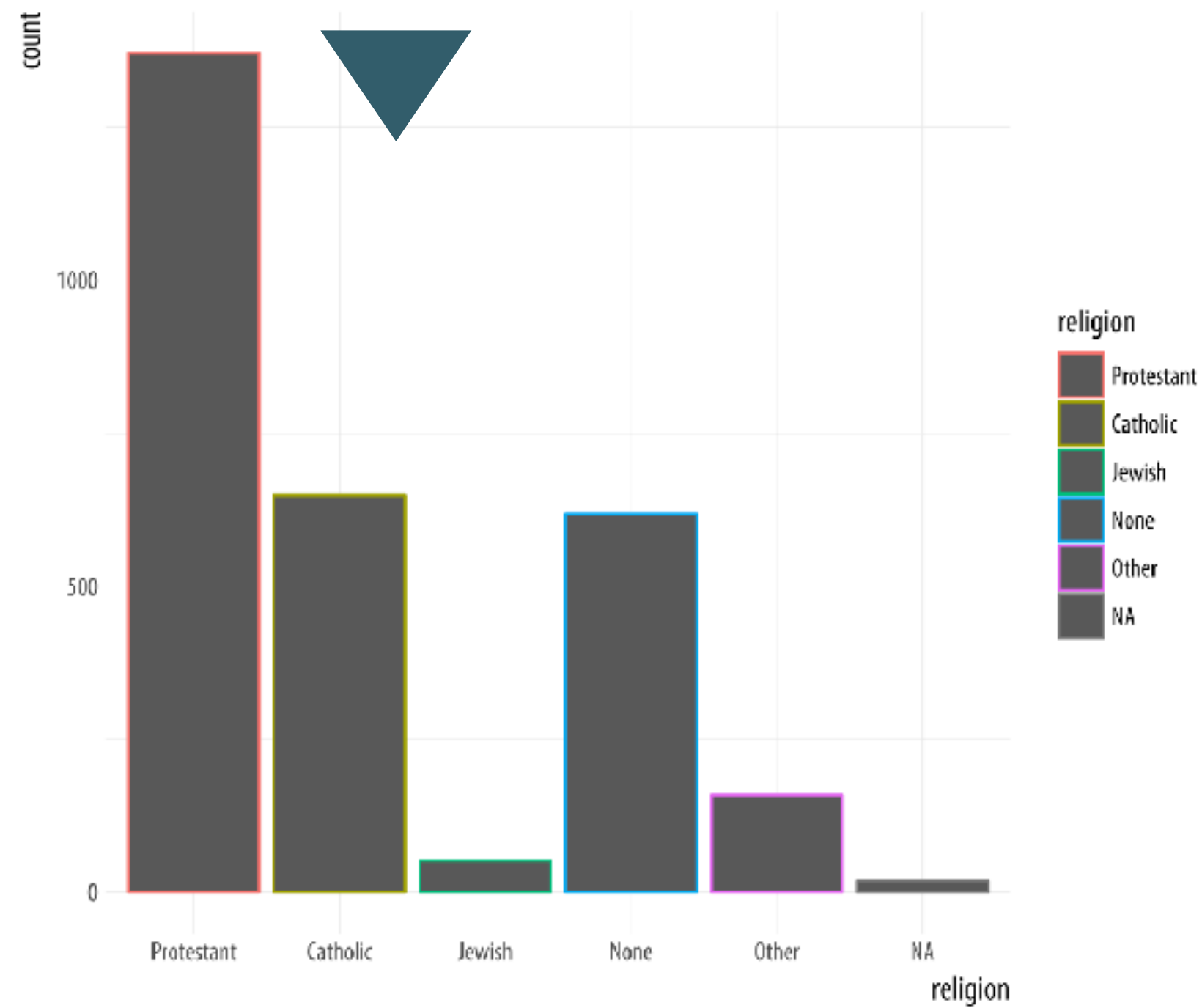
Difference between US and OECD average life expectancies, 1960-2015



Data: OECD. After a chart by Christopher Ingraham,  
Washington Post, December 27th 2017.

CROSSTABULATION  
THE **AWKWARD** WAY

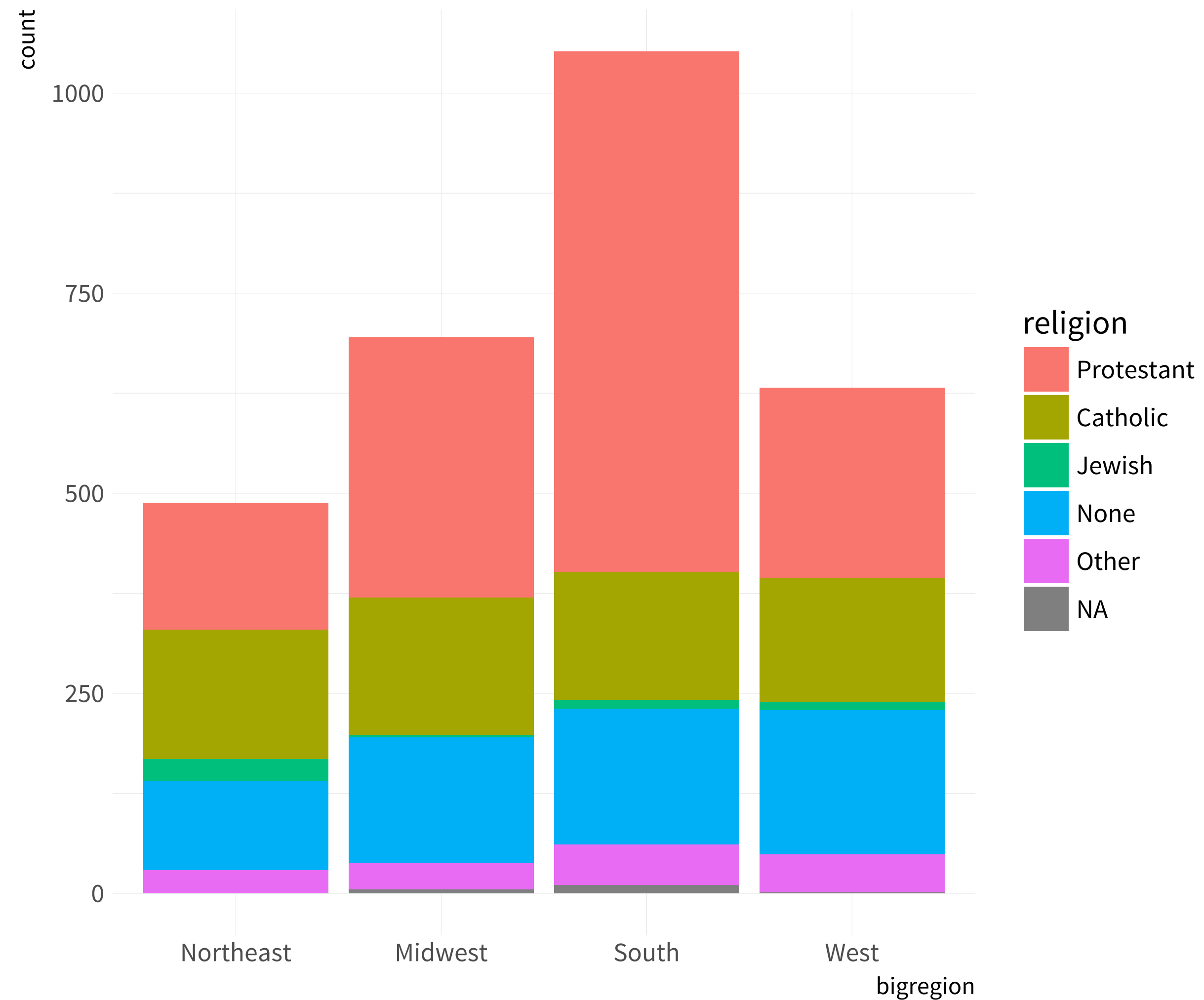
```
p <- ggplot(data = gss_sm,
            mapping = aes(x = religion, color = religion))
p + geom_bar()
```

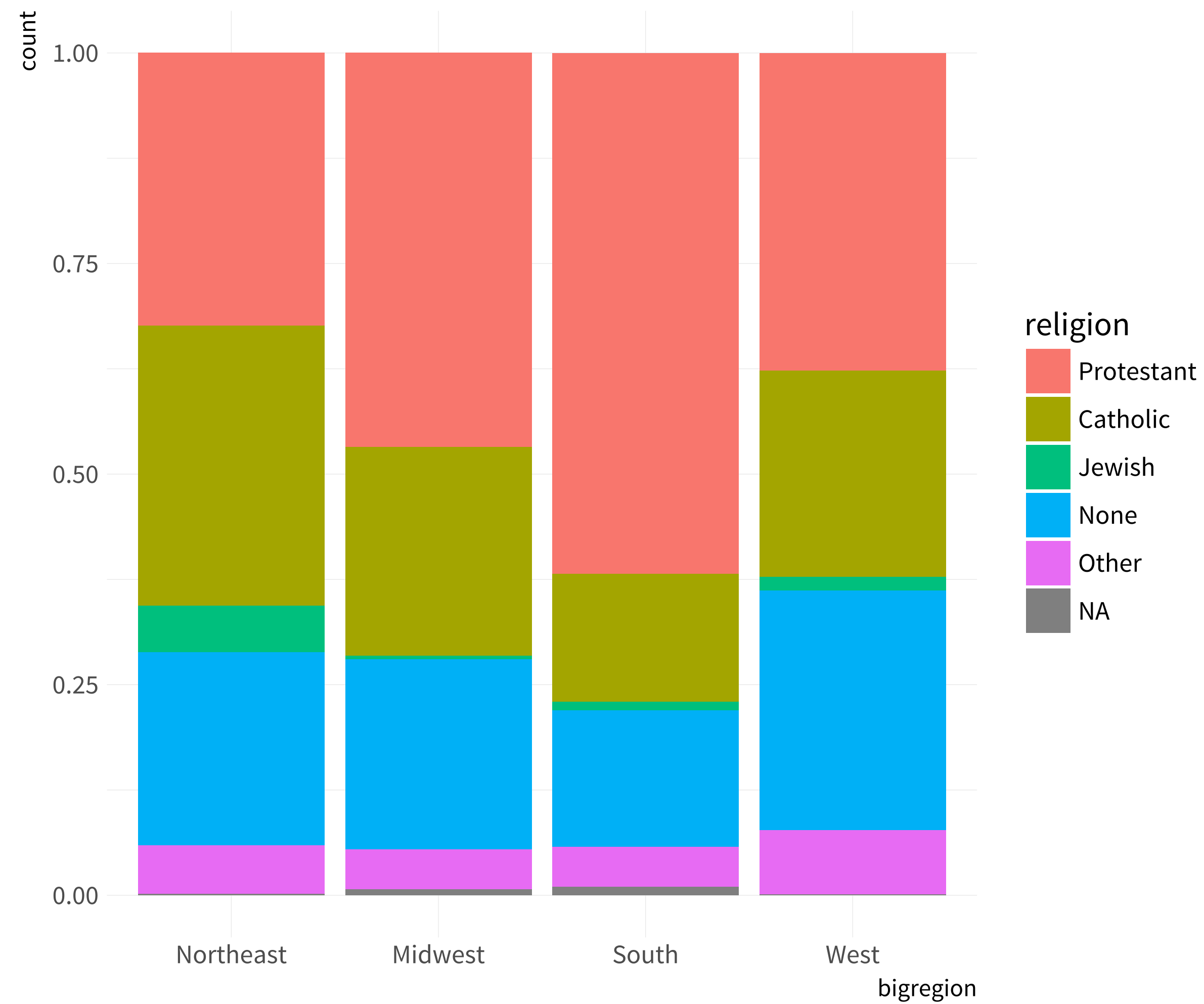


```
p <- ggplot(data = gss_sm,
            mapping = aes(x = religion, fill = religion))
p + geom_bar() + guides(fill = FALSE)
```

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion,  
                          fill = religion))
```

```
p + geom_bar()
```

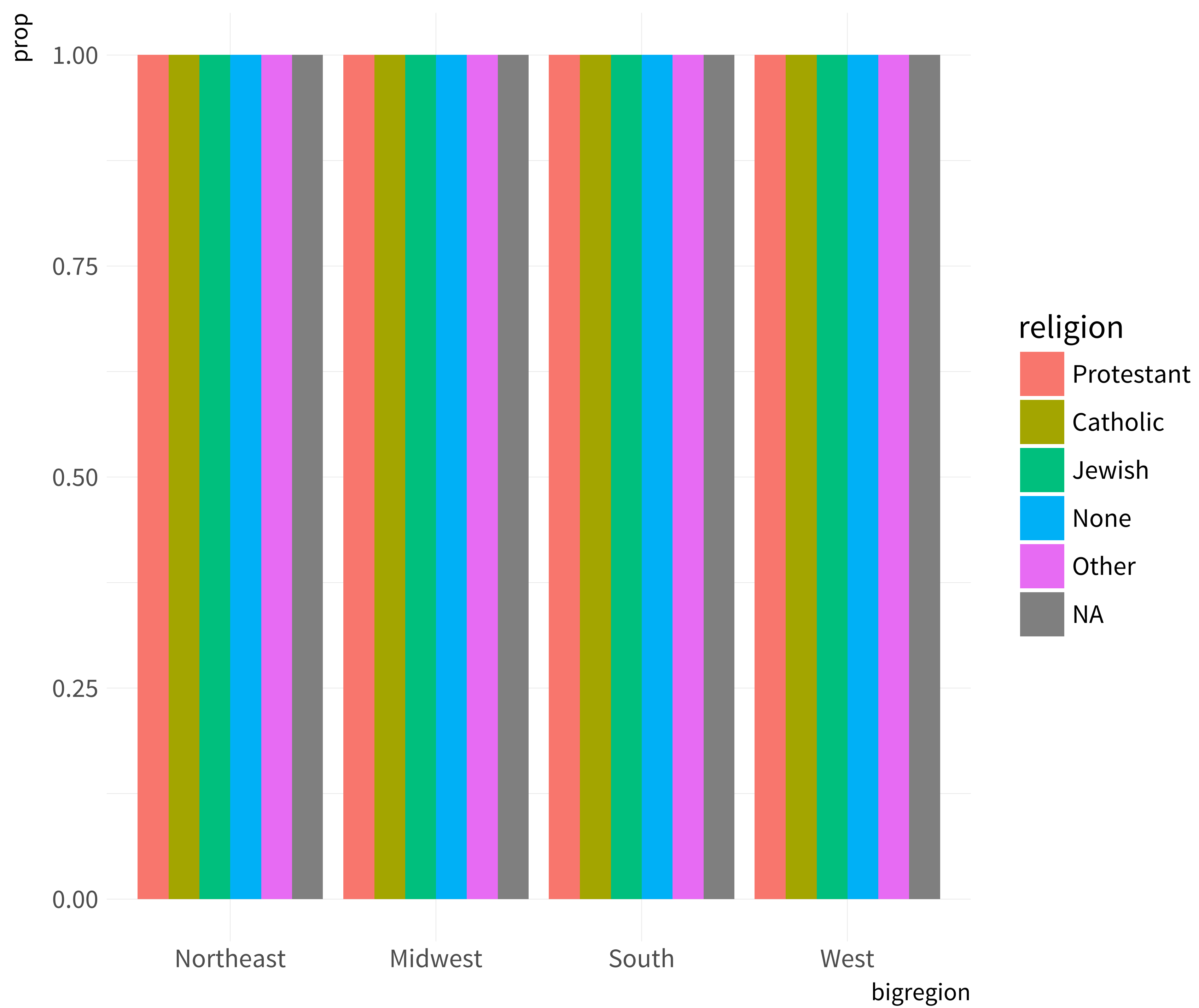




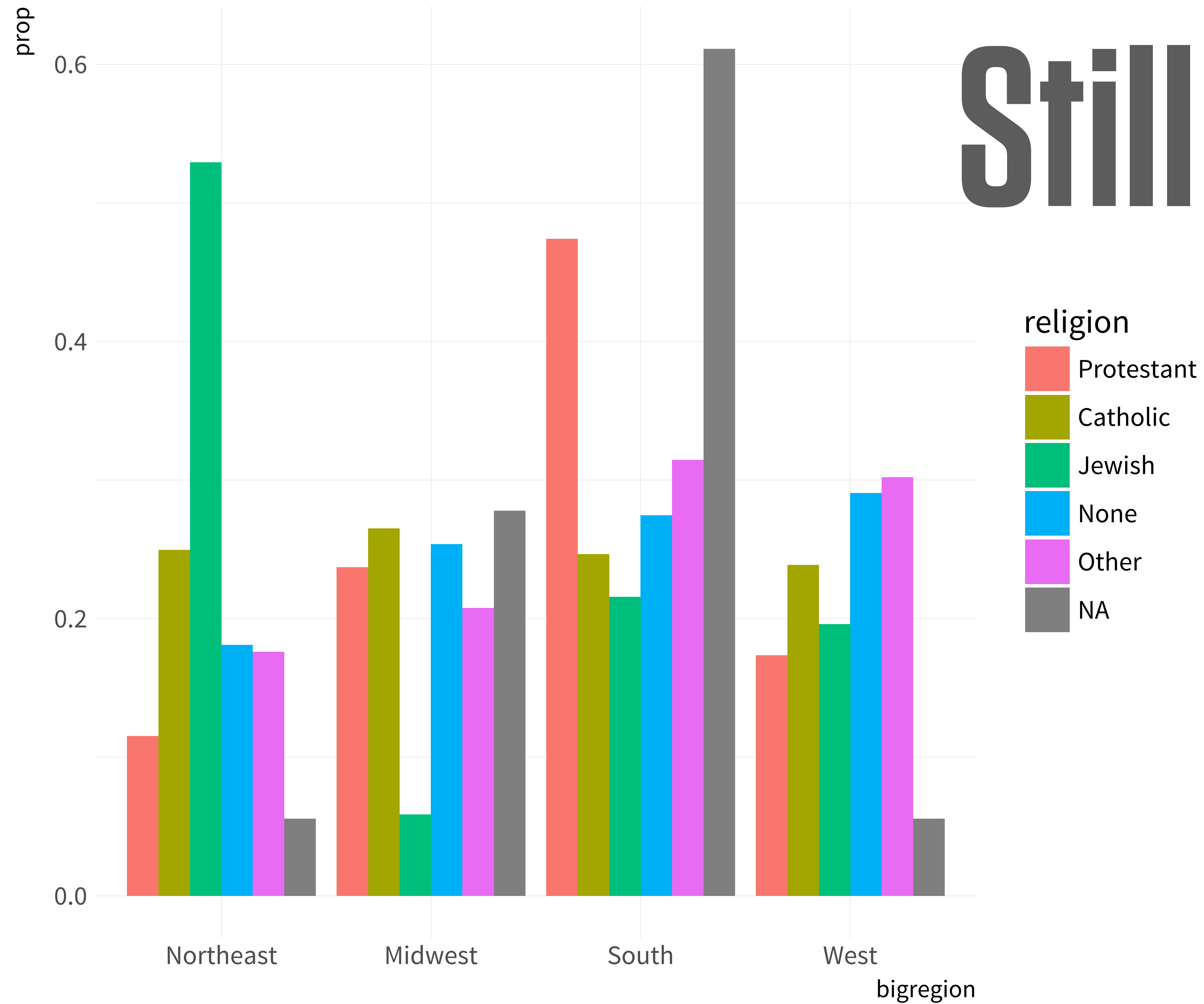
```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion,  
                          fill = religion))  
  
p + geom_bar(position = "fill")
```



```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion,  
                           fill = religion))  
p + geom_bar(position = "dodge",  
             mapping = aes(y = ..prop..))
```



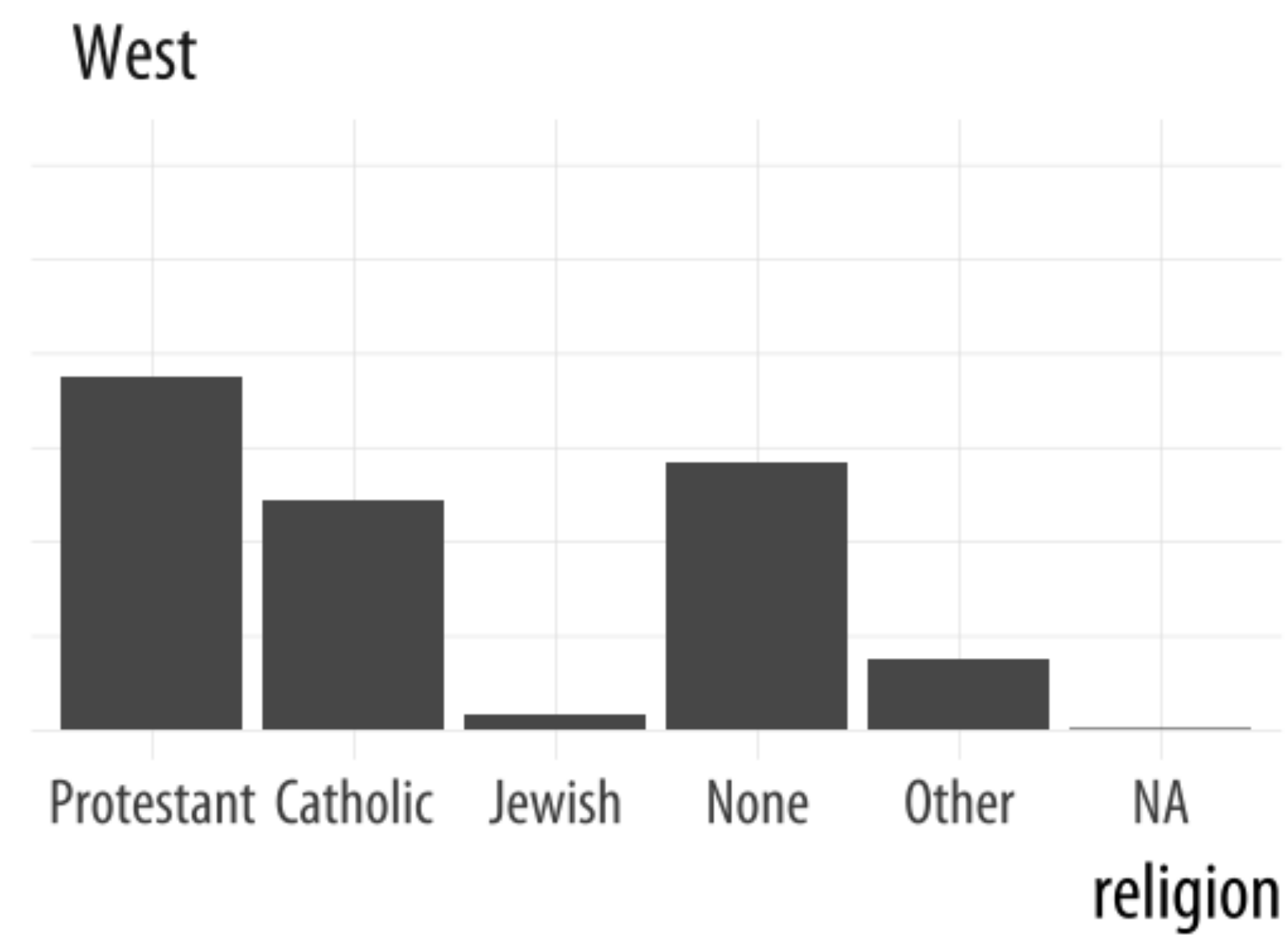
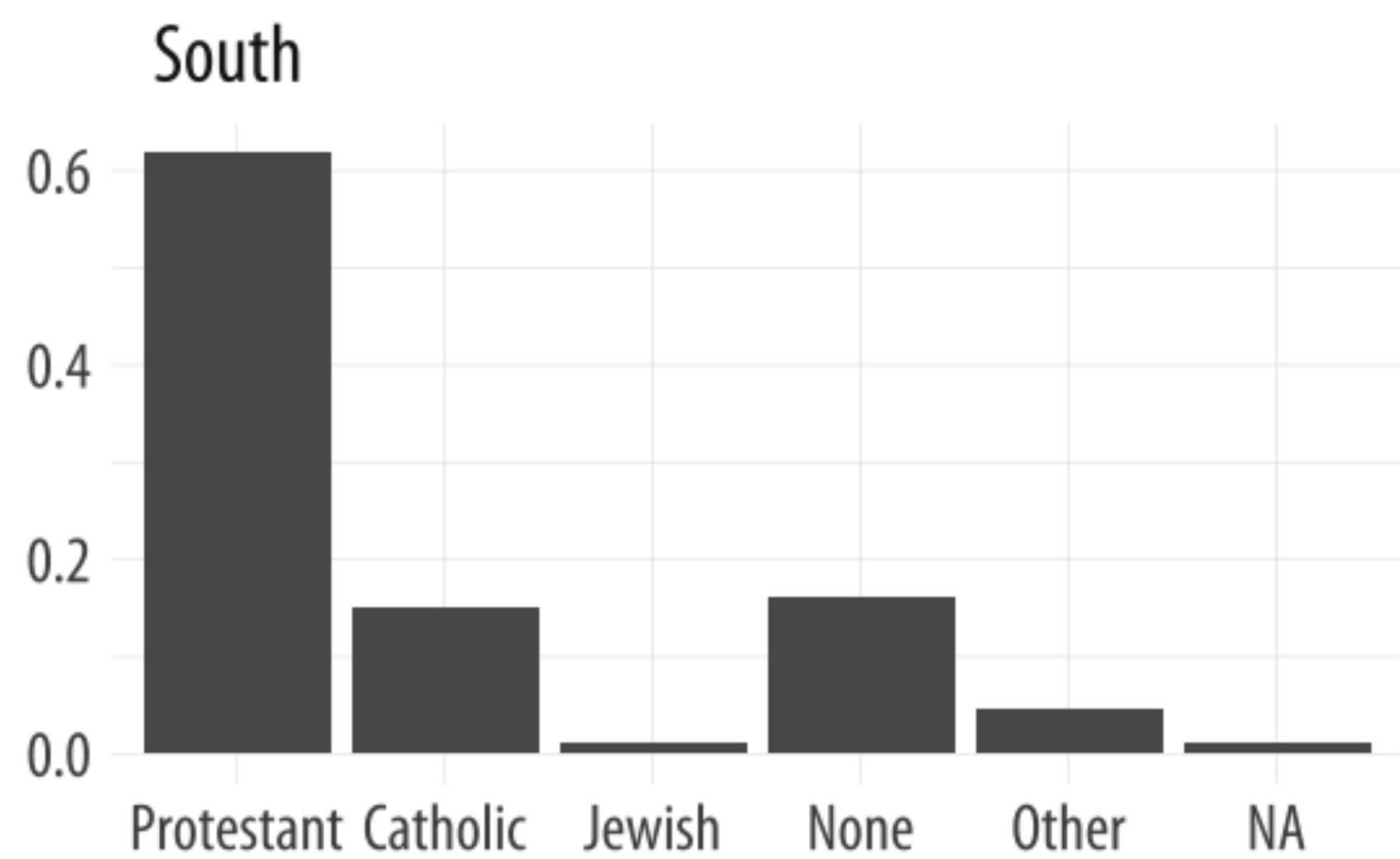
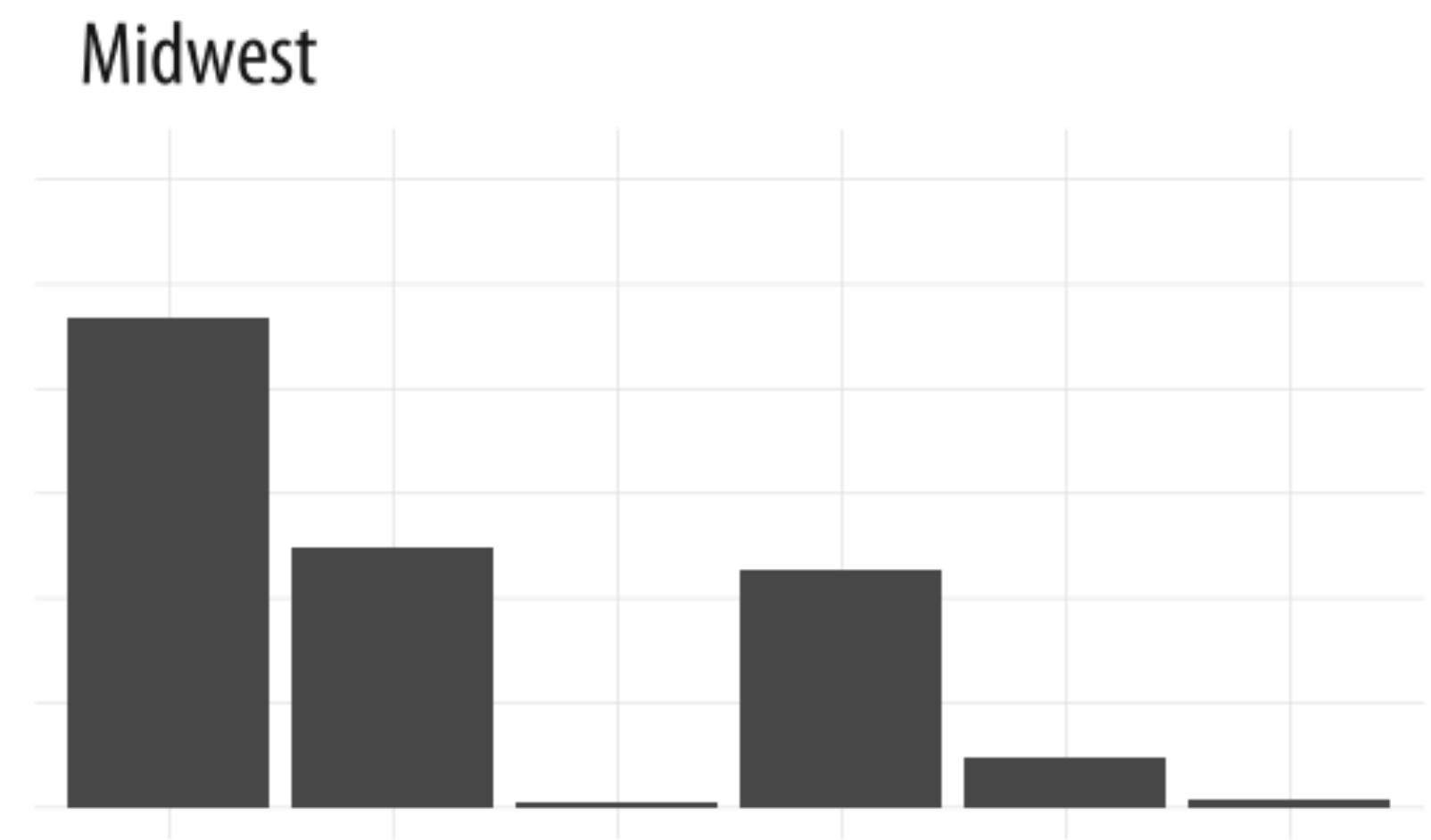
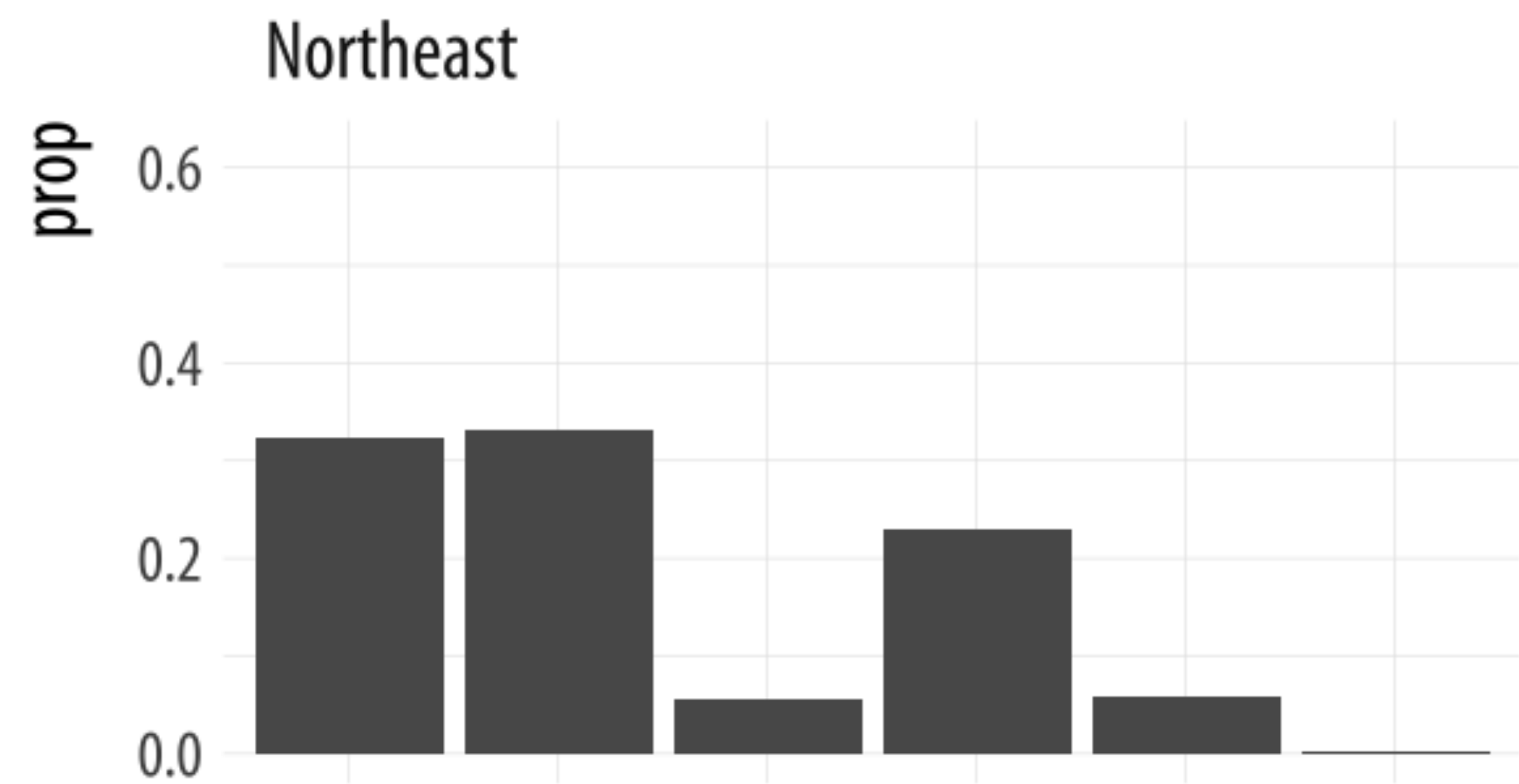
```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = bigregion,  
                          fill = religion))  
p + geom_bar(position = "dodge",  
            mapping = aes(y = ..prop..,  
                          group = religion))
```



Still not right!

# Time to take a step back

```
p <- ggplot(data = gss_sm,  
            mapping = aes(x = religion))  
p + geom_bar(position = "dodge",  
            mapping = aes(y = ..prop..,  
                          group = bigregion)) +  
  facet_wrap(~ bigregion, ncol = 2)
```



**SURELY THINGS  
CAN BE EASIER  
THAN THIS?**