

The cbcTools Package

Tools for Designing and Testing
Choice-Based Conjoint Surveys in 

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Designing a Choice-Based Conjoint Survey is Hard

Design Parameters

- What are my attributes and levels?
- Sample size (# respondents)
- Choice questions per respondent
- Alternative per choice question
- Labeled or unlabeled design?

Designing a Choice-Based Conjoint Survey is Hard

Design Parameters

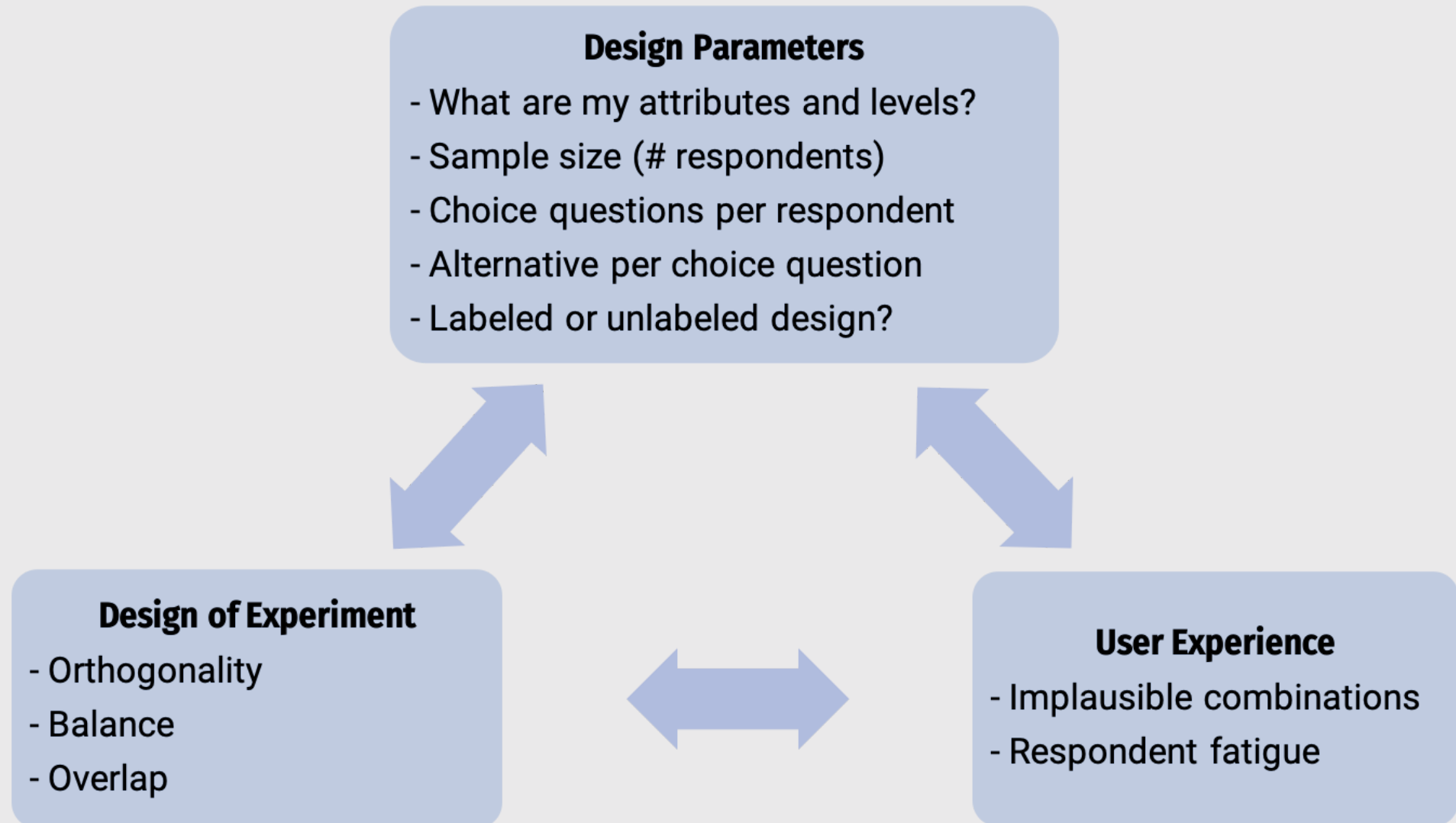
- What are my attributes and levels?
- Sample size (# respondents)
- Choice questions per respondent
- Alternative per choice question
- Labeled or unlabeled design?



Design of Experiment

- Orthogonality
- Balance
- Overlap

Designing a Choice-Based Conjoint Survey is Hard



A simple conjoint experiment about *cars*

Attribute	Levels
Brand	GM, BMW, Ferrari
Price	\$20k, \$40k, \$100k

Design: 9 choice sets, 3 alternatives each

Attribute counts:

brand:

GM	BMW	Ferrari
10	11	6

price:

20k	40k	100k
9	9	9

Pairwise attribute counts:

brand & price:

	20k	40k	100k
GM	3	0	7
BMW	4	5	2
Ferrari	2	4	0

A simple conjoint experiment about *cars*

Attribute	Levels
Brand	GM, BMW, Ferrari
Price	\$20k, \$40k, \$100k

Design: 90 choice sets, 3 alternatives each

Attribute counts:

brand:

GM	BMW	Ferrari
92	80	98

price:

20k	40k	100k
91	84	95

Pairwise attribute counts:

brand & price:

	20k	40k	100k
GM	31	31	30
BMW	25	25	30
Ferrari	35	28	35

Bayesian D-efficient designs

Maximize information on "Main Effects" according to priors

Attribute	Levels	Prior
Brand	GM, BMW, Ferrari	0, 1, 2
Price	\$20k, \$40k, \$100k	0, -1, -4

Attribute counts:

brand:

GM	BMW	Ferrari
93	90	86

price:

20k	40k	100k
97	93	78

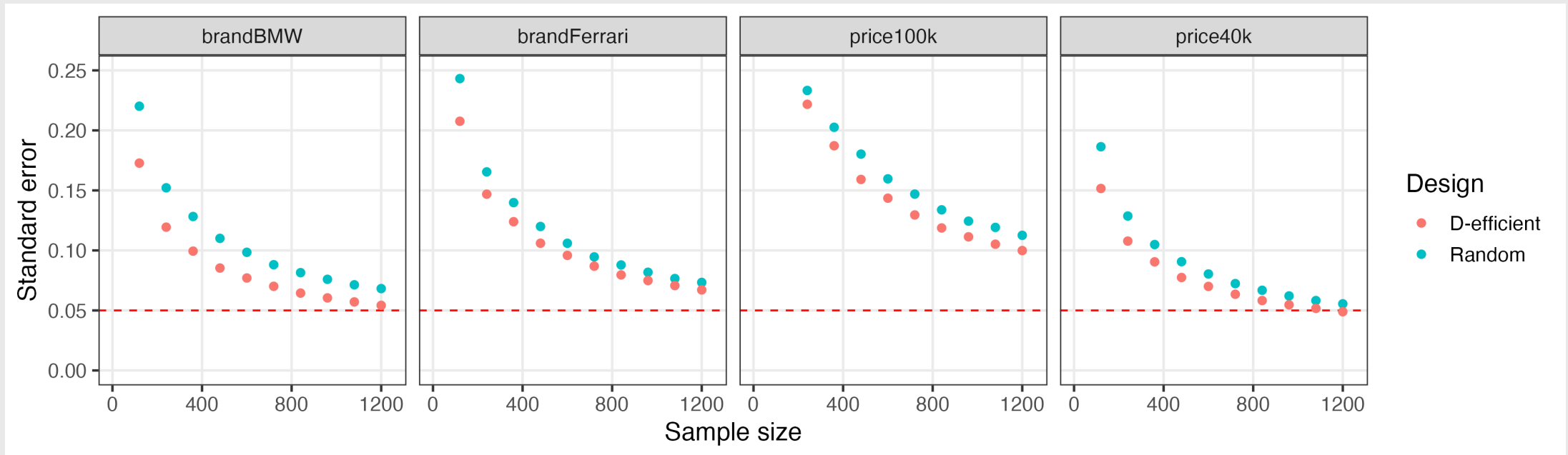
Pairwise attribute counts:

brand & price:

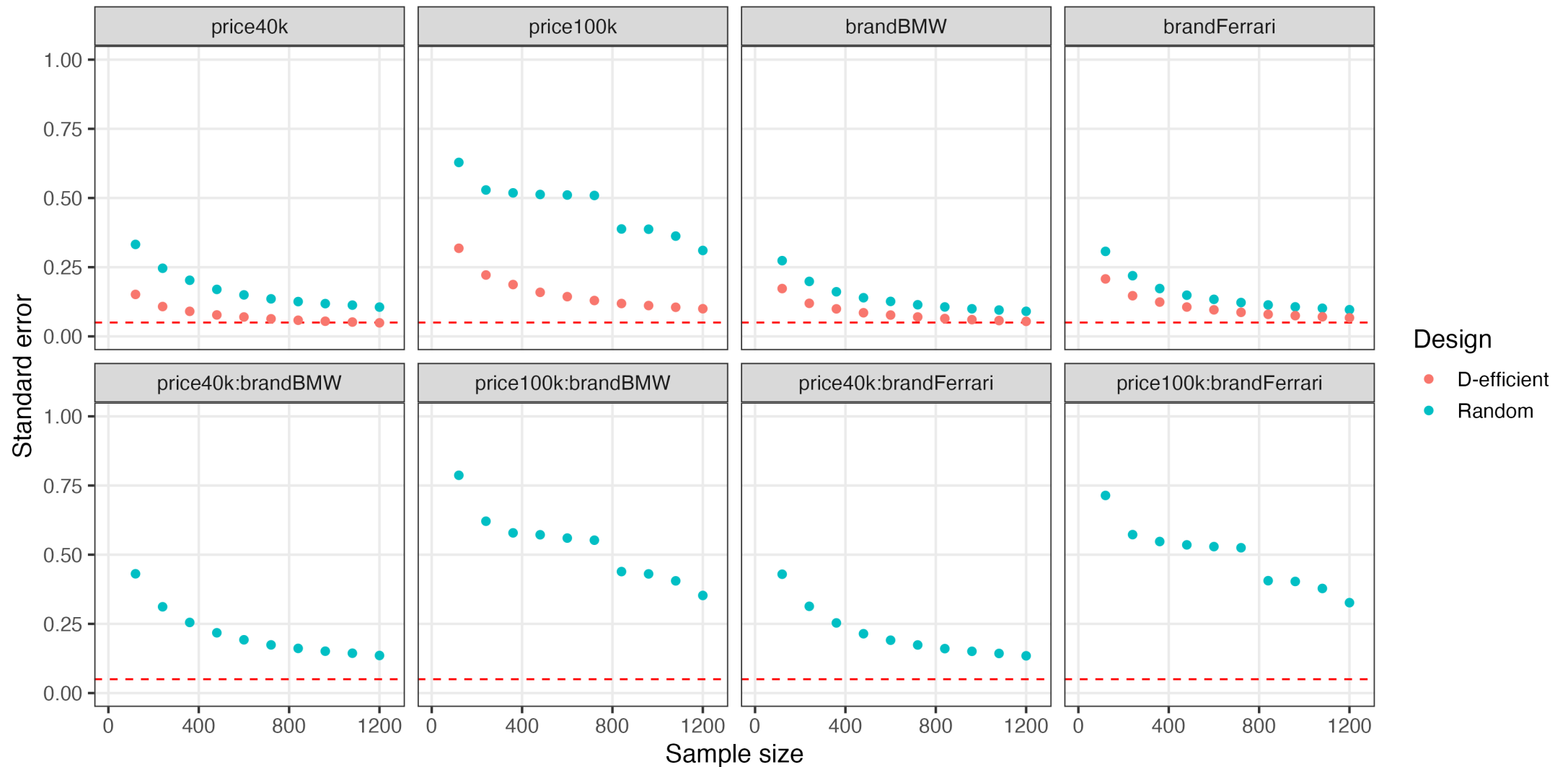
	20k	40k	100k
GM	52	41	0
BMW	30	30	30
Ferrari	15	22	49

Bayesian D-efficient designs

Attempts to maximize information on **Main Effects**



...but **interaction effects** are confounded in D-efficient designs



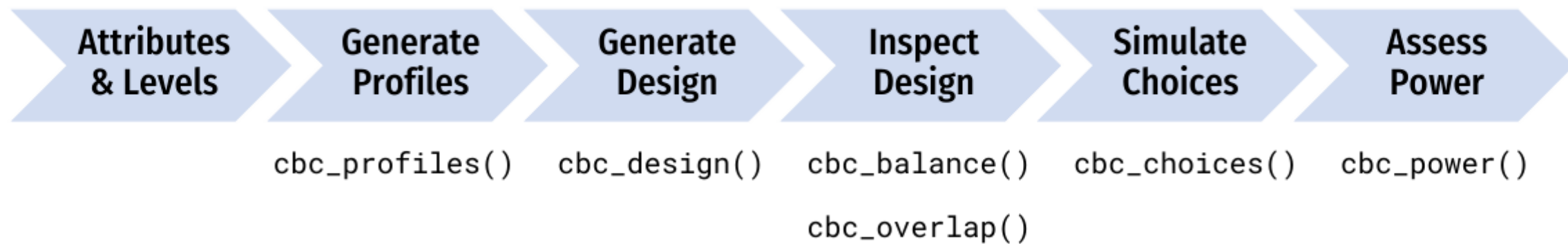
But what about other factors?

- What if I add one more choice question to each respondent?
- What if I increase the number of alternatives per choice question?
- What if I use a labeled design (aka "alternative-specific design")?
- What if there are interaction effects?

The cbcTools Package







Attribu
& Level

```
1 library(cbcTools)
```

```
2
```

```
3 cbc_|
```

◆ cbc_balance	{cbcTools}
◆ cbc_choices	{cbcTools}
◆ cbc_design	{cbcTools}
◆ cbc_overlap	{cbcTools}
◆ cbc_power	{cbcTools}
◆ cbc_profiles	{cbcTools}

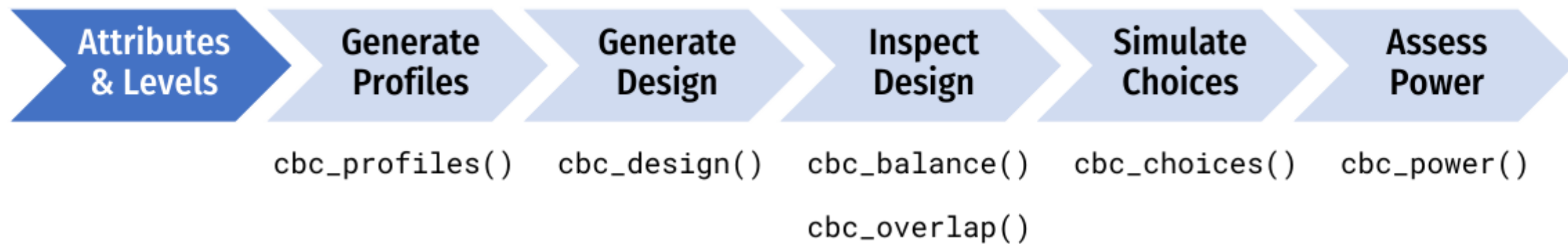
cbc_balance(design, atts = NULL)

This function prints out a summary of the counts of each level for each attribute across all choice questions as well as the two-way counts across all pairs of attributes for a given design.

Press F1 for additional help

Assess
Power

_power()

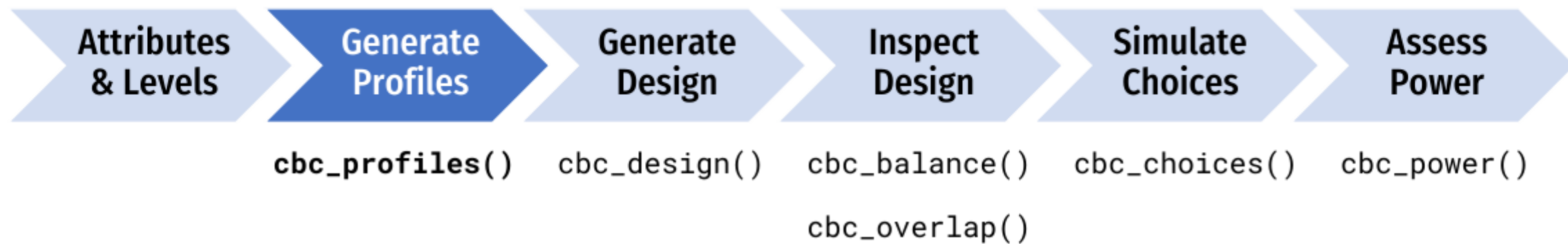


Define the attributes and levels

```
levels <- list(  
  price      = c(1.00, 1.50, 2.00, 2.50, 3.00, 3.50, 4.00), # $ per pound  
  type       = c("Fuji", "Gala", "Honeycrisp"),  
  freshness  = c("Excellent", "Average", "Poor")  
)
```

```
levels
```

```
#> $price  
#> [1] 1.0 1.5 2.0 2.5 3.0 3.5 4.0  
#>  
#> $type  
#> [1] "Fuji"      "Gala"      "Honeycrisp"  
#>  
#> $freshness  
#> [1] "Excellent" "Average"   "Poor"
```



Generate all possible profiles

```
profiles <- cbc_profiles(levels)
```

```
head(profiles)
```

```
#>   profileID price type freshness
#> 1         1  1.0 Fuji  Excellent
#> 2         2  1.5 Fuji  Excellent
#> 3         3  2.0 Fuji  Excellent
#> 4         4  2.5 Fuji  Excellent
#> 5         5  3.0 Fuji  Excellent
#> 6         6  3.5 Fuji  Excellent
```

```
tail(profiles)
```

```
#>   profileID price      type freshness
#> 58         58  1.5 Honeycrisp    Poor
#> 59         59  2.0 Honeycrisp    Poor
#> 60         60  2.5 Honeycrisp    Poor
#> 61         61  3.0 Honeycrisp    Poor
#> 62         62  3.5 Honeycrisp    Poor
#> 63         63  4.0 Honeycrisp    Poor
```

Attribute-specific levels

```
levels <- list(  
  price = c(1.00, 1.50, 2.00, 2.50, 3.00, 3.50, 4.00),  
  freshness = c("Excellent", "Average", "Poor"),  
  type = list(  
    "Fuji" = list(  
      price = c(2.00, 2.50, 3.00)  
    ),  
    "Gala" = list(  
      price = c(1.00, 1.50, 2.00)  
    ),  
    "Honeycrisp" = list(  
      price = c(2.50, 3.00, 3.50, 4.00),  
      freshness = c("Excellent", "Average")  
    )  
  )  
)
```

Generate restricted set of profiles

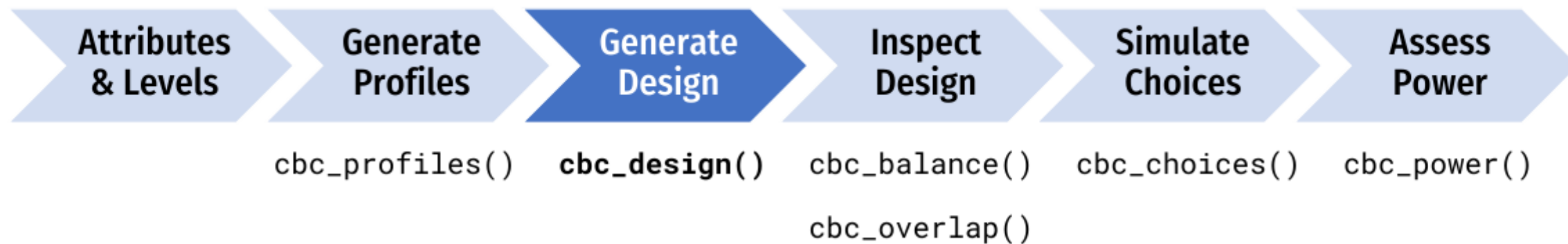
```
profiles <- cbc_profiles(levels)
```

```
head(profiles)
```

```
#>   profileID price freshness type
#> 1         1   2.0  Excellent Fuji
#> 2         2   2.5  Excellent Fuji
#> 3         3   3.0  Excellent Fuji
#> 4         4   2.0    Average Fuji
#> 5         5   2.5    Average Fuji
#> 6         6   3.0    Average Fuji
```

```
tail(profiles)
```

```
#>   profileID price freshness      type
#> 21         21   3.5  Excellent Honeycrisp
#> 22         22   4.0  Excellent Honeycrisp
#> 23         23   2.5    Average Honeycrisp
#> 24         24   3.0    Average Honeycrisp
#> 25         25   3.5    Average Honeycrisp
#> 26         26   4.0    Average Honeycrisp
```



Generate a survey design

```
design <- cbc_design(  
  profiles = profiles,  
  n_resp   = 300, # Number of respondents  
  n_alts   = 3,   # Number of alternatives per question  
  n_q      = 6    # Number of questions per respondent  
)
```

```
head(design)
```

```
#>   respID qID altID obsID profileID price      type freshness  
#> 1      1  1    1      1        16   1.5 Honeycrisp Excellent  
#> 2      1  1    2      1        23   1.5      Fuji    Average  
#> 3      1  1    3      1        44   1.5      Fuji      Poor  
#> 4      1  2    1      2        18   2.5 Honeycrisp Excellent  
#> 5      1  2    2      2        14   4.0      Gala    Excellent  
#> 6      1  2    3      2         8   1.0      Gala    Excellent
```


Include a "no choice" option

```
design <- cbc_design(  
  profiles = profiles,  
  n_resp   = 300, # Number of respondents  
  n_alts   = 3,   # Number of alternatives per question  
  n_q      = 6,   # Number of questions per respondent  
  no_choice = TRUE  
)
```

```
head(design)
```

```
#>      respID qID altID obsID profileID price type_Fuji type_Gala type_Honeycrisp freshn  
#> 1          1  1    1      1         37   1.5         0         0             1  
#> 2          1  1    2      1         50   1.0         0         1             0  
#> 3          1  1    3      1         21   4.0         0         0             1  
#> 11000      1  1    4      1          0   0.0         0         0             0  
#> 4          1  2    1      2          3   2.0         1         0             0  
#> 5          1  2    2      2         28   4.0         1         0             0
```

Make a labeled design

(aka "Alternative-specific design")

```
design <- cbc_design(  
  profiles = profiles,  
  n_resp   = 300, # Number of respondents  
  n_alts   = 3,   # Number of alternatives per question  
  n_q      = 6,   # Number of questions per respondent  
  label    = "type"  
)
```

```
head(design)
```

```
#>   respID qID altID obsID profileID price      type freshness  
#> 1      1  1    1      1         22  1.0      Fuji    Average  
#> 2      1  1    2      1         10  2.0      Gala    Excellent  
#> 3      1  1    3      1         58  1.5 Honeycrisp    Poor  
#> 4      1  2    1      2         49  4.0      Fuji    Poor  
#> 5      1  2    2      2         32  2.5      Gala    Average  
#> 6      1  2    3      2         59  2.0 Honeycrisp    Poor
```

Make a Bayesian D-efficient design

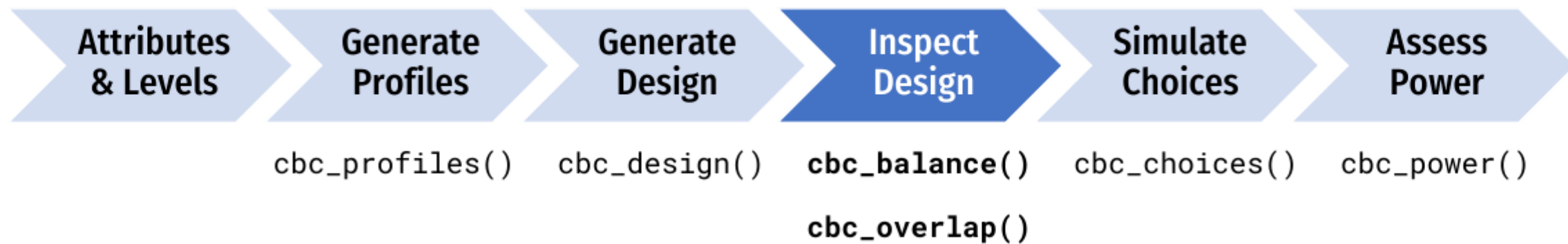
(coming soon!)

```
design <- cbc_design(  
  profiles = profiles,  
  n_resp   = 300, # Number of respondents  
  n_alts   = 3,   # Number of alternatives per question  
  n_q      = 6,   # Number of questions per respondent  
  priors = list(  
    price      = -0.1,  
    type       = c(0.1, 0.2),  
    freshness  = c(0.1, -0.2)  
  )  
)
```

Make a Bayesian D-efficient design

(coming soon!)

- Check out the `idefix` package
- Import a design: `Sawtooth` →  → 



Check design **balance**

```
cbc_balance(design)
```

Attribute counts:

price:

	1	1.5	2	2.5	3	3.5	4
	825	797	743	743	767	779	746

type:

	Fuji	Gala	Honeycrisp
	1842	1769	1789

freshness:

	Excellent	Average	Poor
	1813	1775	1812

Pairwise attribute counts:

price & type:

	Fuji	Gala	Honeycrisp
1	304	252	269
1.5	274	251	272
2	257	254	232
2.5	240	254	249
3	249	263	255
3.5	257	250	272
4	261	245	240

Check design **overlap**

```
cbc_overlap(design)
```

```
Counts of attribute overlap:  
(# of questions with N unique levels)
```

```
price:
```

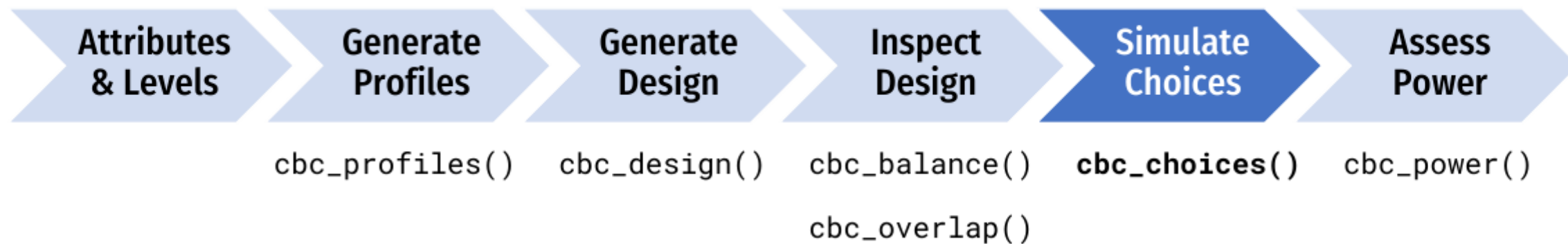
1	2	3
31	630	1139

```
type:
```

1	2	3
156	1248	396

```
freshness:
```

1	2	3
175	1189	436



Simulate random choices

```
data <- cbc_choices(  
  design = design,  
  obsID   = "obsID"  
)
```

```
head(data)
```

```
#>   respID qID altID obsID profileID price      type freshness choice  
#> 1      1  1    1      1        22   1.0      Fuji   Average      1  
#> 2      1  1    2      1        10   2.0      Gala  Excellent      0  
#> 3      1  1    3      1        58   1.5 Honeycrisp    Poor      0  
#> 4      1  2    1      2        49   4.0      Fuji    Poor      0  
#> 5      1  2    2      2        32   2.5      Gala   Average      1  
#> 6      1  2    3      2        59   2.0 Honeycrisp    Poor      0
```

Simulate choices according to a prior

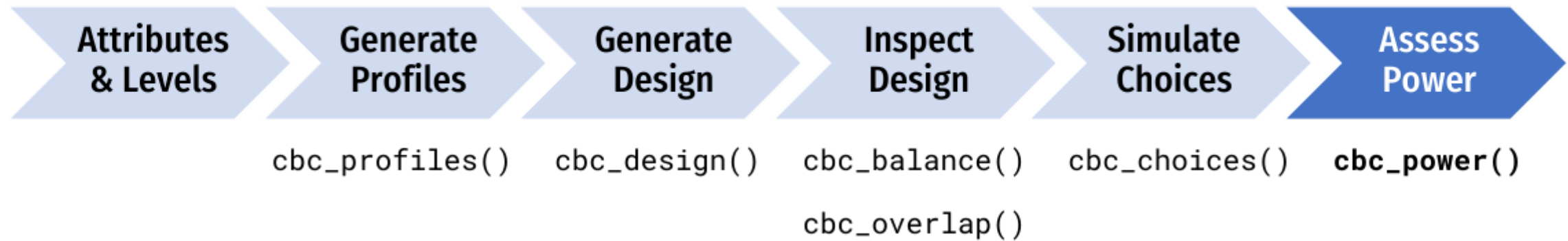
```
data <- cbc_choices(  
  design = design,  
  obsID = "obsID",  
  priors = list(  
    price      = -0.1,  
    type       = c(0.1, 0.2),  
    freshness  = c(0.1, -0.2)  
  )  
)
```

Attribute	Level	Utility
Price	Continuous	-0.1
Type	Fuji	0
	Gala	0.1
	Honeycrisp	0.2
Freshness	Average	0
	Excellent	0.1
	Poor	-0.2

Simulate choices according to a prior

```
data <- cbc_choices(  
  design = design,  
  obsID = "obsID",  
  priors = list(  
    price = -0.1,  
    type = randN(  
      mu = c(0.1, 0.2),  
      sigma = c(0.5, 1)  
    ),  
    freshness = c(0.1, -0.2)  
  )  
)
```

Attribute	Level	Utility
Price	Continuous	-0.1
Type	Fuji	0
	Gala	N(0.1, 0.5)
	Honeycrisp	N(0.2, 1)
Freshness	Average	0
	Excellent	0.1
	Poor	-0.2



Conduct a power analysis

```
power <- cbc_power(  
  nbreaks = 10,  
  n_q     = 3,  
  data    = data,  
  pars    = c("price", "type", "freshness"),  
  outcome = "choice",  
  obsID   = "obsID"  
)
```

head(power)

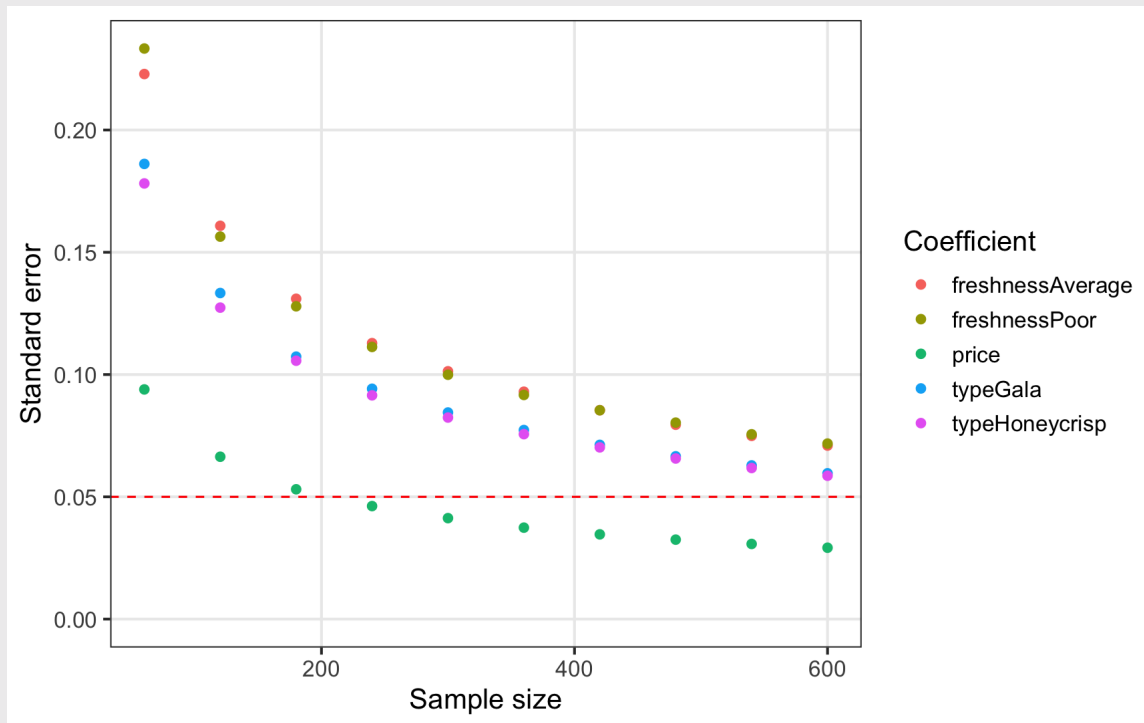
```
#>   sampleSize      coef      es  
#> 1         60      price -0.00751708  
#> 2         60    typeGala -0.19045129  
#> 3         60 typeHoneycrisp -0.02352382  
#> 4         60 freshnessAverage -0.05873052  
#> 5         60  freshnessPoor -0.25428352  
#> 6        120      price -0.07826151
```

tail(power)

```
#>   sampleSize      coef      es  
#> 45         540  freshnessPoor -0.1263108  
#> 46         600      price -0.0854275  
#> 47         600    typeGala  0.1563946  
#> 48         600 typeHoneycrisp  0.2336239  
#> 49         600 freshnessAverage  0.0884197  
#> 50         600  freshnessPoor -0.1156453
```

Conduct a power analysis

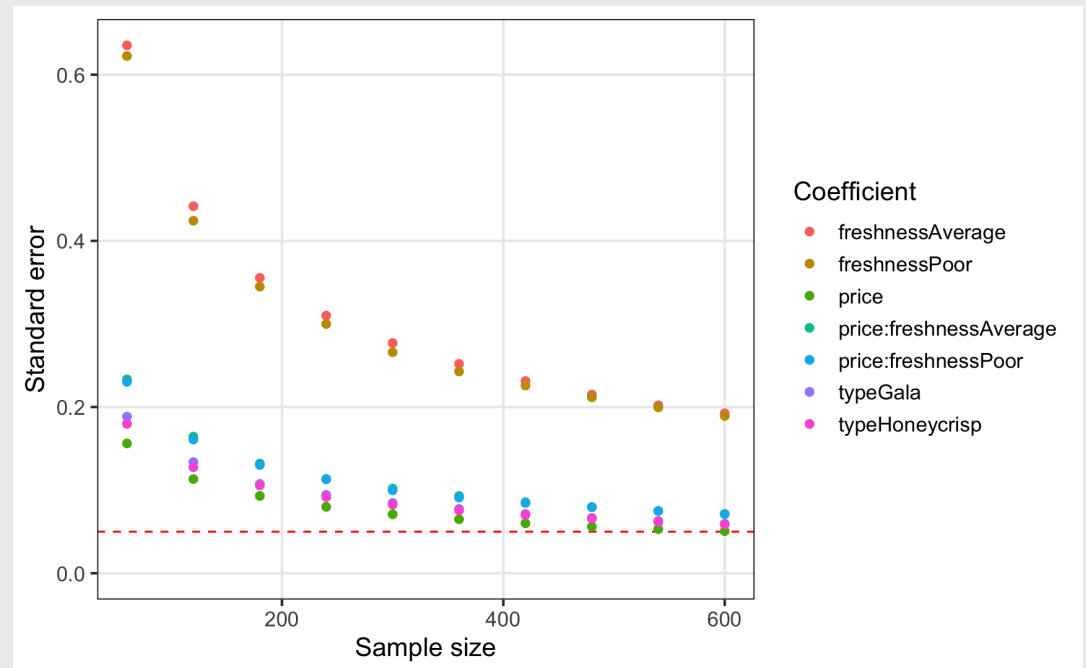
```
plot(power)
```



Conduct a power analysis

```
power_int <- cbc_power(  
  nbreaks = 10,  
  n_q     = 3,  
  data    = data,  
  pars    = c(  
    "price",  
    "type",  
    "freshness",  
    "price*freshness"  
  ),  
  outcome = "choice",  
  obsID   = "obsID"  
)
```

```
plot(power_int)
```





**Attributes
& Levels**

**Generate
Profiles**

**Generate
Design**

**Inspect
Design**

**Simulate
Choices**

**Assess
Power**

`cbc_profiles()`

`cbc_design()`

`cbc_balance()`

`cbc_choices()`

`cbc_power()`

`cbc_overlap()`



**Attributes
& Levels**

**Generate
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`cbc_profiles()`

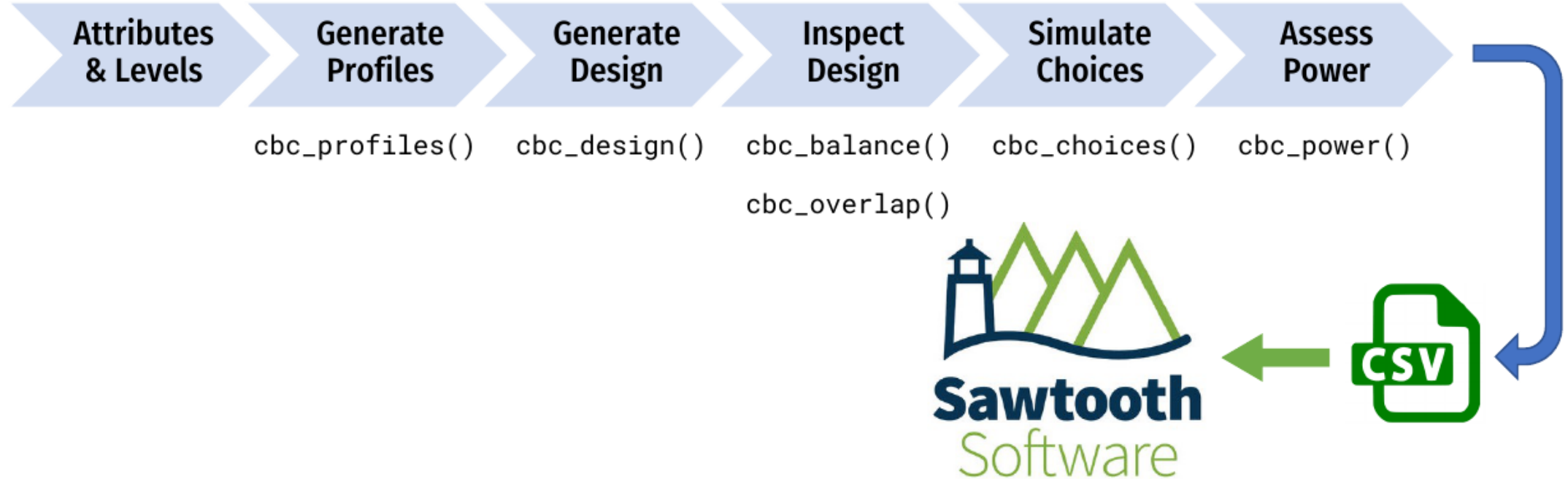
`cbc_design()`

`cbc_balance()`

`cbc_choices()`

`cbc_power()`

`cbc_overlap()`



Thanks!

cbcTools documentation: <https://jhelvy.github.io/cbcTools/>

Slides: <https://jhelvy.github.io/2022-sawtooth-conf>

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Extra slides