

GenSamp: RESULTS

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## GenSamp: RESULTS

**Setup****Packages and Data****Organize Objects**

```
## Joining, by = "DSID"
```

```
##
```

```
## -- Column specification -----
```

```
## cols(
```

```
##   vnames = col_character(),
```

```
##   Variables = col_character(),
```

```
##   Sub = col_character(),
```

```
##   Category = col_character(),
```

```
##   Type = col_character()
```

```
## )
```

```
##
```

```
##
```

```
## -- Column specification -----
```

```
## cols(
```

```
##   vnames = col_character(),
```

```
##   Variables = col_character(),
```

```
##   Sub = col_character(),
```

```
##   Category = col_character(),
```

```
##   Type = col_character()
```

```
## )
```

```
## Joining, by = "vnames"
```

## Data Summary

### Covaraite Statistics

```
## 'summarise()' ungrouping output (override with '.groups' argument)
```

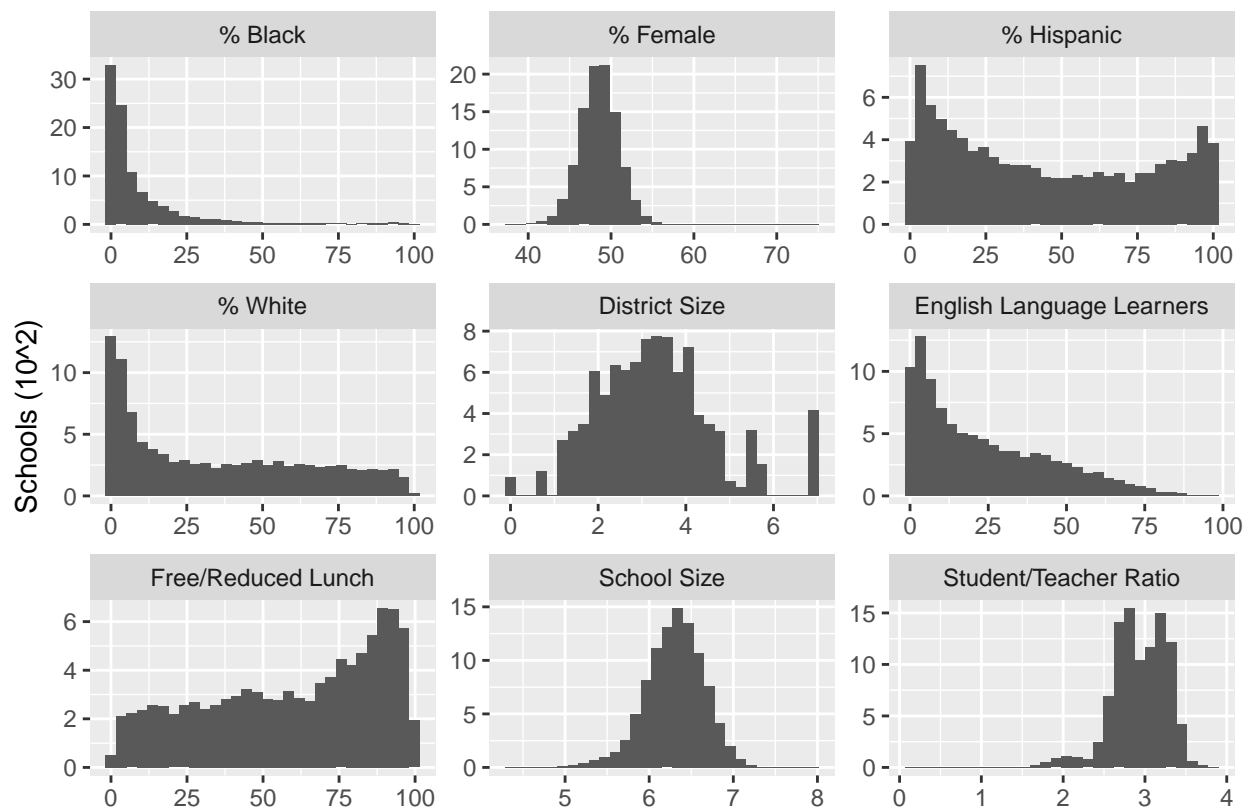
```
## Joining, by = "vnames"
```

```
## Joining, by = c("vnames", "Variables", "Sub", "Category", "Type")
```

### Continuous variable distributions

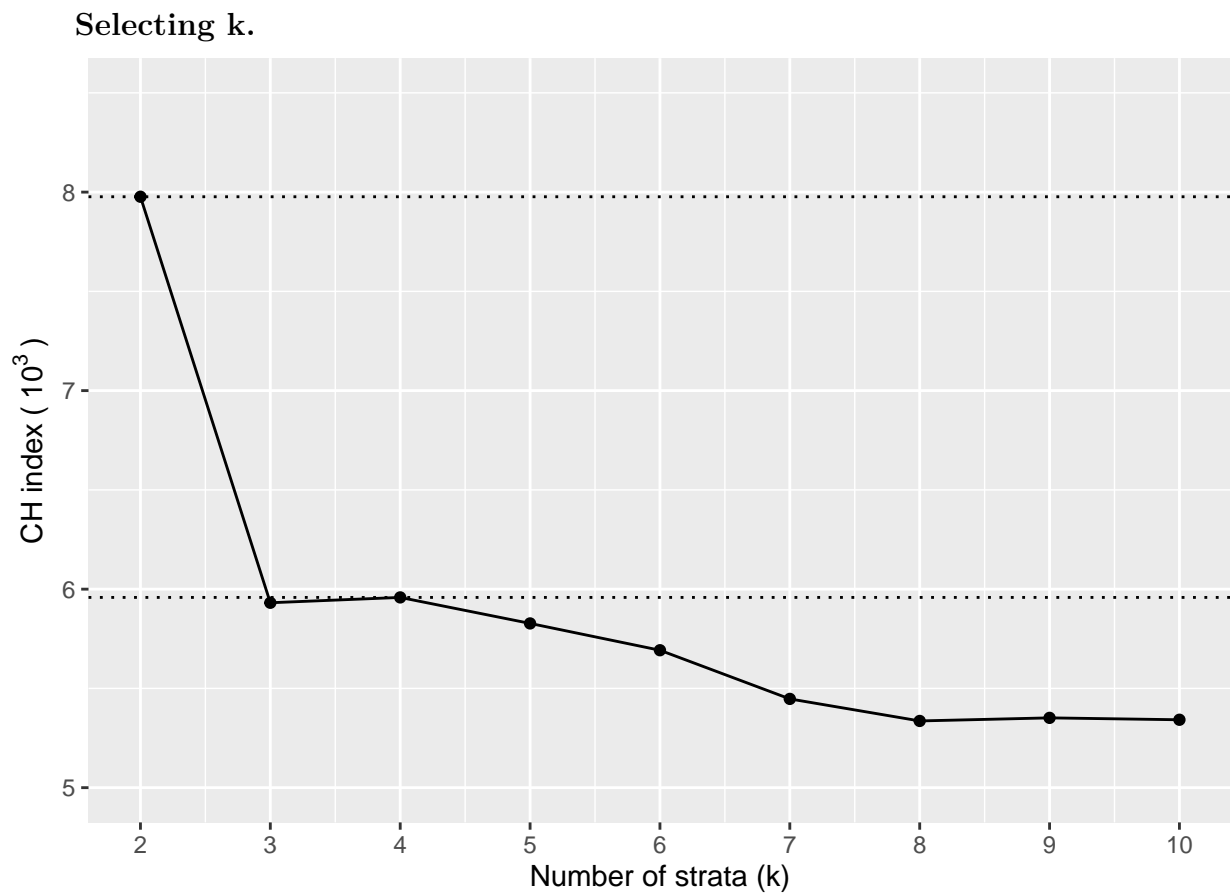
```
## Joining, by = "vnames"
```

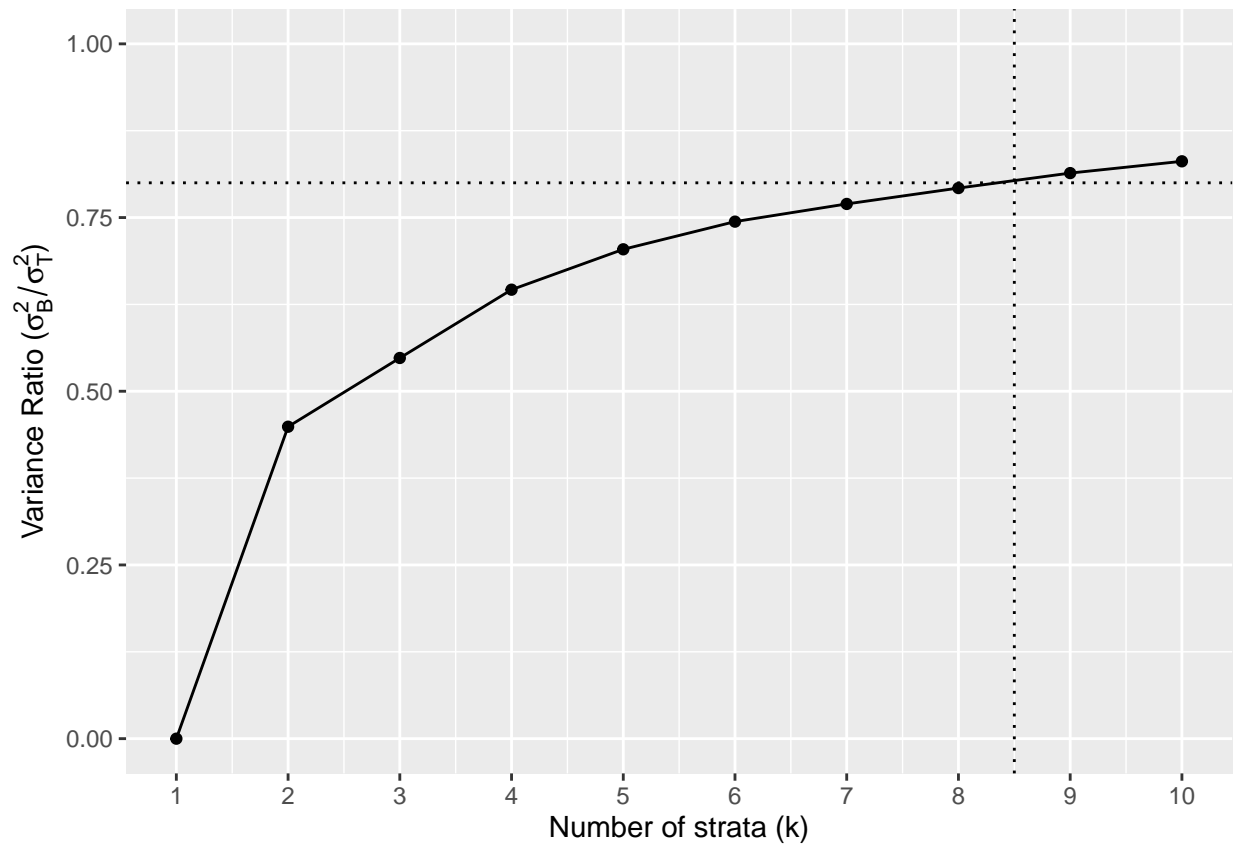
```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```



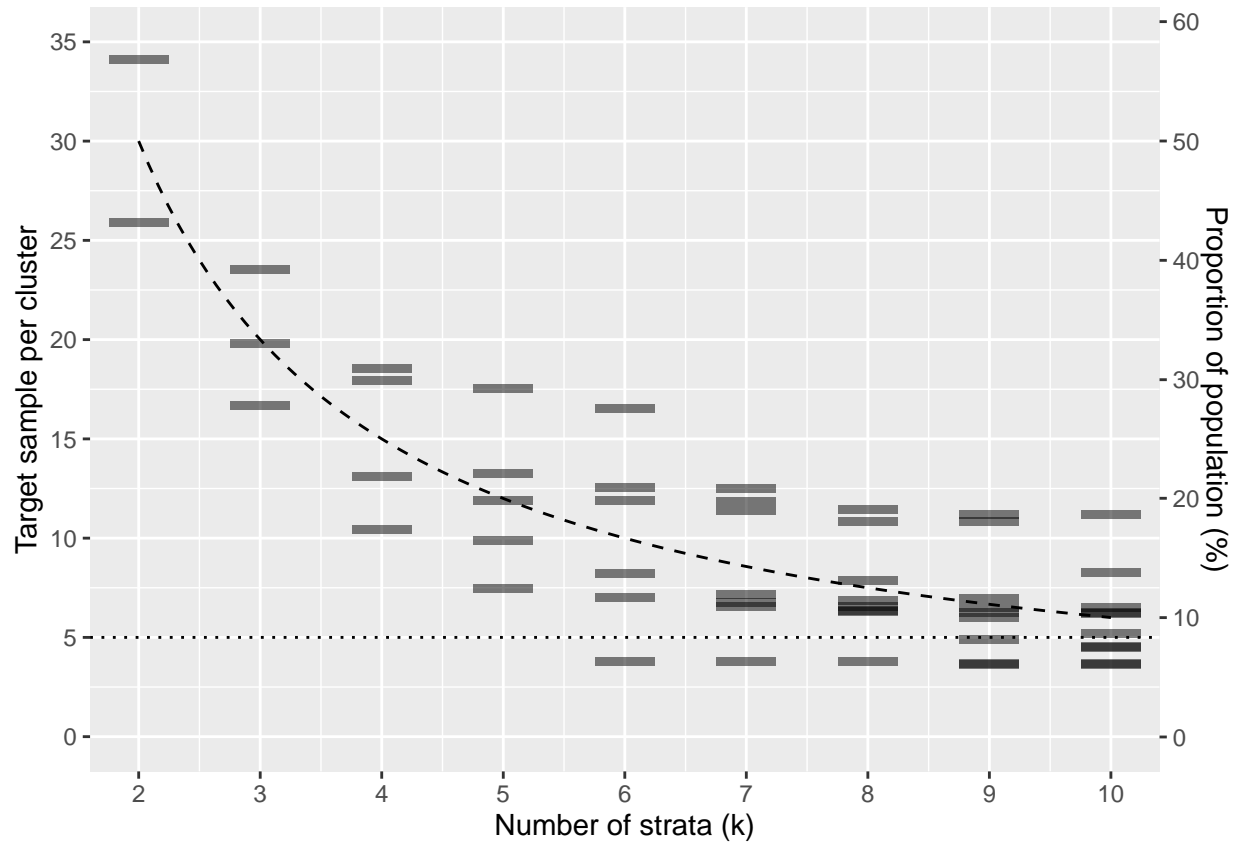
Methods Summary

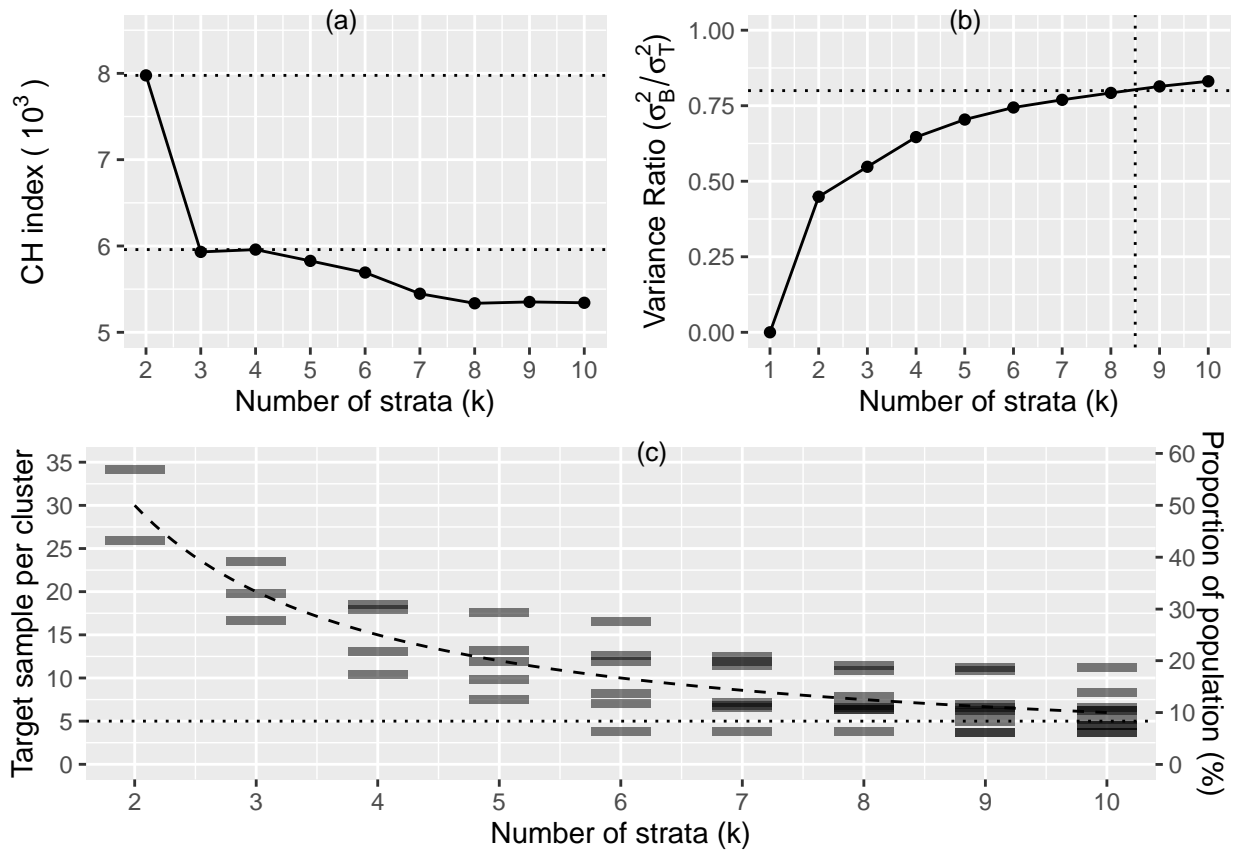
Cluster Analysis





```
## 'summarise()' regrouping output by 'k' (override with '.groups' argument)
```



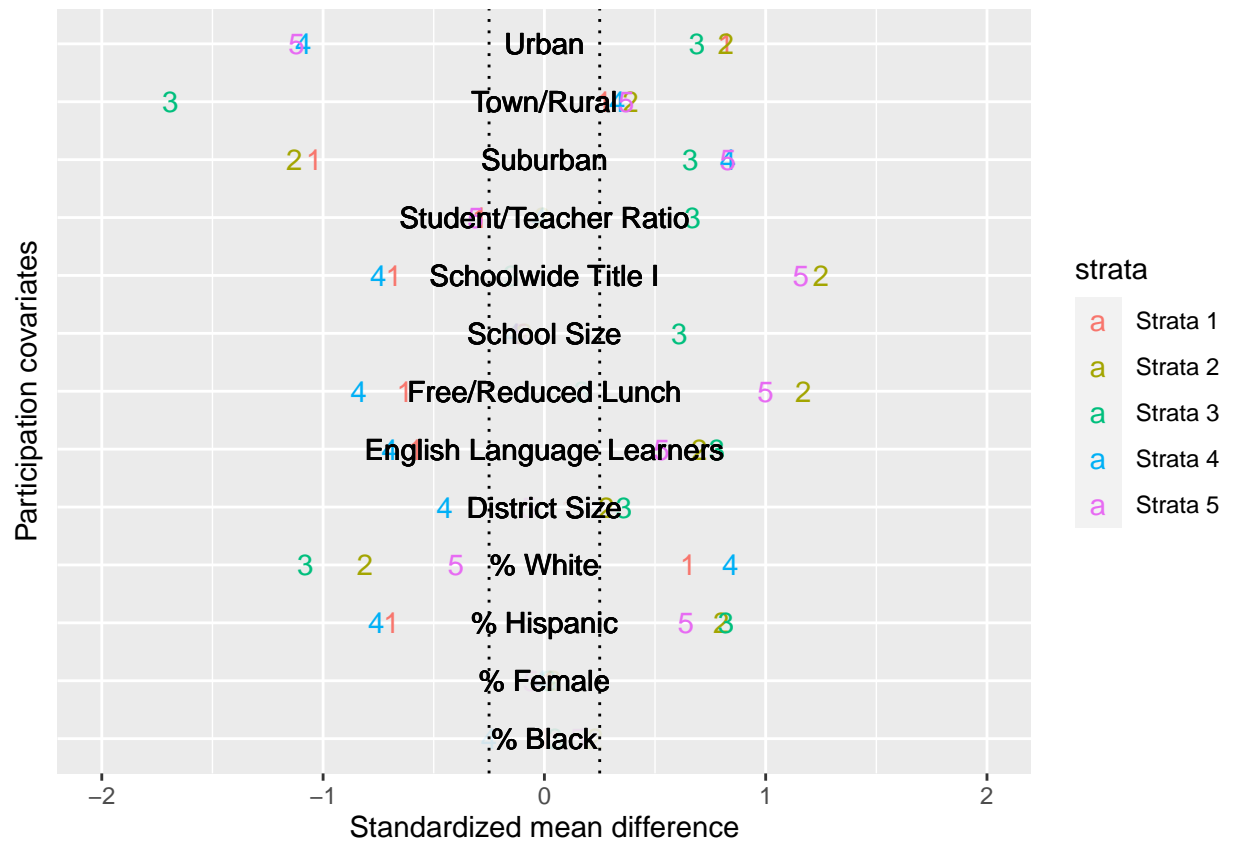


```
## 'summarise()' regrouping output by 'strata' (override with '.groups' argument)
```

```
## Joining, by = "var"
```

```
## Joining, by = "vnames"
```

```
## Warning: Using alpha for a discrete variable is not advised.
```



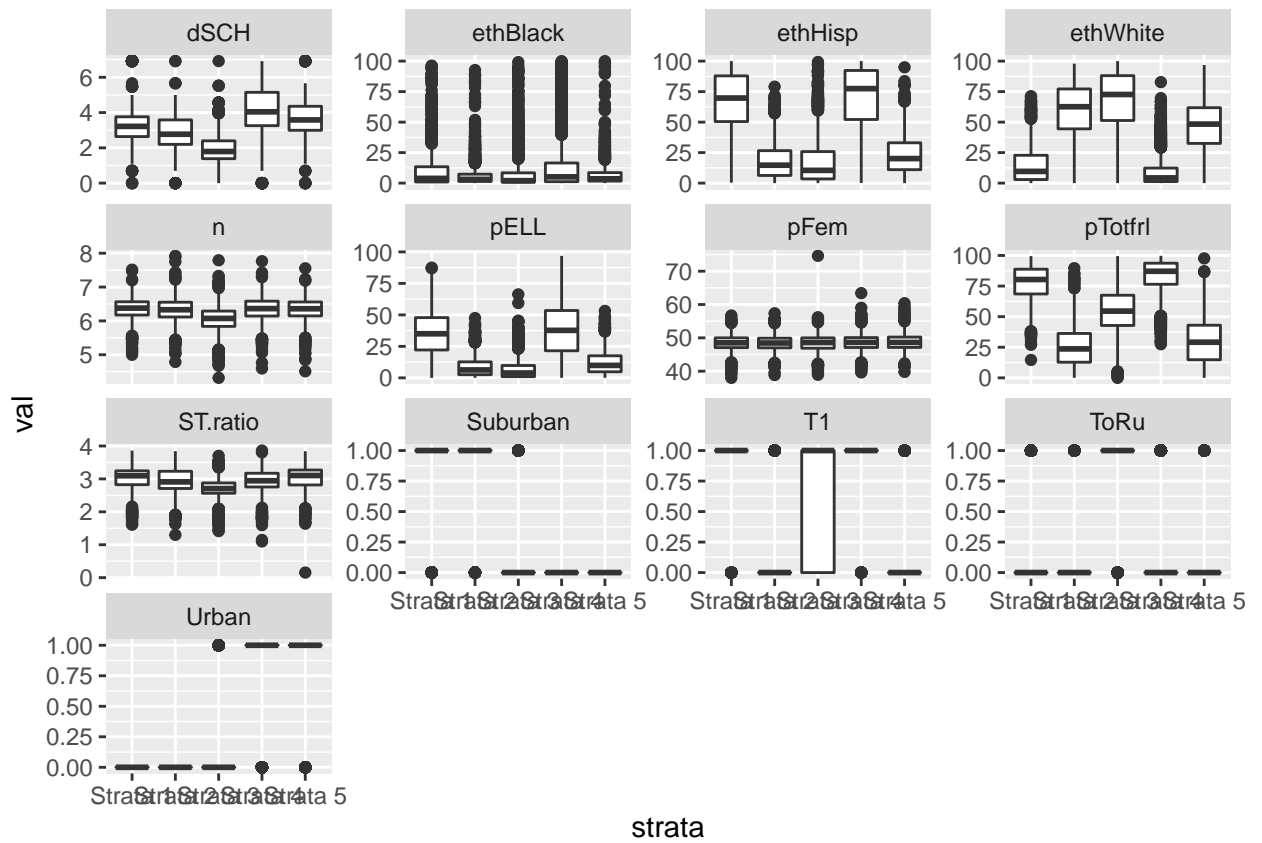
## Note: Using an external vector in selections is ambiguous.

## i Use 'all\_of(covariates)' instead of 'covariates' to silence this message.

## i See <<https://tidyselect.r-lib.org/reference/faq-external-vector.html>>.

## This message is displayed once per session.

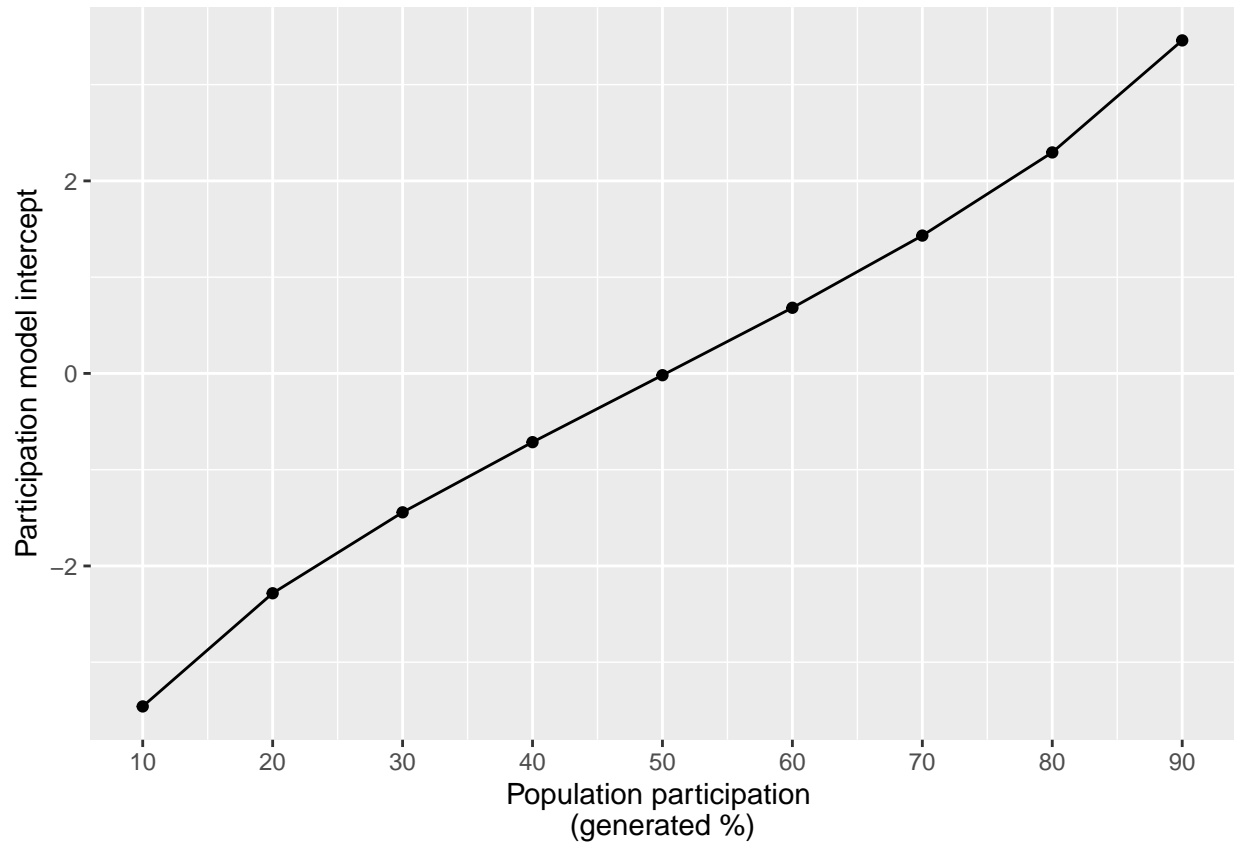




### Variation explained by the strata

### Participation Generating Model

log_odds	Variables	Sub	Category	Type
0.019	Schoolwide Title I	Status	School Data	Prop
0.374	School Size	Enrollment	School Data	Mean
0.081	Free/Reduced Lunch	Status	Student Data	Mean
0.433	Urban	Urbanicity	School Data	Prop
0.007	Suburban	Urbanicity	School Data	Prop
-0.403	Town/Rural	Urbanicity	School Data	Prop
-0.538	% White	Ethnicity	Student Data	Mean
0.291	% Black	Ethnicity	Student Data	Mean
0.395	% Hispanic	Ethnicity	Student Data	Mean
-0.019	% Female	Gender	Student Data	Mean
-0.101	Student/Teacher Ratio	Enrollment	School Data	Mean
0.520	District Size	District	School Data	Mean
0.412	English Language Learners	Status	Student Data	Mean



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

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

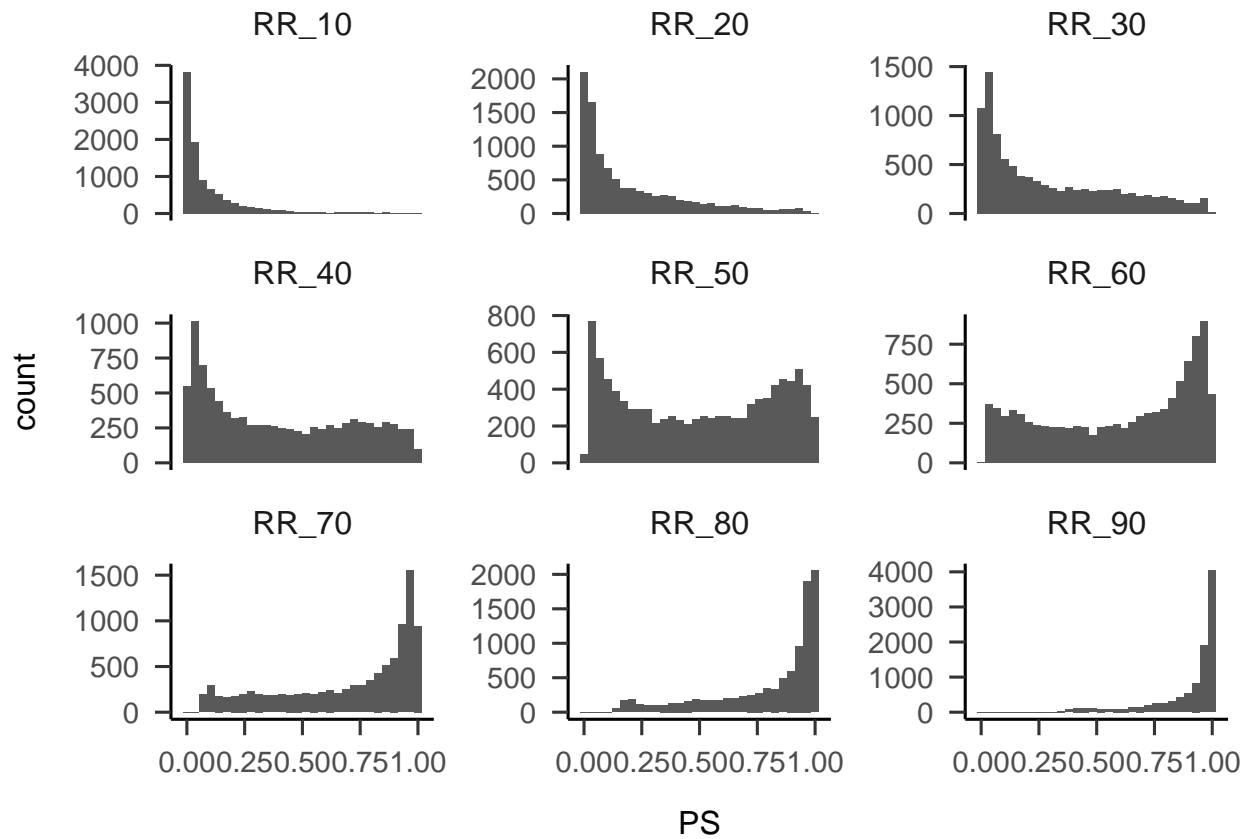
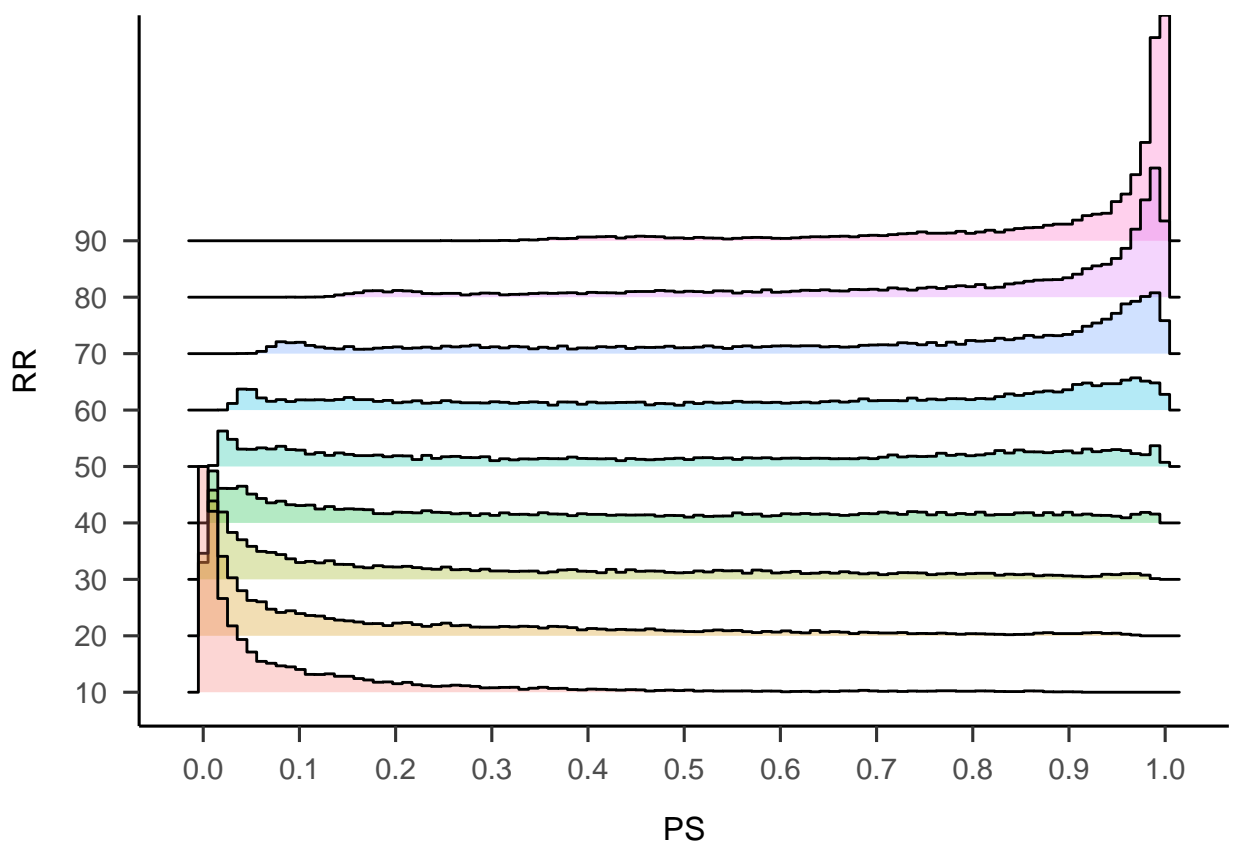


Figure 1. Distributions of Participation Propensity Scores



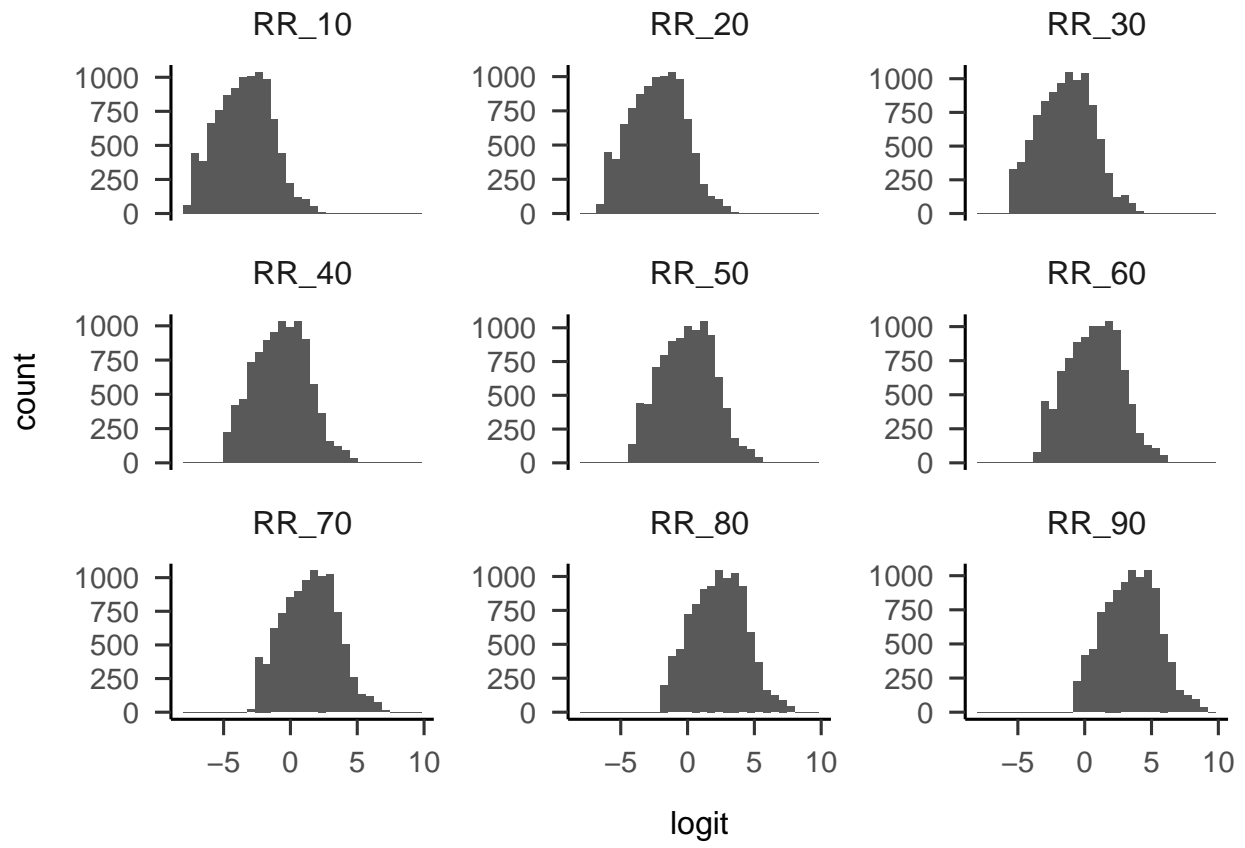
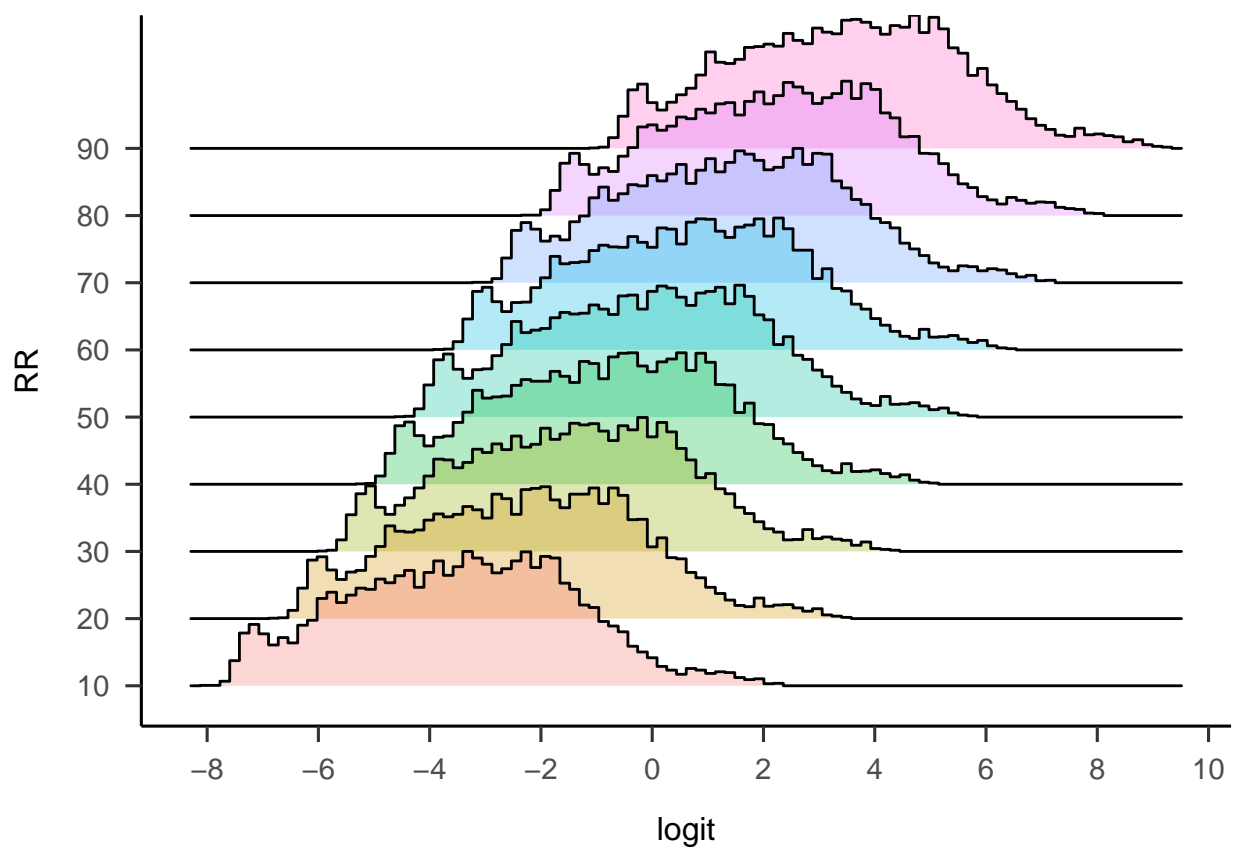
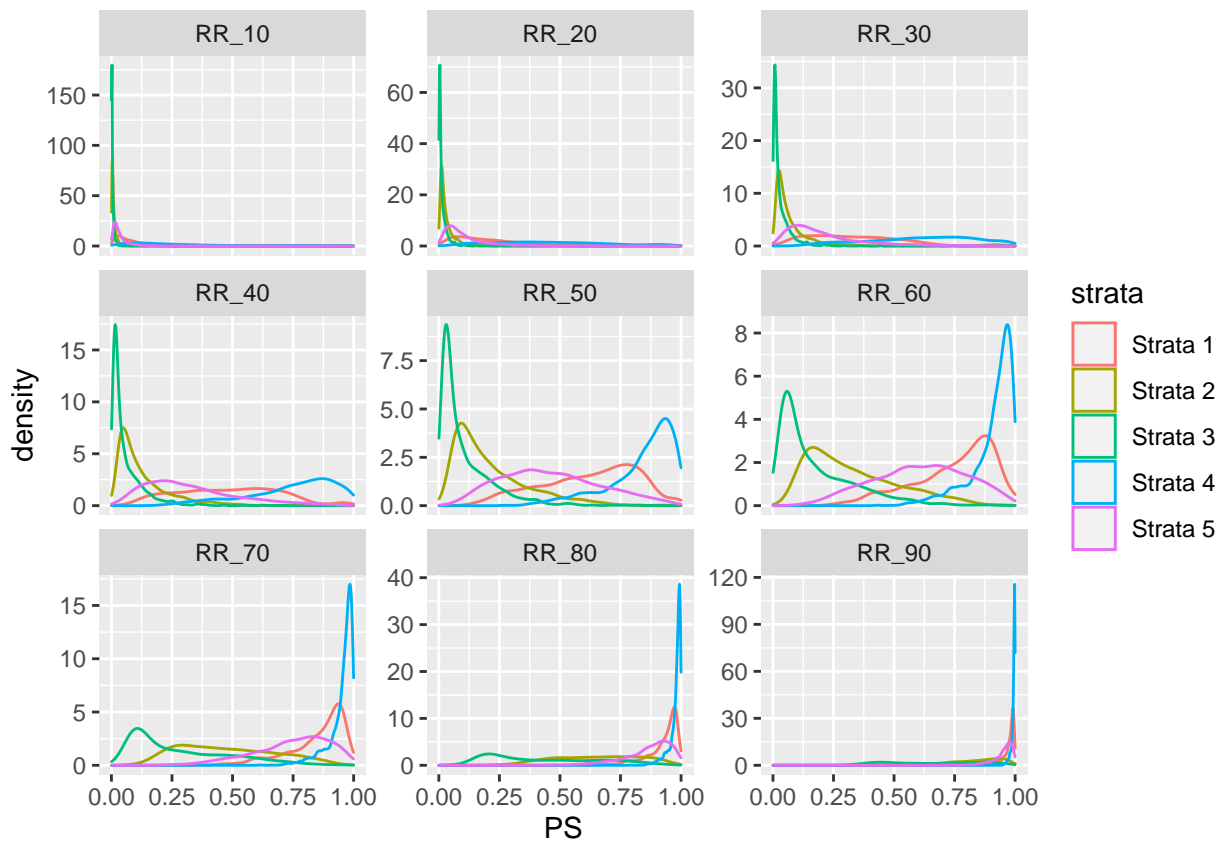


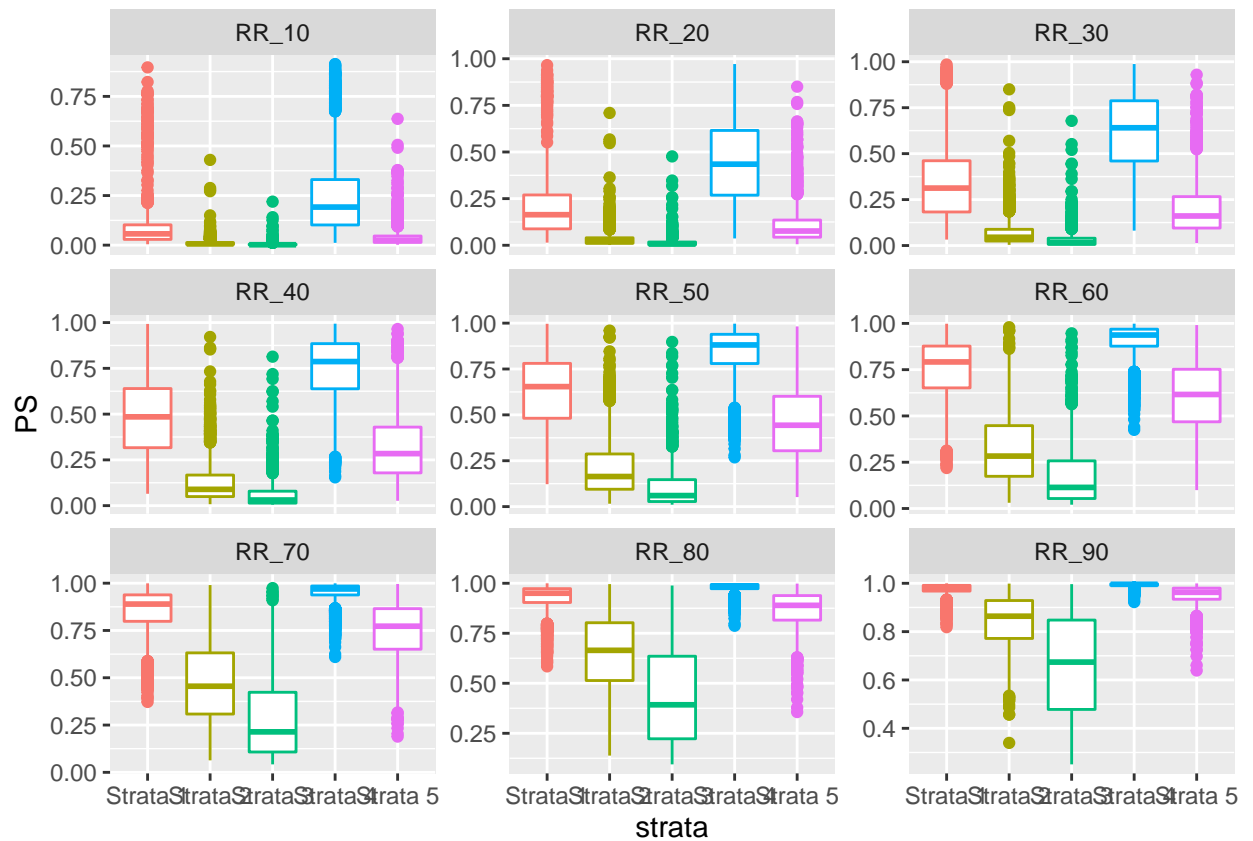
Figure 2. Distributions of Participation Propensity Scores



```
## Joining, by = "DSID"
```



```
## Joining, by = "DSID"
```



## Results

### Generalizability

#### B Index.

```
## 'summarise()' regrouping output by 'sample_method', 'RR' (override with '.groups' arg
```

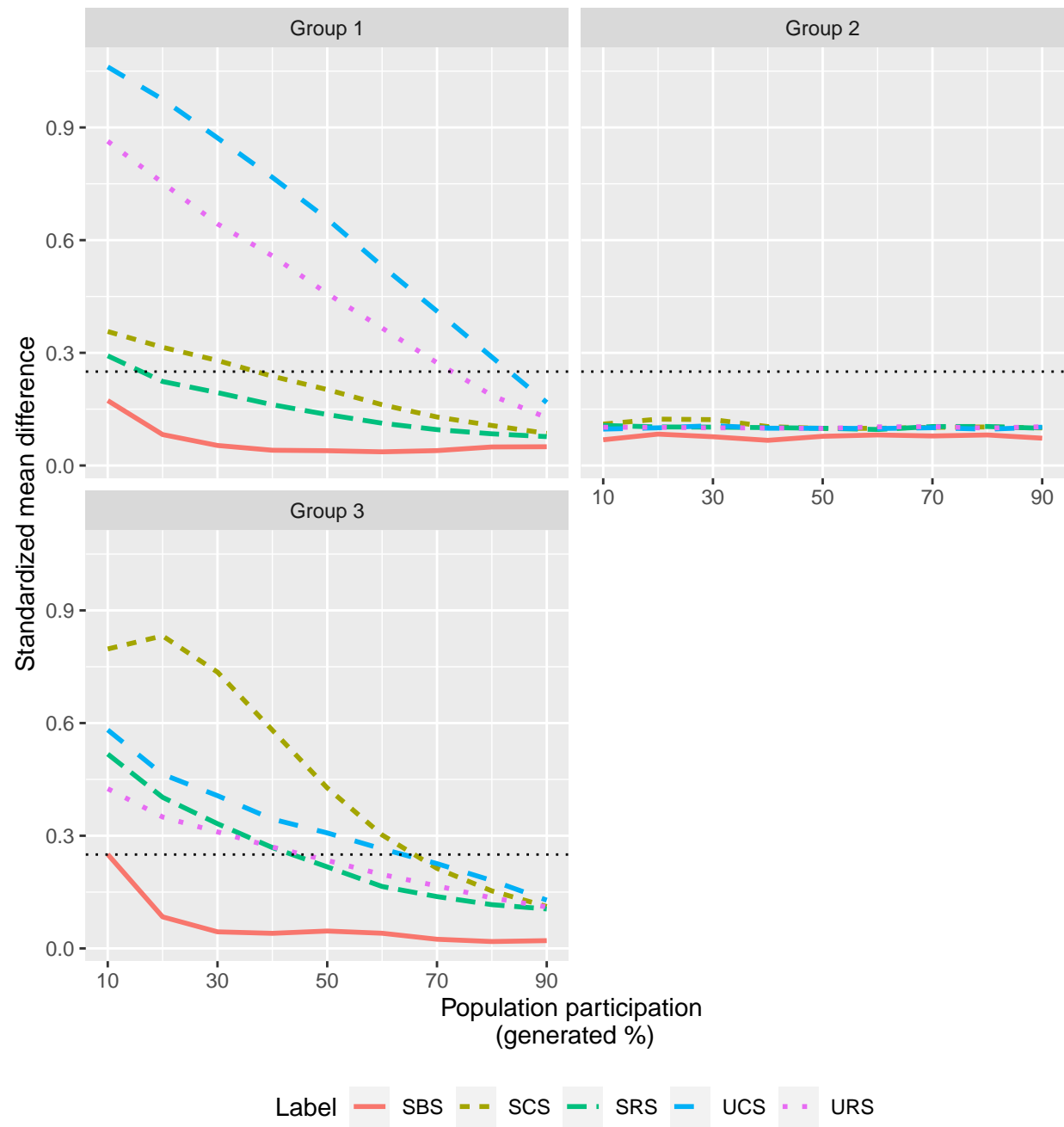
#### Standardized mean differences.

```
## Joining, by = "var"
```

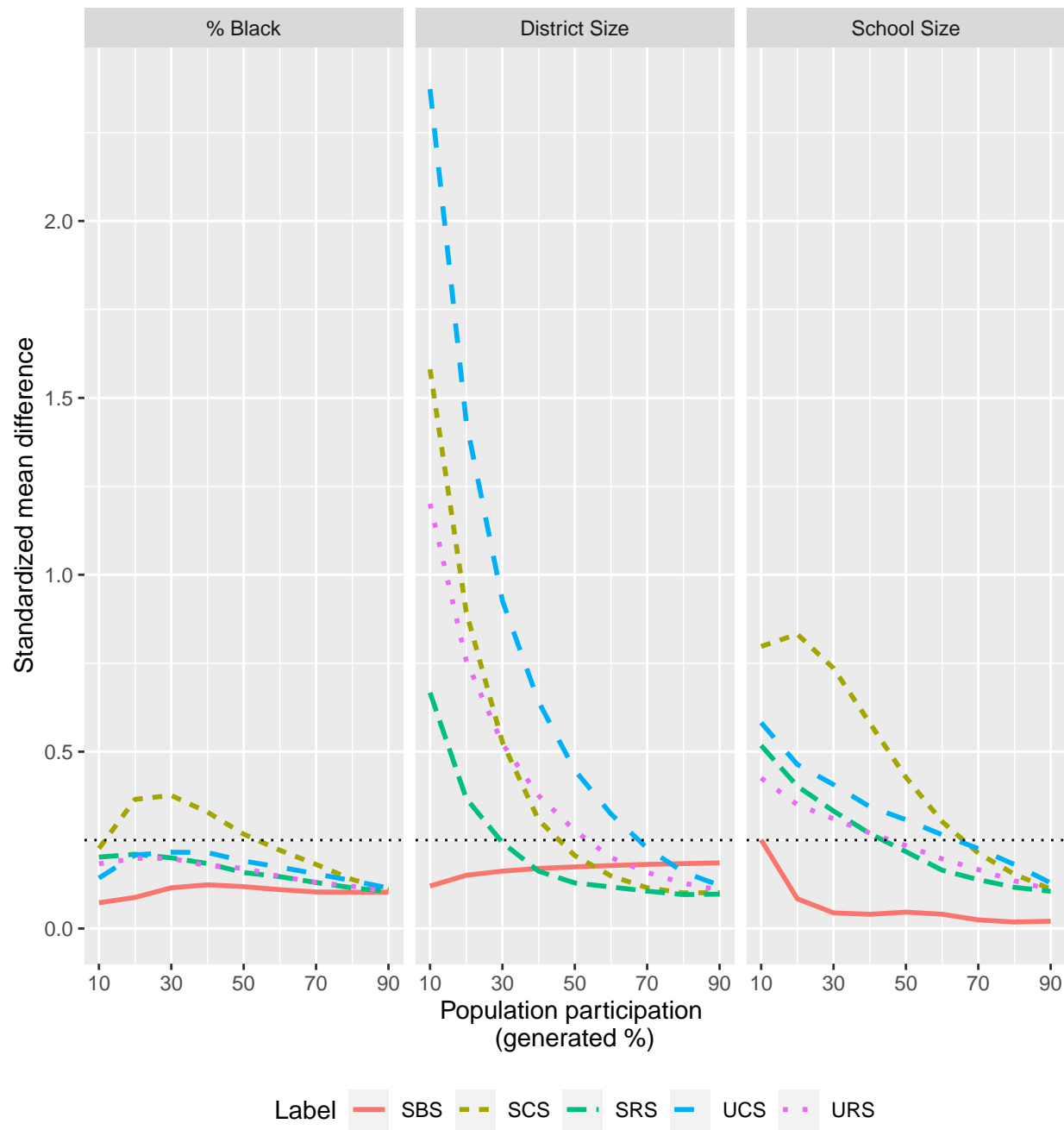
```
## 'summarise()' regrouping output by 'sample_method', 'RR', 'var' (override with '.grou
```

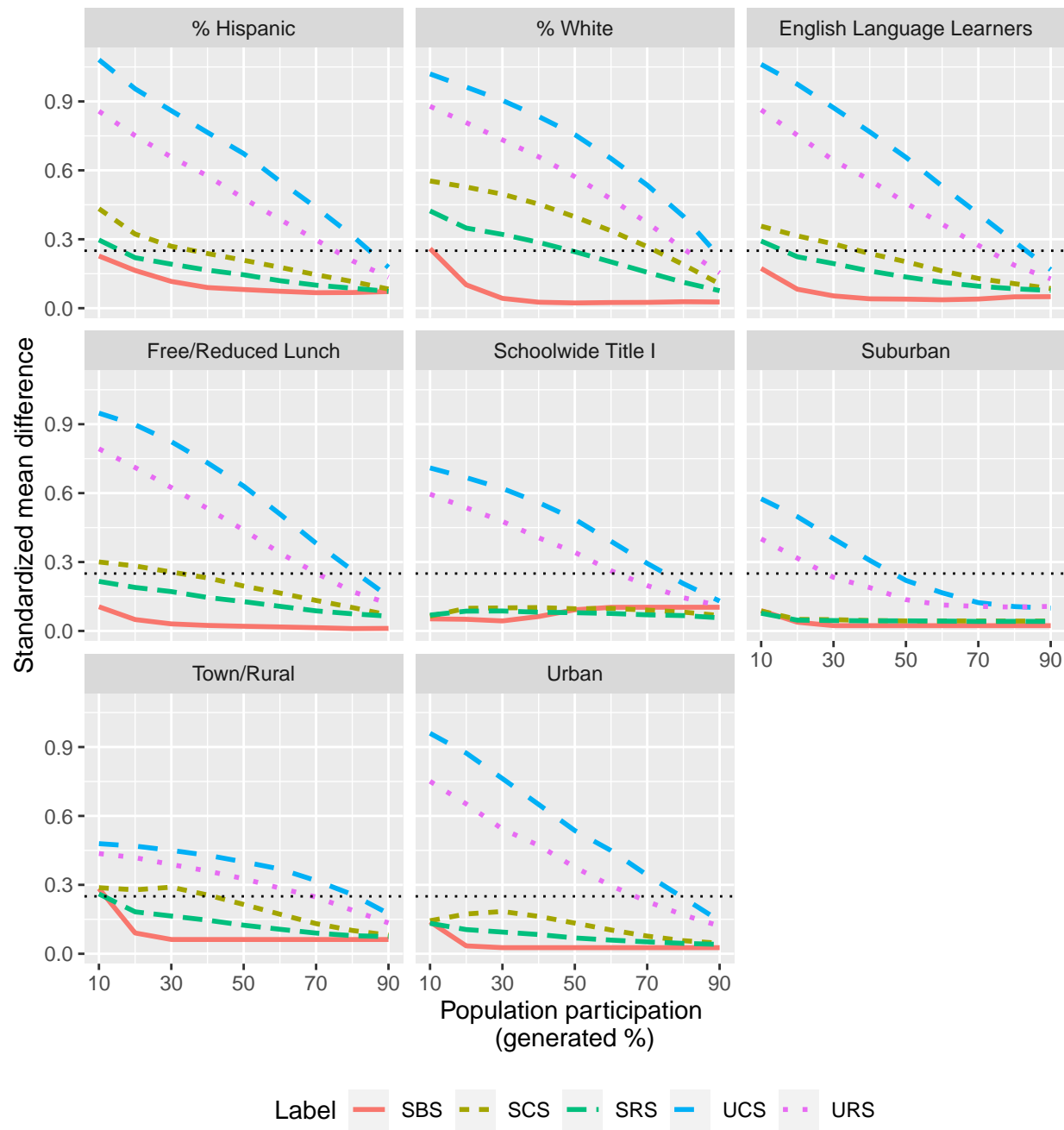
```
## Joining, by = "vnames"
```

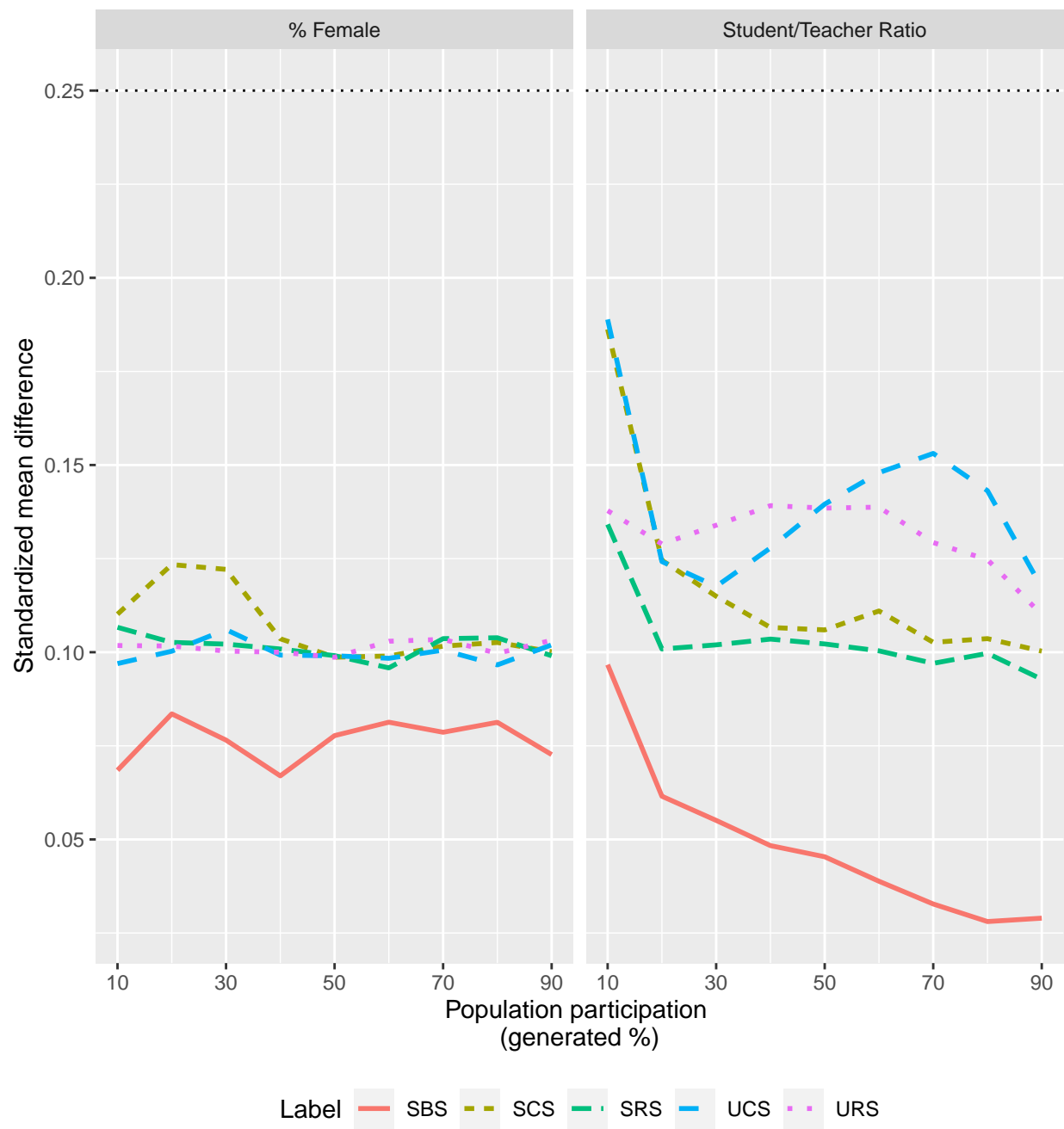
```
## Joining, by = "vnames"
```

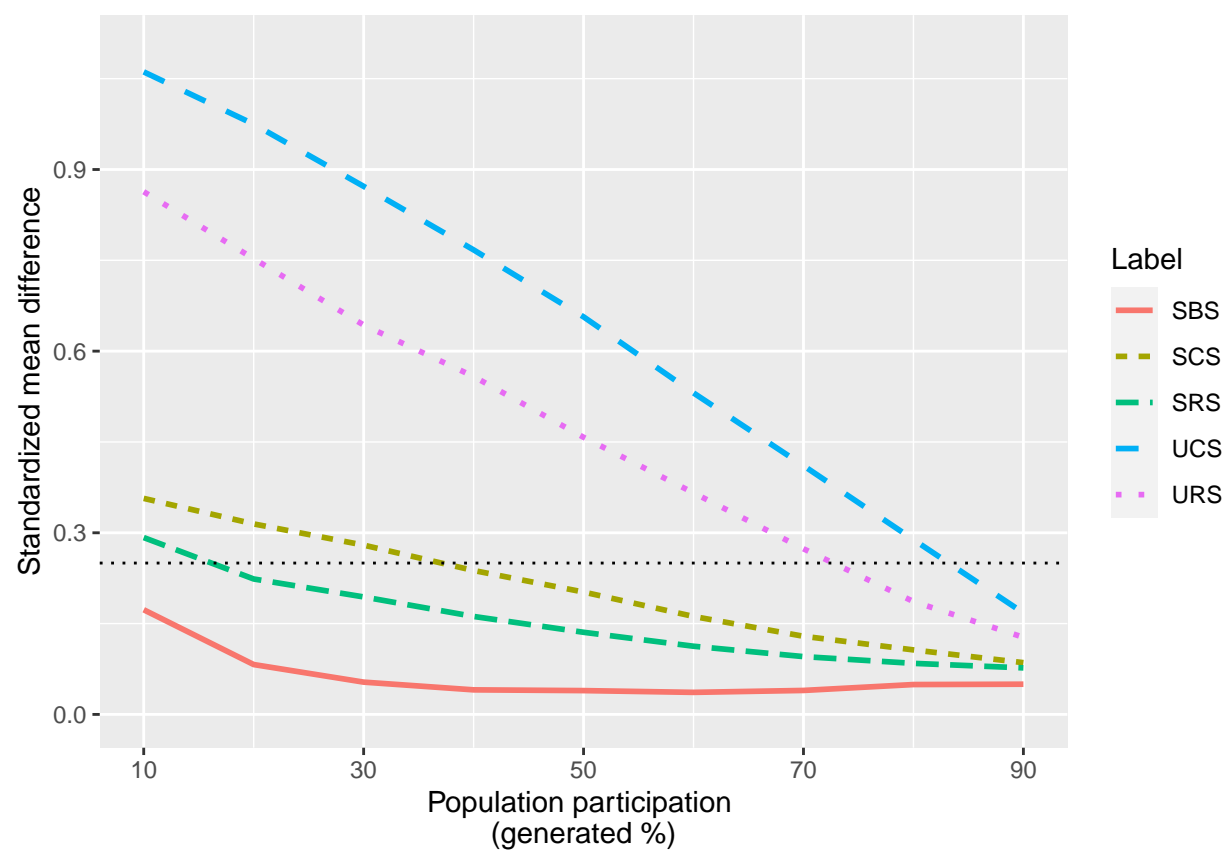


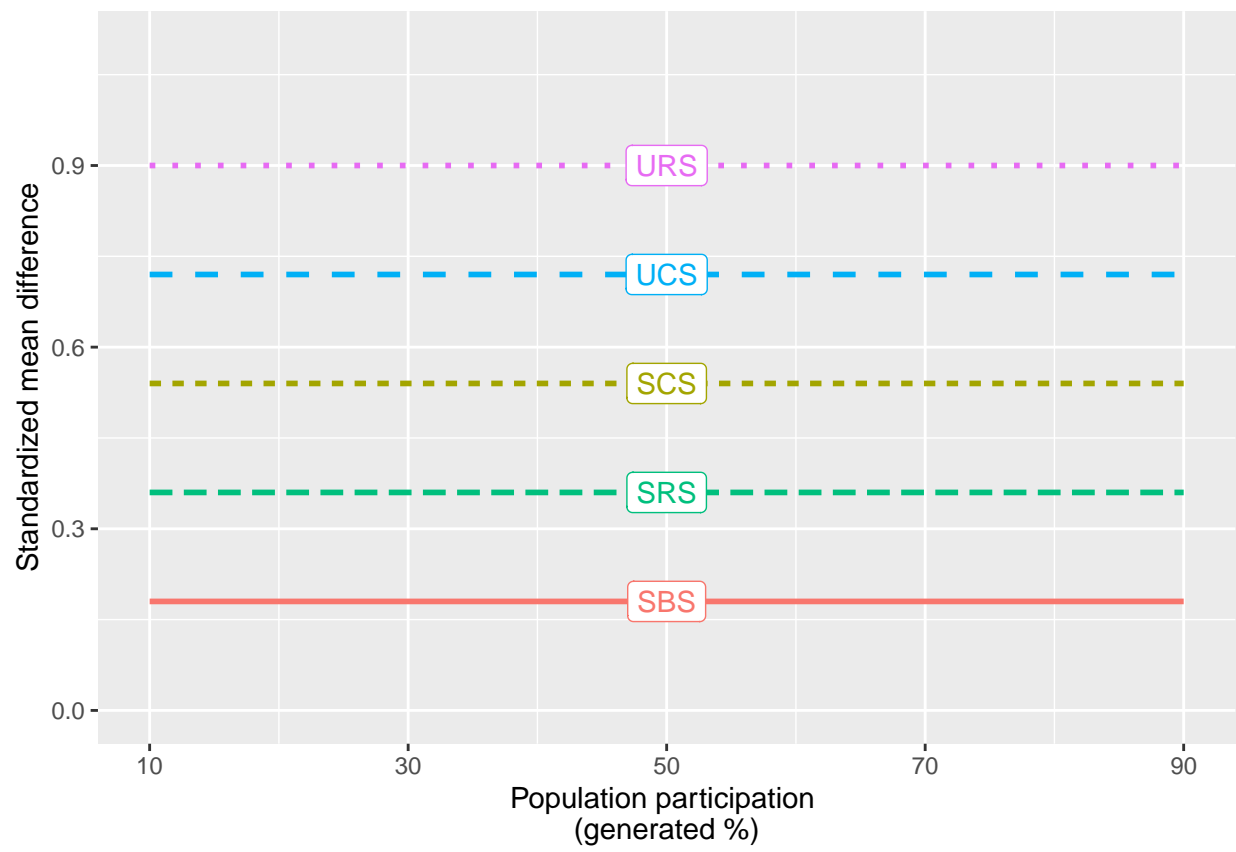


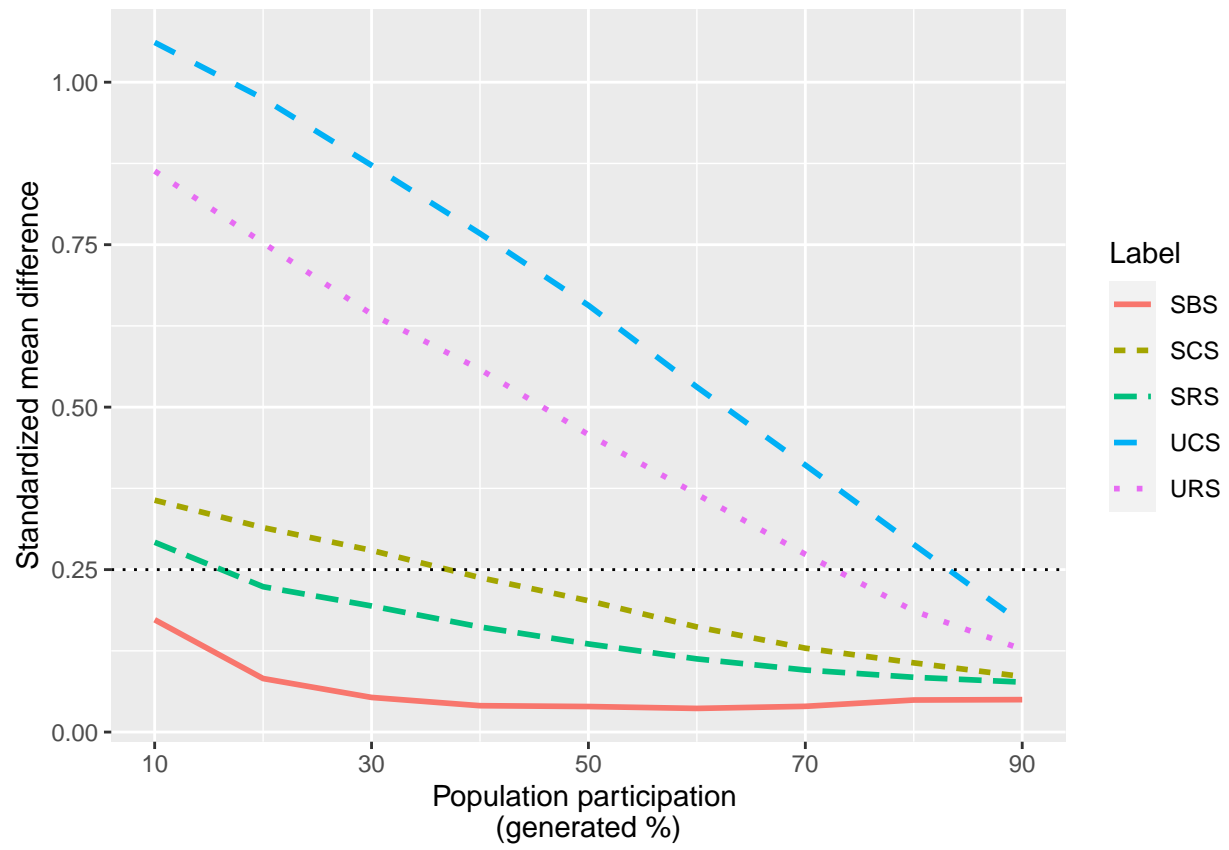


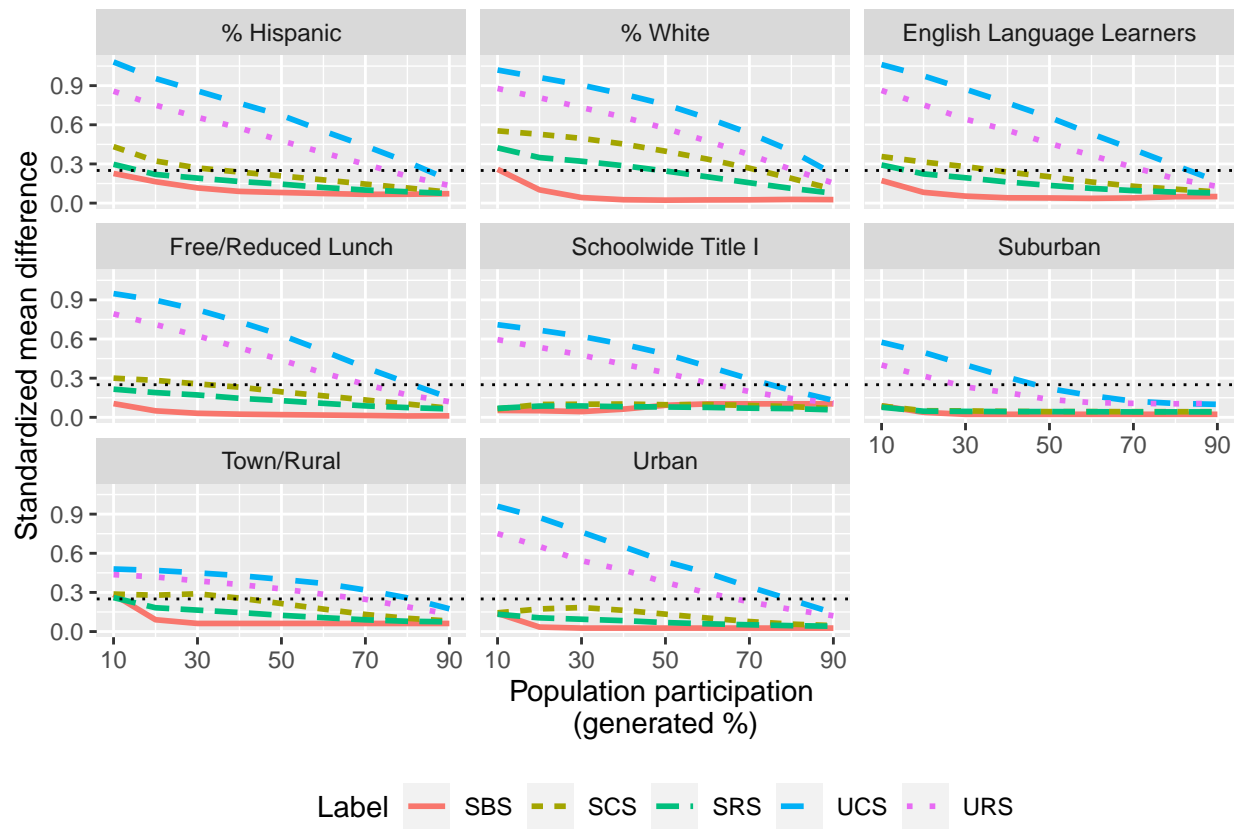


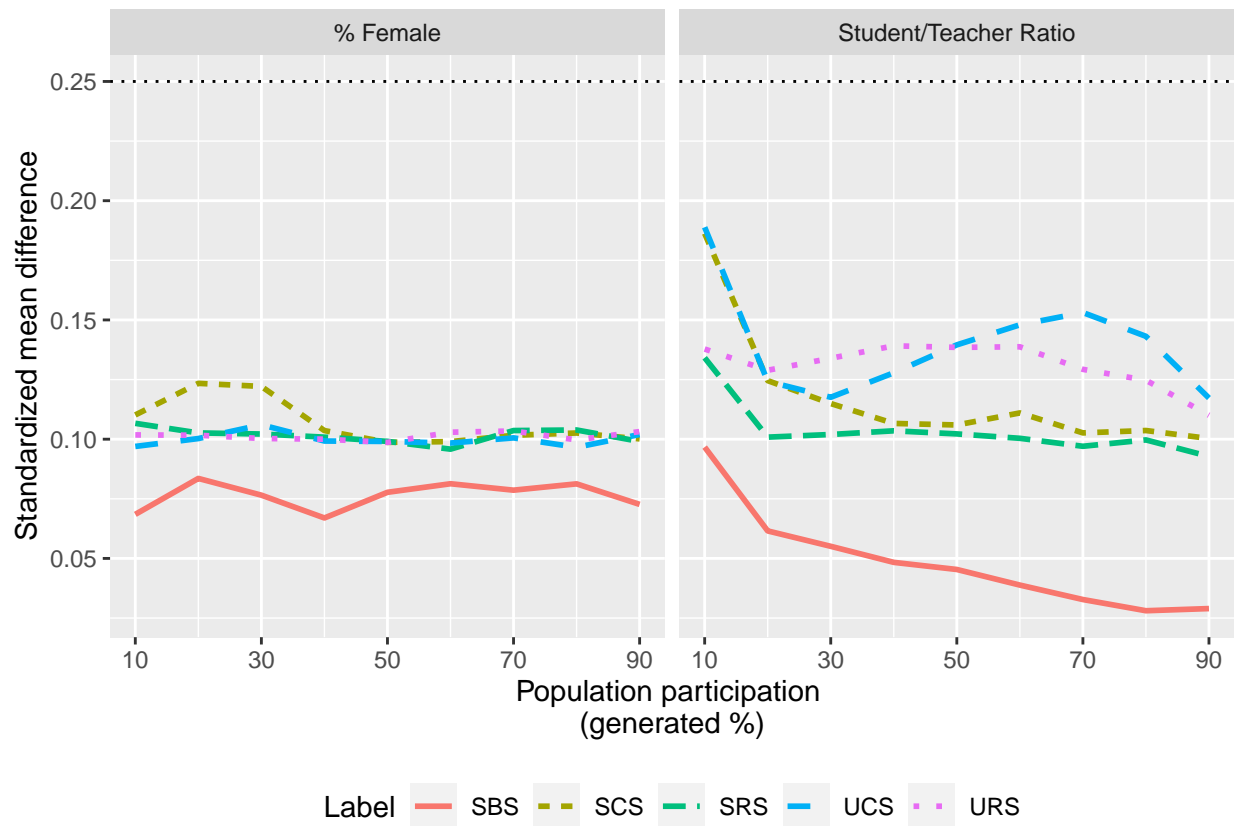












### Examples for presentations

**V-ratio and Log odds.**

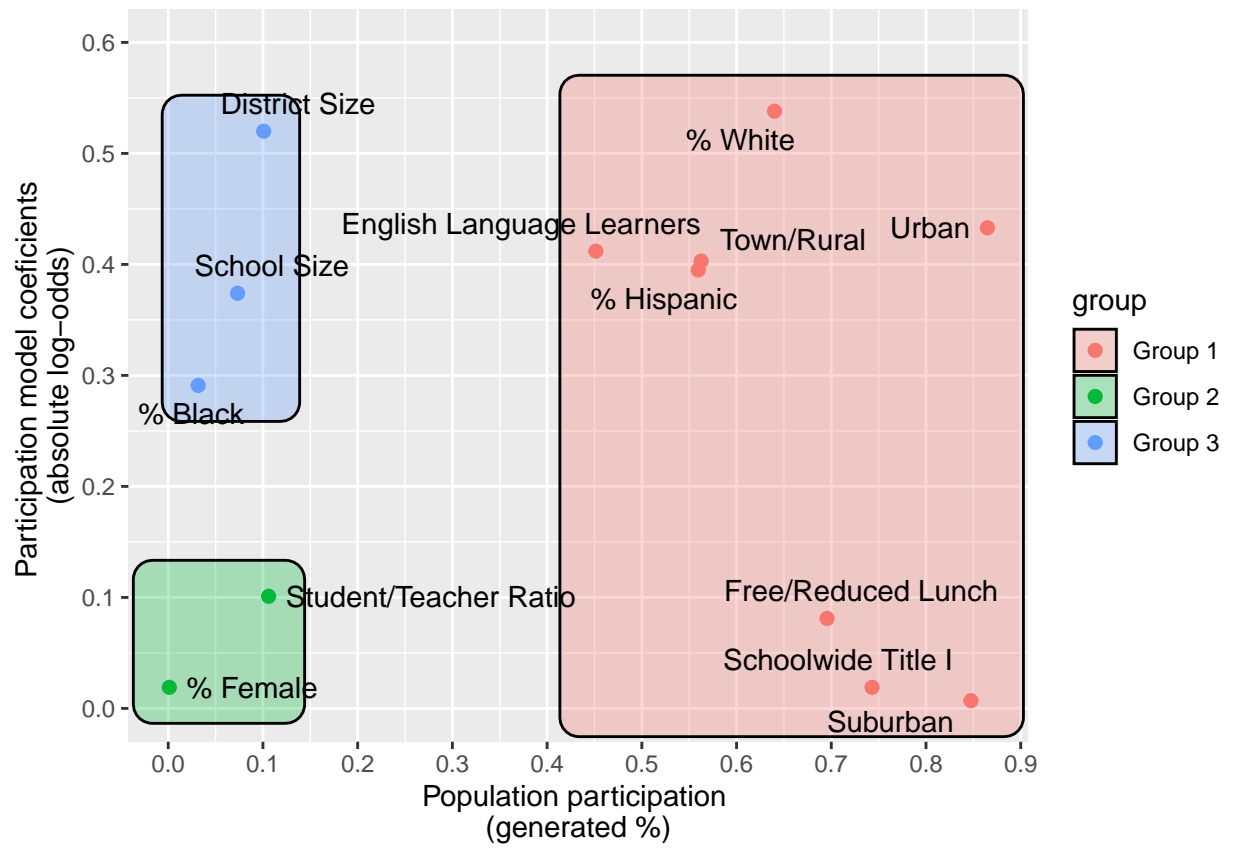
```
## 'summarise()' regrouping output by 'vnames', 'T.SS' (override with '.groups' argument)
```

```
## 'summarise()' regrouping output by 'vnames' (override with '.groups' argument)
```

```
## Joining, by = "vnames"
```

```
## Joining, by = "vnames"
```

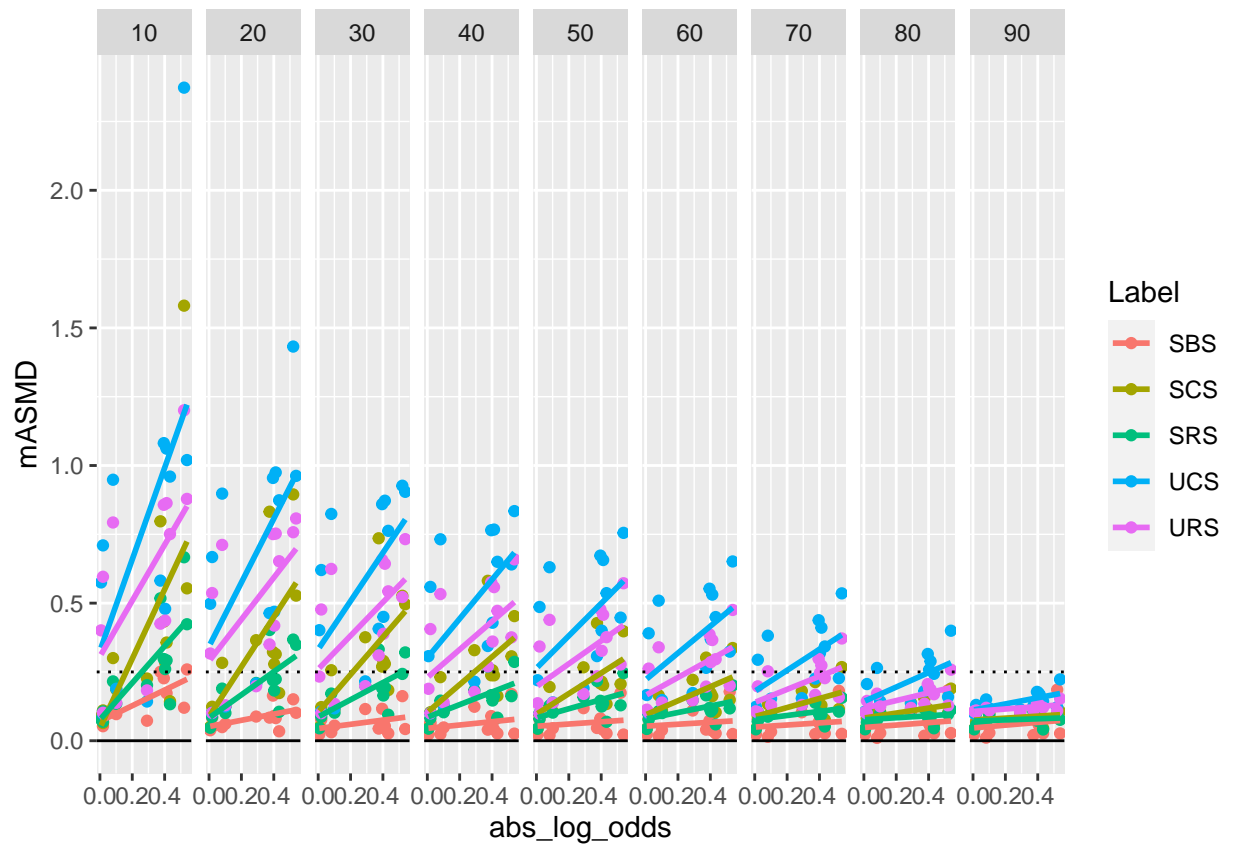




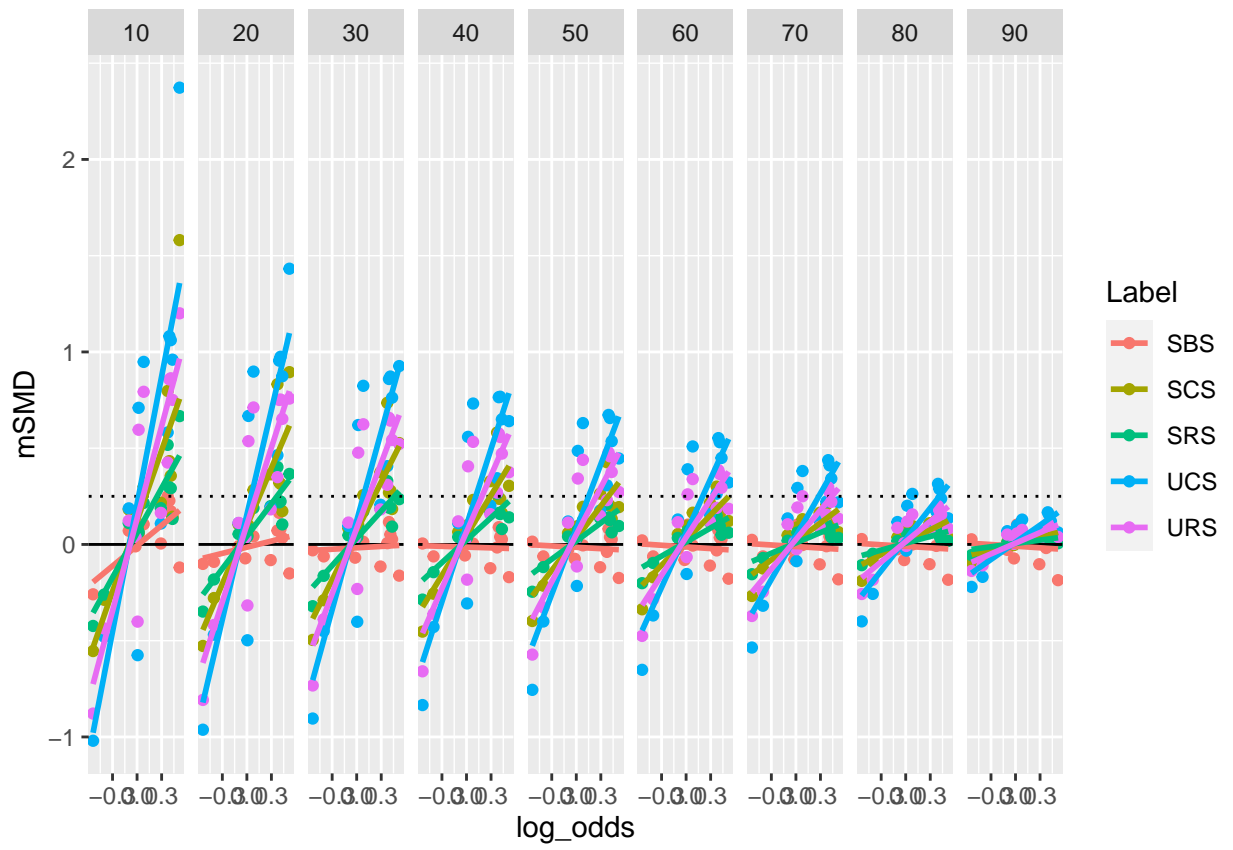
Test.

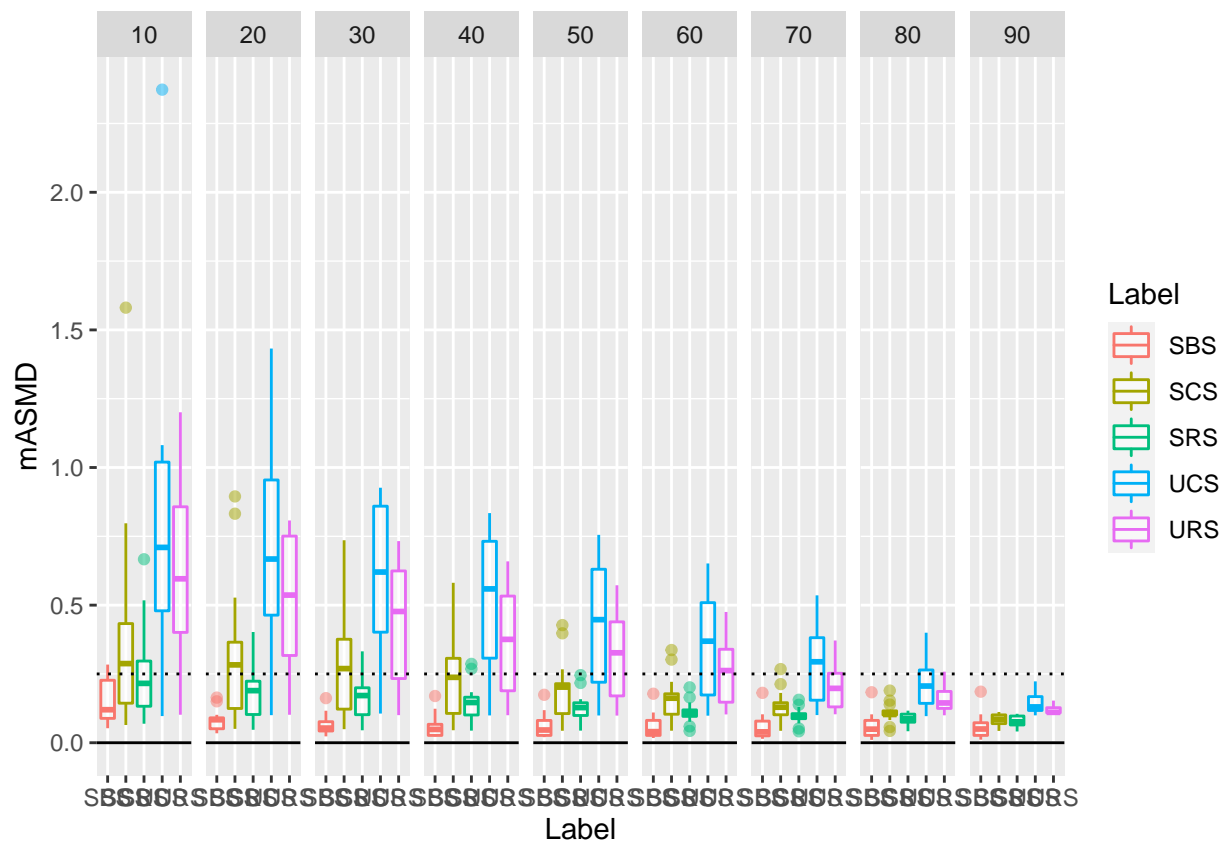
```
## Joining, by = c("vnames", "log_odds")
```

```
## 'geom_smooth()' using formula 'y ~ x'
```



```
## 'geom_smooth()' using formula 'y ~ x'
```





Analysis of Variance Table

Model 1: mSMD ~ RR Model 2: mSMD ~ log_odds + vratio + RR Model 3: mSMD									
~ log_odds * vratio * RR Res.Df RSS Df Sum of Sq F Pr(>F)									
1	583	59.336							
2	581	38.107	2	21.2295	196.574	< 2.2e-16	<b>3</b>	<b>577</b>	<b>31.157</b>
							<b>4</b>	<b>6.9494</b>	<b>32.174</b>
<b>2.2e-16</b> — Signif. codes: 0 “’ <b>0.001</b> ” 0.01 ” 0.05 “.” 0.1 ’ ’ 1									

Model 1 Model 2 Model 3

(Intercept)			0.24 ***	0.15 ***	0.11 *
(0.03)			(0.03)	(0.05)	
RR			-0.00 ***	-0.00 ***	-0.00
(0.00)			(0.00)	(0.00)	(0.00)

log\_odds 0.61 \*\*\* 1.24 (*0.03*) (*0.16*)

*vratio 0.04 -0.02*

(*0.03*) (*0.08*)

*log\_odds:vratio -0.00*

(*0.28*)

*log\_odds:RR -0.02* (0.00)

vratio:RR 0.00

(0.00)

log\_odds:vratio:RR 0.01

(0.00)

————— R<sup>2</sup> 0.04 0.39 0.50

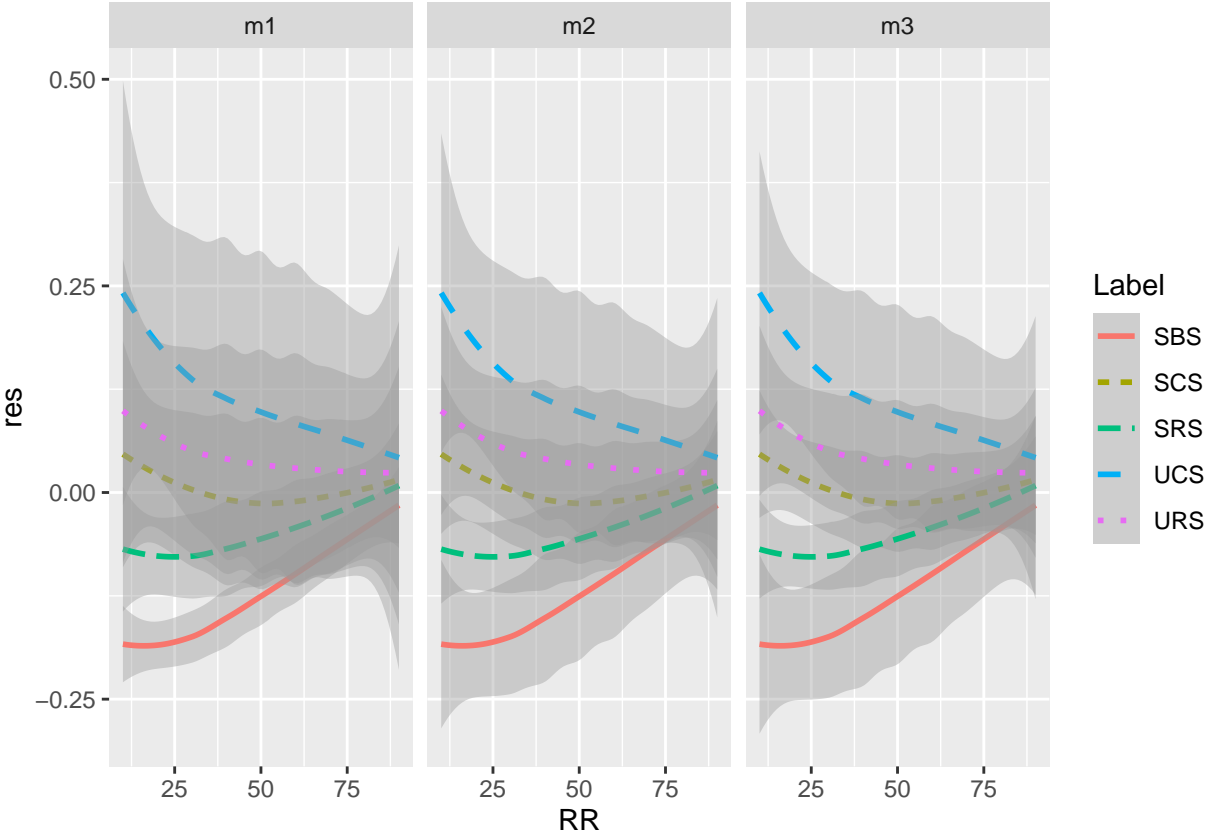
Adj. R<sup>2</sup> 0.04 0.38 0.49

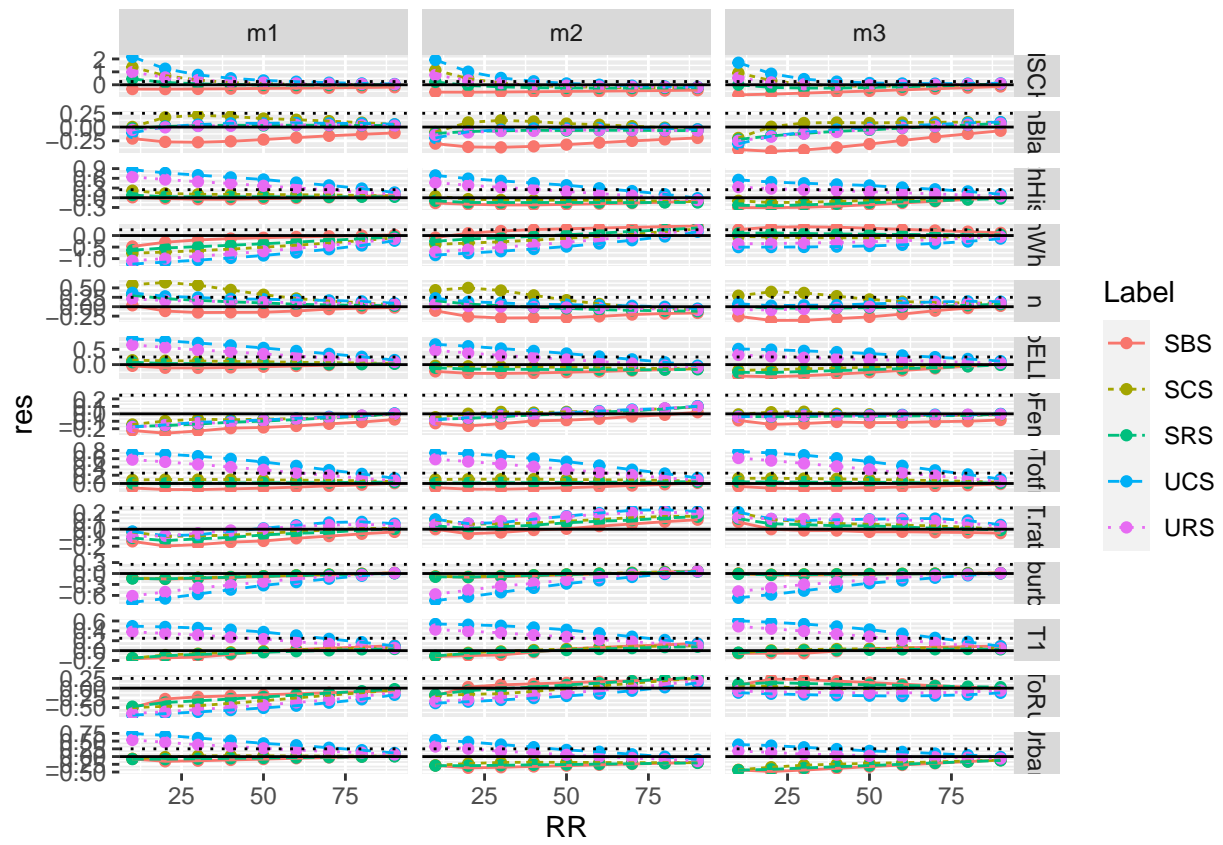
Num. obs. 585 585 585

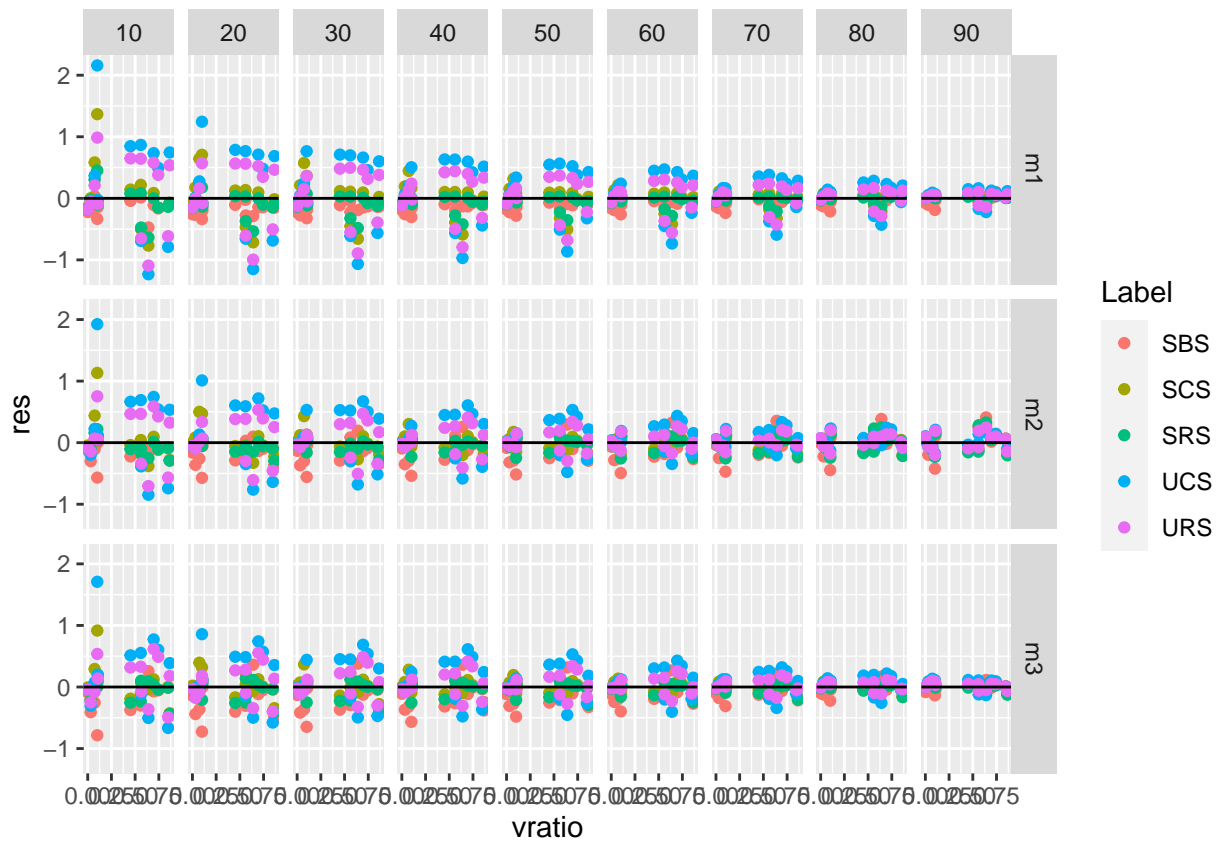
=====

\*\*\* p < 0.001; \*\* p < 0.01; \* p < 0.05

## 'geom\_smooth()' using method = 'loess' and formula 'y ~ x'





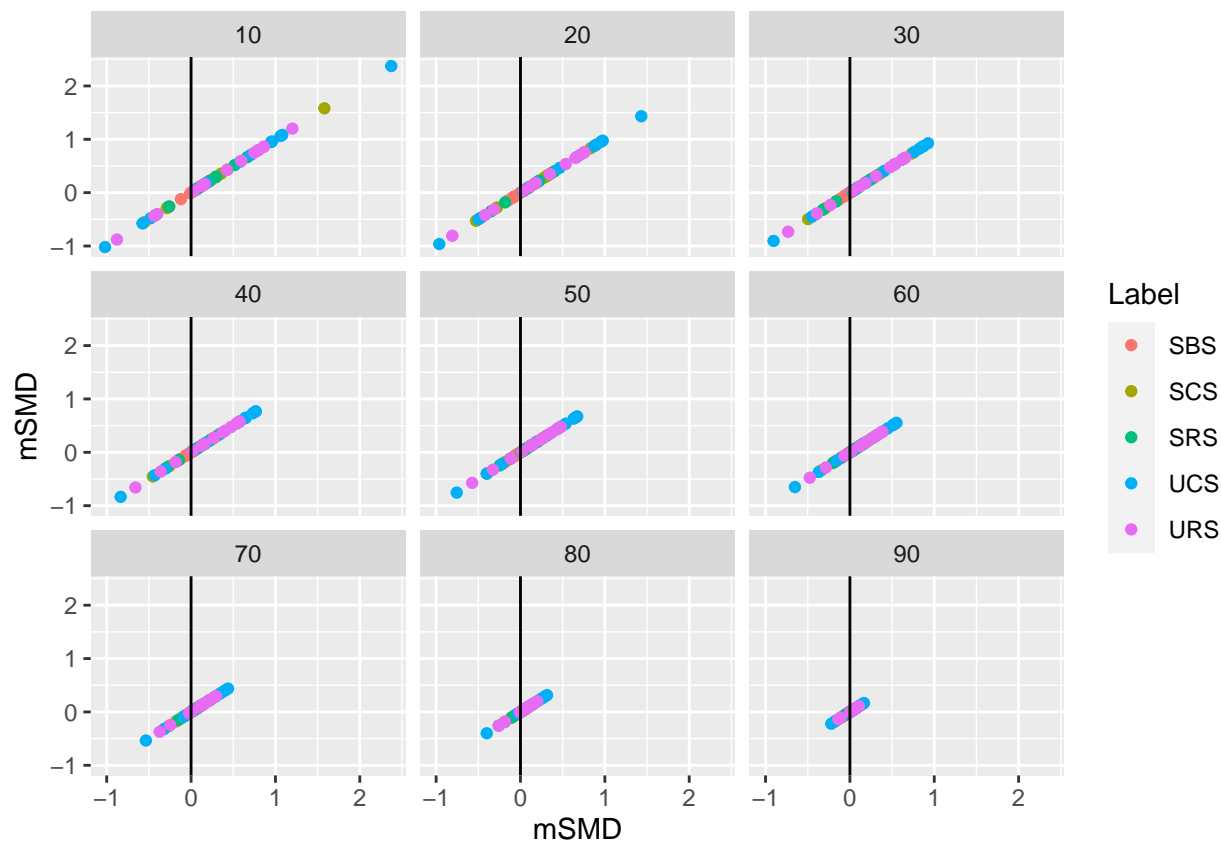


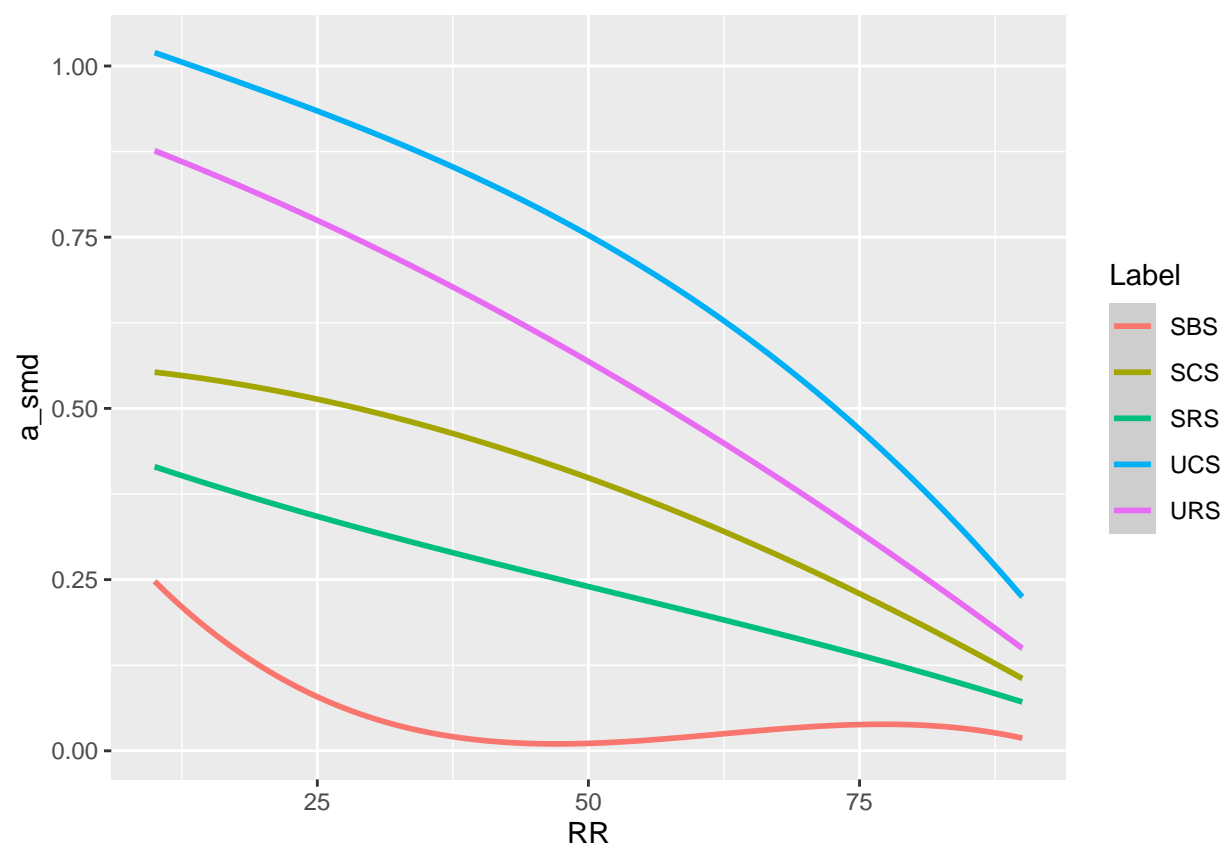
```
### Test 2
```

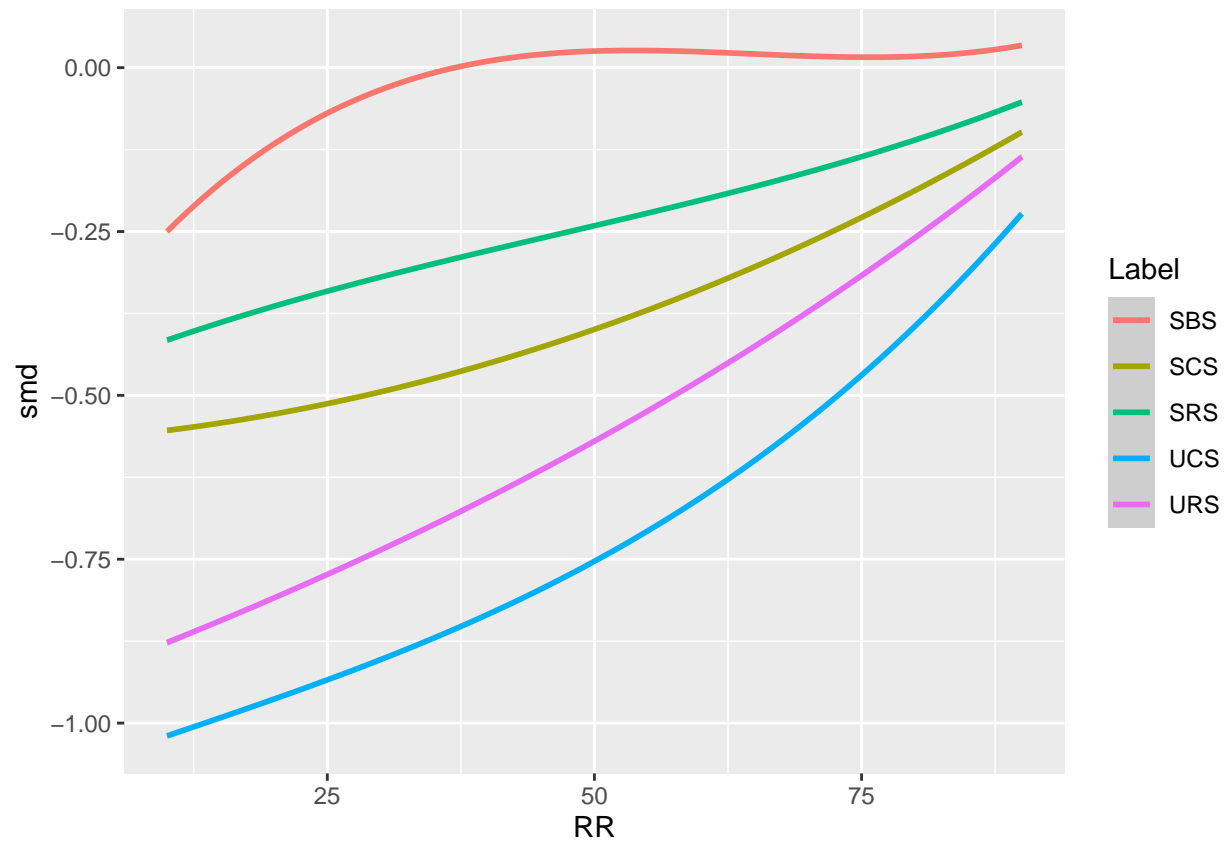
```
## Joining, by = "var"
```

```
## 'summarise()' regrouping output by 'sample_method', 'RR', 'var' (override with '.group
```









## Feasibility

### Sampling Difficulty.

```
## 'summarise()' regrouping output by 'sample_method', 'RR', 'measure', 'Label' (override)
```

```
## 'summarise()' ungrouping output (override with '.groups' argument)
```

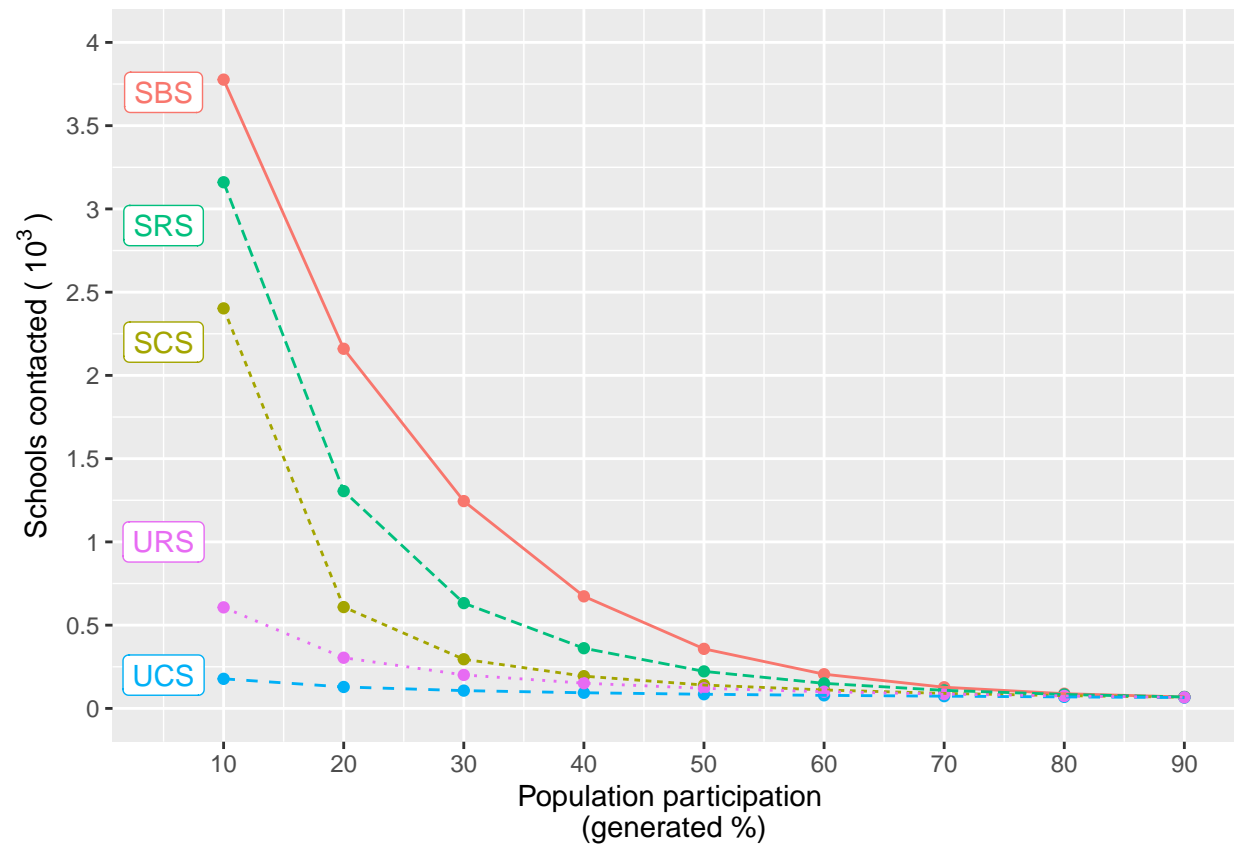
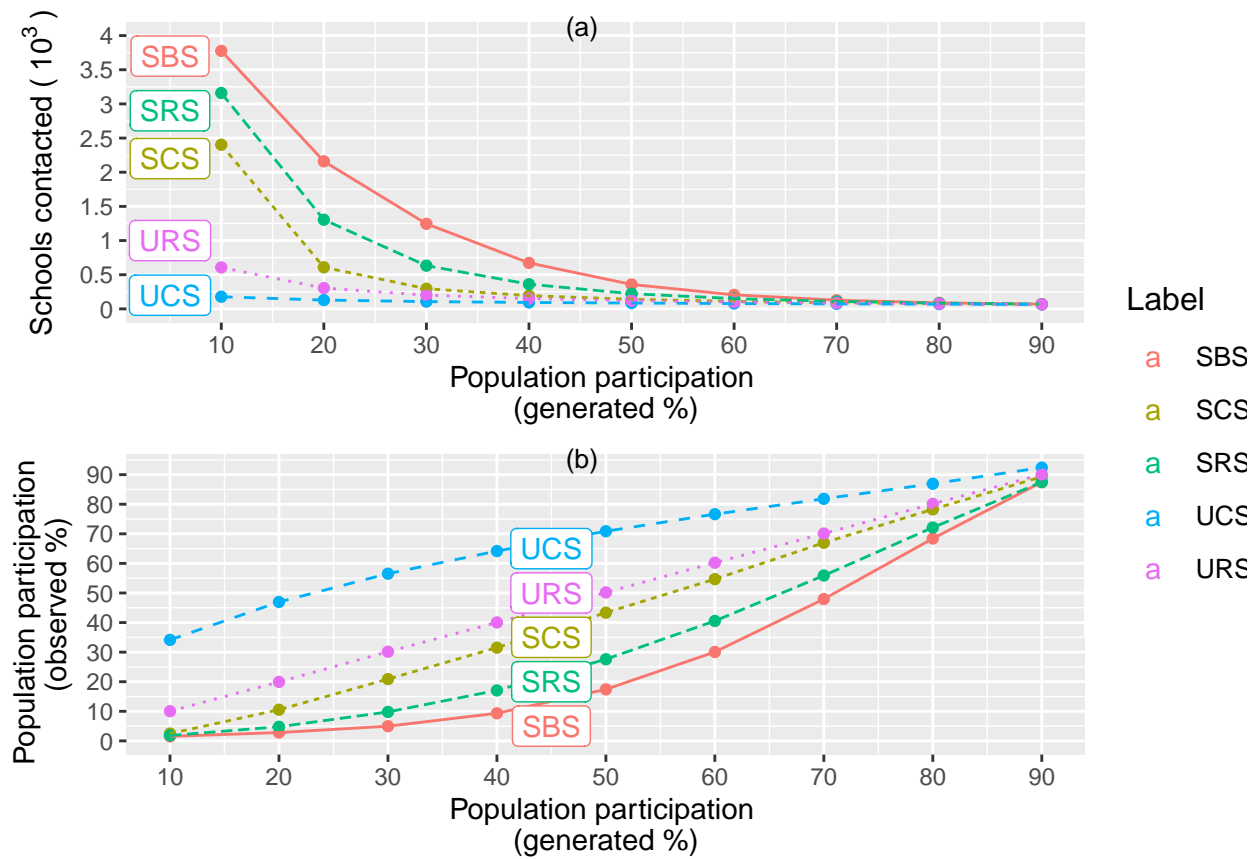


Figure 3. Schools Contacted



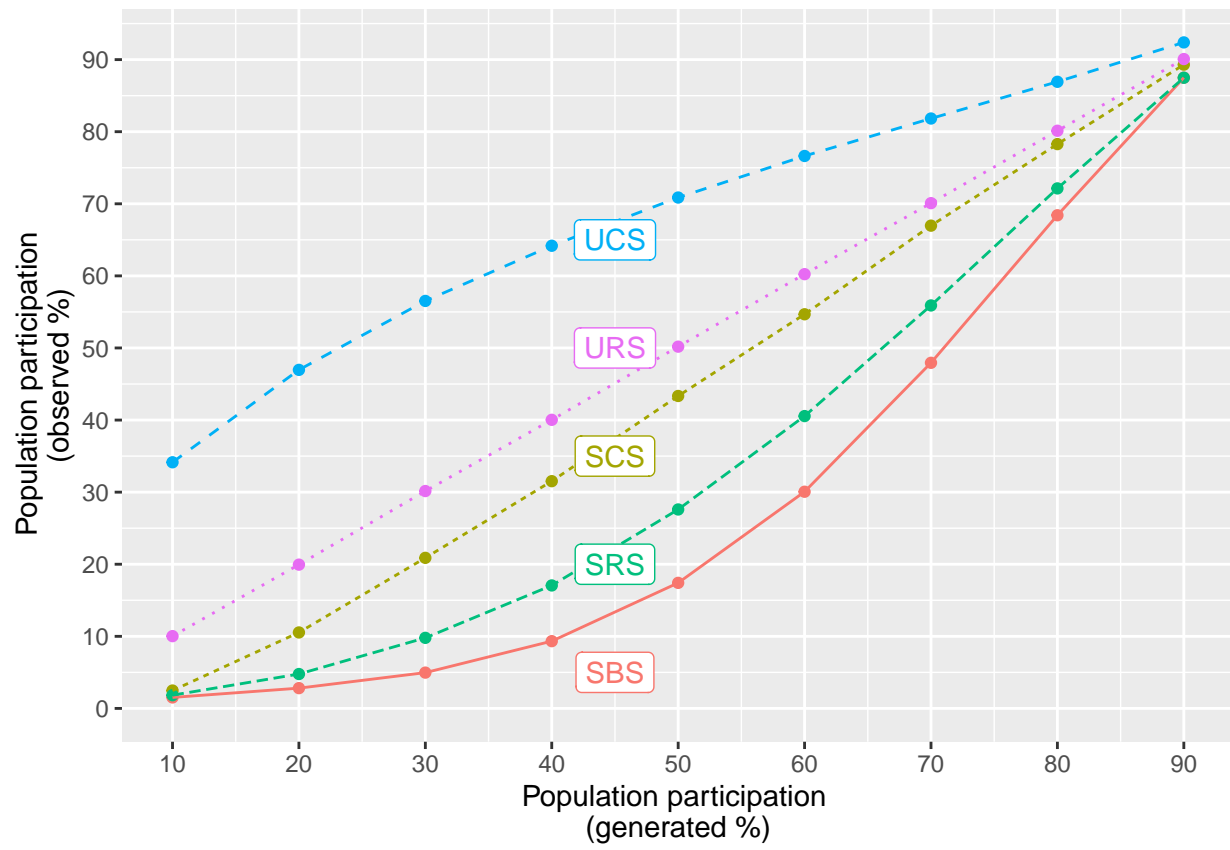


Figure 4. Sampling response rates

### Relative Performance.

```
## Joining, by = c("RR", "measure", "K")
```



[illegible]

```
## Joining, by = "DSID"
```

```
## Joining, by = "DSID"
```

```
## Joining, by = "DSID"
```

```
## Joining, by = "DSID"
```

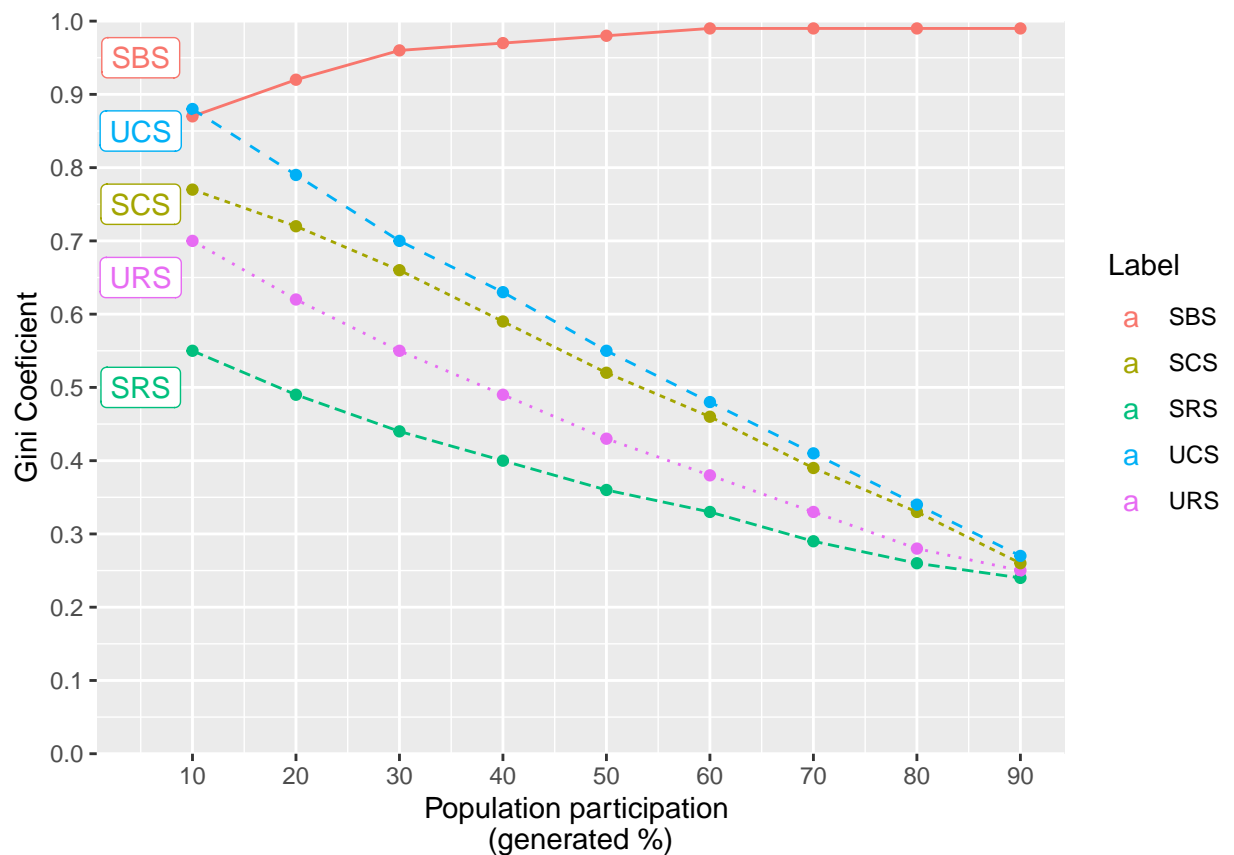
```
## Joining, by = "DSID"
```

```
## Joining, by = "DSID"
```

```
## Warning: 'cols' is now required when using unnest().
```

```
## Please use 'cols = c(data)'
```

```
## 'summarise()' regrouping output by 'sample_method', 'Label' (override with '.groups'
```





Export Plots

APPAM PLOTS

B-index

Sample

Non-Response

Combined

Individual

## Scale for 'x' is already present. Adding another scale for 'x', which will  
## replace the existing scale.

