



skpr

R Package for Generation and Evaluation of
Experimental Designs

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Outline

1. Demo
2. Introduction
3. Features of skpr
4. skprGUI

Live Demo

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4. skprGUI

Motivation for skpr

- Create an open-source design of experiments package to allow for the creation and analysis of designs **all within R**.
- Implement optimal design algorithms using modern linear algebra libraries to make the underlying code **simple, reliable, and auditable**.
- Integrate Monte Carlo power evaluation as a standard feature and make it **flexible, extensible, and easy to use**.

Preparation matching execution

Plan your experiment using the same models and tools that you plan to use when analyzing them.

Shareable and repeatable

With skpr, there is never a question about *how* you ran an analysis—it's all there in the script.

Reproducing your work is as simple as re-running the code.

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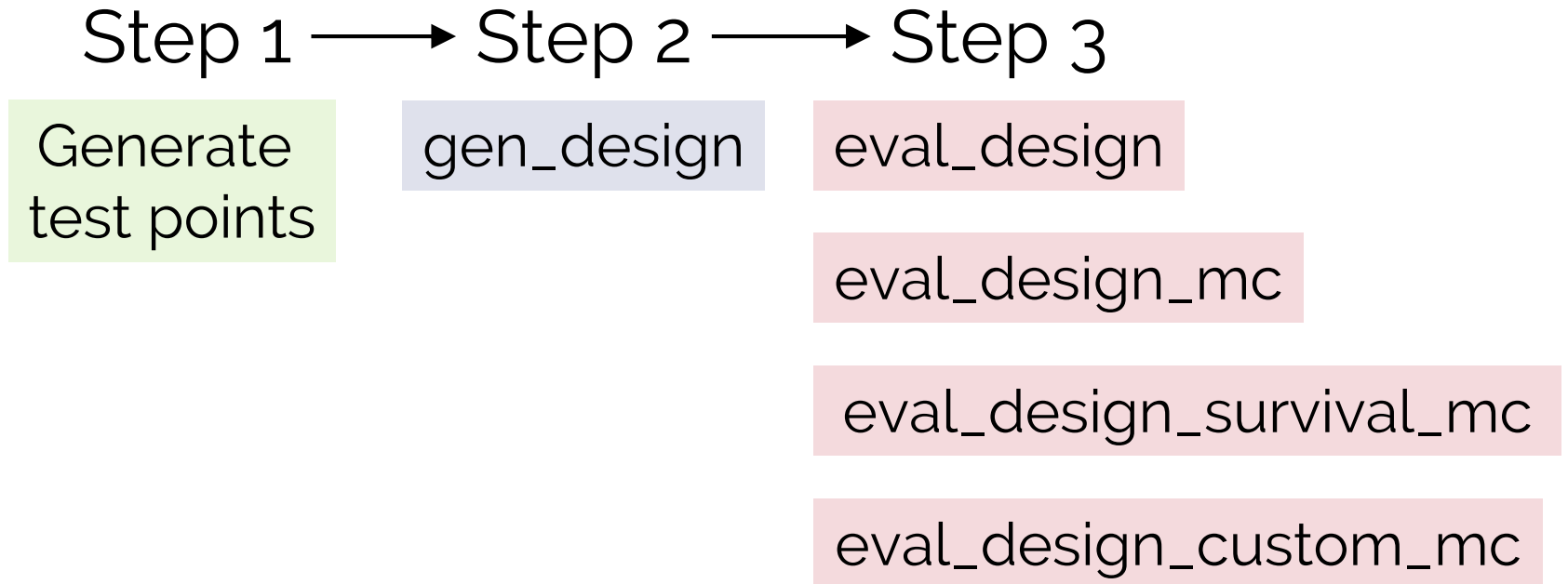
Five main functions + plotting

- **gen_design**: Optimal design generation
- **eval_design**: Parametric power evaluation
- **eval_design_mc**: Monte Carlo power evaluation
- **eval_design_survival_mc**: Censored Monte Carlo power evaluation
- **eval_design_custom_mc**: Custom library Monte Carlo power evaluation
- **plot_correlations & plot_fds**: Plot correlation map and fraction of design space plots

Five main functions + plotting

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Workflow in skpr



Example: How much caffeine extracted from coffee?

Factors:

Bean size: Large, Small

Bean type: Kona, Java

Water temperature: 80°C, 90°C, 100°C

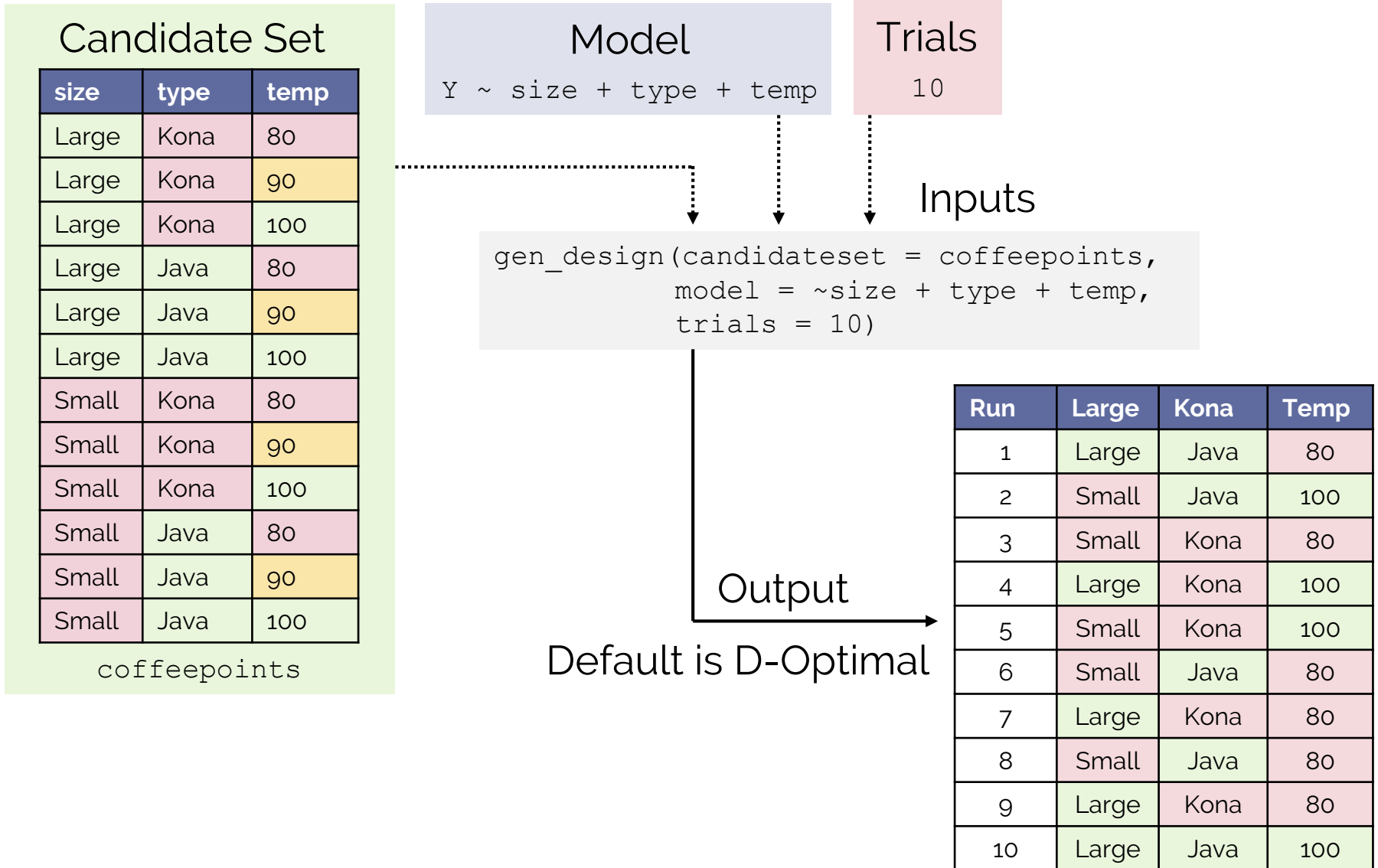
Roast Darkness (hard to change): Light, Medium, Dark



Main functions in skpr

- **gen_design**: Optimal design generation
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gen_design - Basic Functionality



Creation of the candidate set is simple

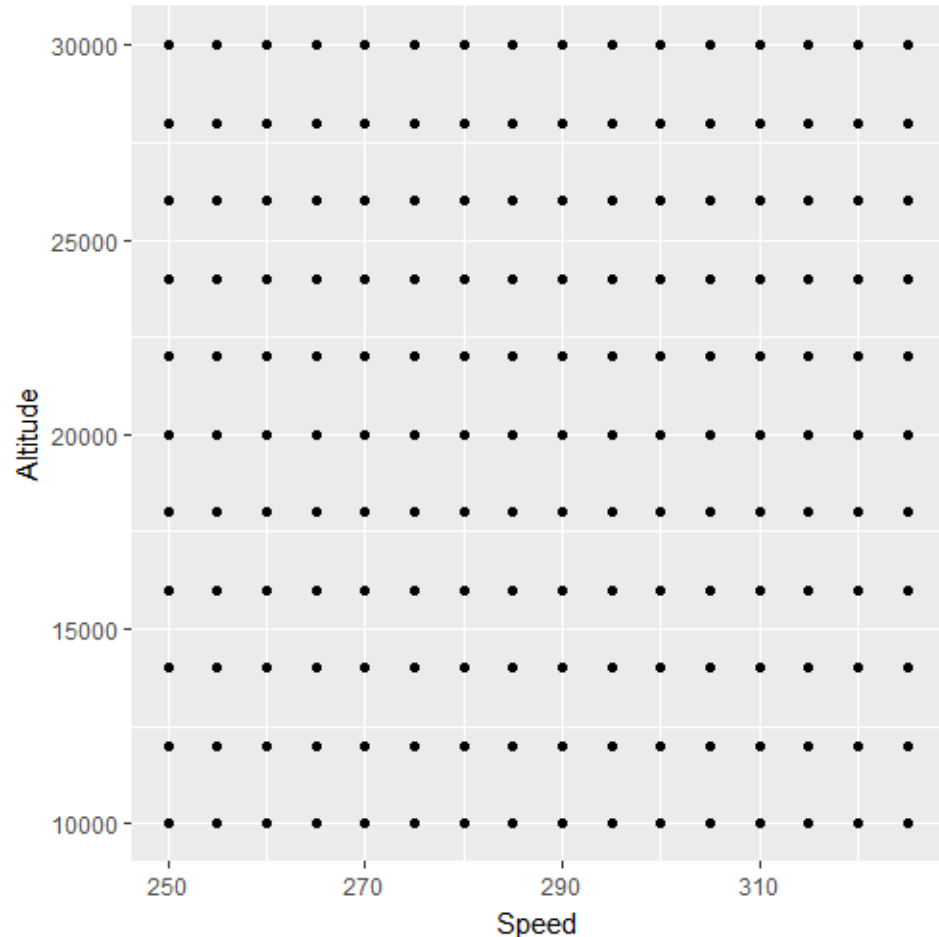
expand.grid function in R automatically generates all combinations of input parameters

```
> expand.grid(size=c("Large","Small"),type=c("Java","Kona"),temp=c(80,90,100))
  size type temp
1 Large Java  80
2 Small Java  80
3 Large Kona  80
4 Small Kona  80
5 Large Java  90
6 Small Java  90
7 Large Kona  90
8 Small Kona  90
9 Large Java 100
10 Small Java 100
11 Large Kona 100
12 Small Kona 100
```

Easily constrain your design space

Simple filtering of disallowed combinations to constrain the design space.

Example:
Testing within a flight envelope

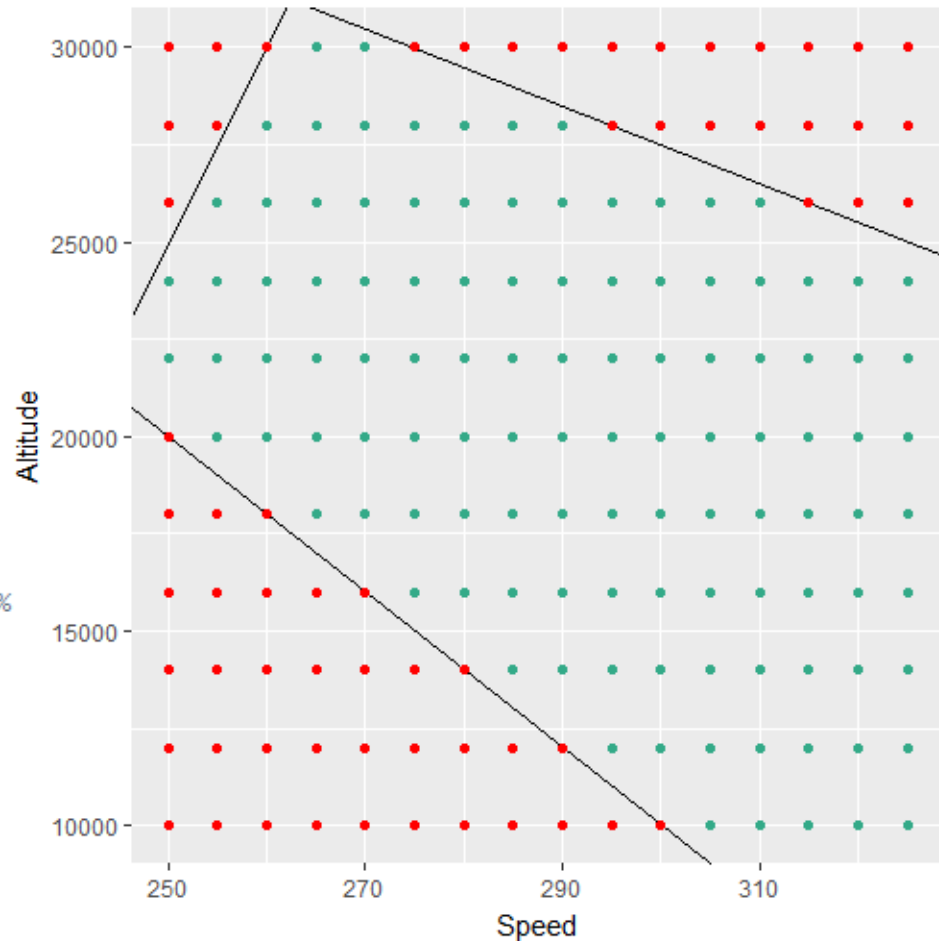


Easily constrain your design space

Simple filtering of disallowed combinations to constrain the design space.

Filter out unwanted test points

```
filteredcandidateset = candidateset %>%  
  filter(70000 - speed * 200 < altitude) %>%  
  filter(57500 - speed * 100 > altitude) %>%  
  filter(-100000 + speed * 500 > altitude)
```

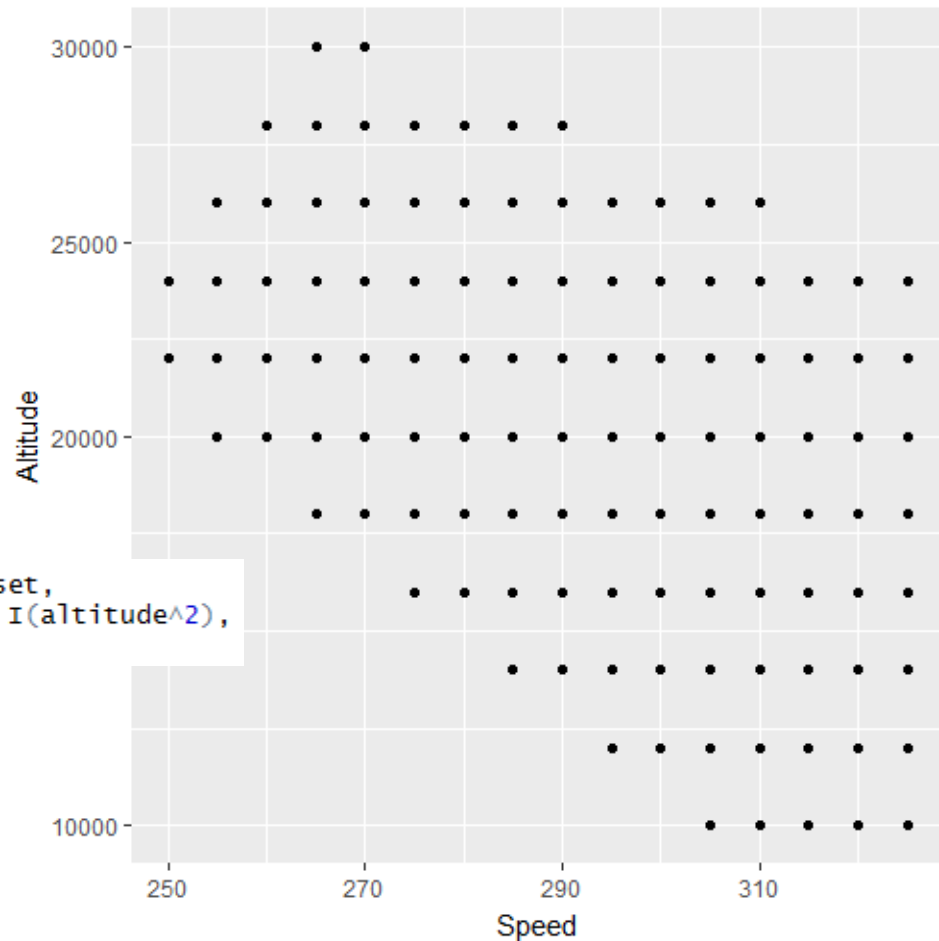


Easily constrain your design space

Simple filtering of disallowed combinations to constrain the design space.

Use new candidate set to generate design

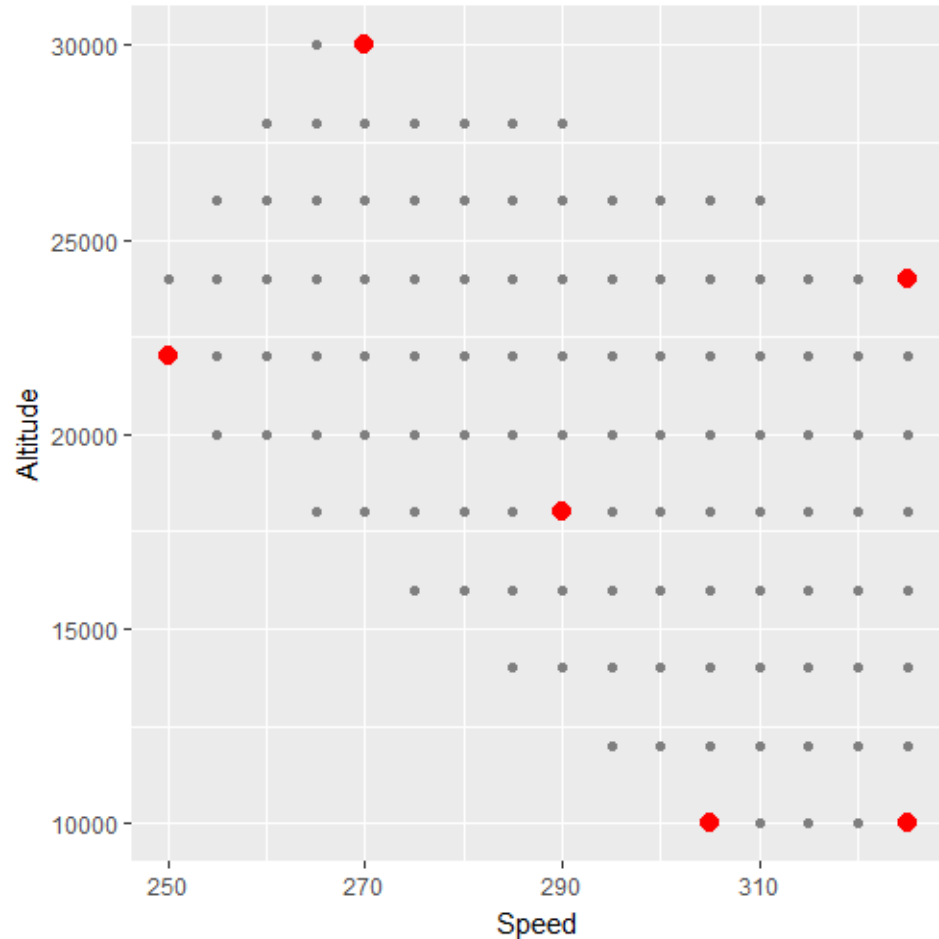
```
gen_design(candidateset = filteredcandidateset,  
           ~speed + altitude + I(speed^2) + I(altitude^2),  
           trials = 18)
```



Easily constrain your design space

Simple filtering of disallowed combinations to constrain the design space.

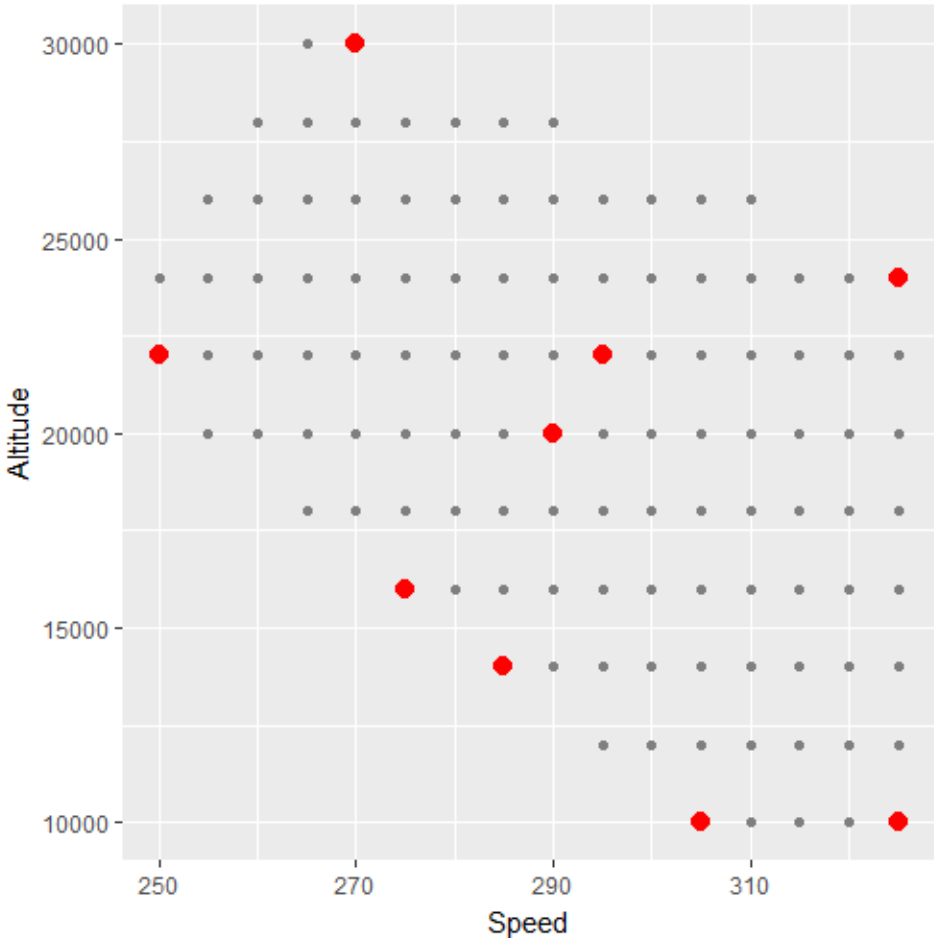
Use new candidate set to generate design



Easily constrain your design space

Simple filtering of disallowed combinations to constrain the design space.

Explore different options (e.g. I-optimal design)



Sequential construction of split-plot designs

```
candidatelist = expand.grid(size=as.factor(c("Large","Small")),  
                             type=as.factor(c("Kona","Java")),  
                             temp=c(80,100),  
                             roastdarkness=c("Light","Medium","Dark"))
```

```
gen_design(candidateset = candidatelist, ~roastdarkness, trials=6) -> htcdesign
```

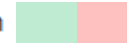
```
gen_design(candidateset = candidatelist, ~size+temp+type, trials=18,  
           splitplotdesign = htcdesign, splitplotsizes=3)
```

	roastdarkness	size	temp	type
1	Dark	Small	100	Java
2	Light	Large	80	Kona
3	Dark	Large	100	Kona
4	Medium	Small	80	Kona
5	Light	Large	100	Java
6	Medium	Large	100	Java

	roastdarkness	size	temp	type
1.1	Dark	Small	100	Java
1.2	Dark	Large	80	Kona
1.3	Dark	Large	100	Kona
2.1	Light	Small	80	Kona
2.2	Light	Large	100	Kona
2.3	Light	Large	100	Java
3.1	Dark	Small	80	Kona
3.2	Dark	Small	80	Java
3.3	Dark	Large	100	Java
4.1	Medium	Large	80	Java
4.2	Medium	Small	100	Kona
4.3	Medium	Small	80	Java
5.1	Light	Large	80	Kona
5.2	Light	Small	80	Java
5.3	Light	Small	100	Java
6.1	Medium	Large	80	Java
6.2	Medium	Large	100	Kona
6.3	Medium	Small	100	Kona

Supports N-depth split-plot designs

```
gen_design(candidateset = candidatelist, ~roastdarkness, trials=4) -> vhtcdesign
```



```
gen_design(candidateset = candidatelist, ~type, trials=8,  
  splitplotdesign = vhtcdesign, splitplotsizes = 2) -> htcdesign
```



```
gen_design(candidateset = candidatelist, ~size+temp, trials=24,  
  splitplotdesign = htcdesign, splitplotsizes = 3)
```

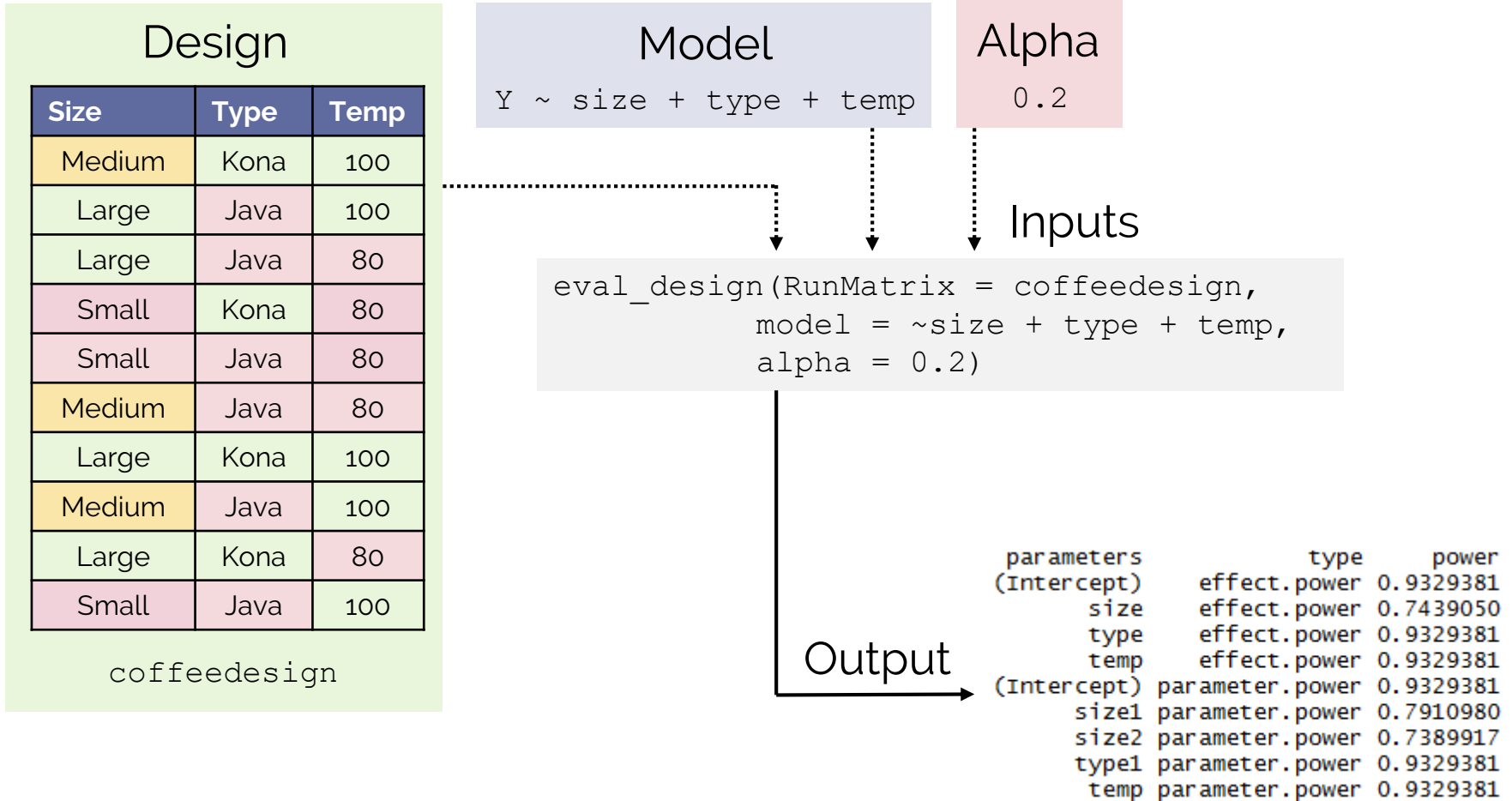


	roastdarkness	type	size	temp
1.1.1	Dark	Kona	Large	100
1.1.2	Dark	Kona	Small	80
1.1.3	Dark	Kona	Small	100
1.2.1	Dark	Java	Large	80
1.2.2	Dark	Java	Small	100
1.2.3	Dark	Java	Large	80
2.1.1	Medium	Java	Small	80
2.1.2	Medium	Java	Small	80
2.1.3	Medium	Java	Large	100
2.2.1	Medium	Kona	Small	100
2.2.2	Medium	Kona	Large	80
2.2.3	Medium	Kona	Large	100
3.1.1	Light	Kona	Small	80
3.1.2	Light	Kona	Large	80
3.1.3	Light	Kona	Large	100
3.2.1	Light	Java	Large	80
3.2.2	Light	Java	Small	100
3.2.3	Light	Java	Small	100
4.1.1	Light	Kona	Large	100
4.1.2	Light	Kona	Small	80
4.1.3	Light	Kona	Small	80
4.2.1	Light	Java	Small	100
4.2.2	Light	Java	Large	80
4.2.3	Light	Java	Large	100

Main functions in skpr

- **gen_design**: Optimal design generation
- **eval_design**: Parametric power evaluation
- **eval_design_mc**: Monte Carlo power evaluation

eval_design - Basic Functionality



Change effect size

Design

Size	Type	Temp
Medium	Kona	100
Large	Java	100
Large	Java	80
Small	Kona	80
Small	Java	80
Medium	Java	80
Large	Kona	100
Medium	Java	100
Large	Kona	80
Small	Java	100

coffeedesign

Model
 $Y \sim \text{size} + \text{type} + \text{temp}$

Alpha
 0.2

Effect size
 1.2

Inputs

```
eval_design(RunMatrix = coffeedesign,
            model = ~size + type + temp,
            alpha = 0.2
            effectsize = 1.2)
```

Output

```
parameters      type      power
(Intercept)    effect.power 0.6546176
size            effect.power 0.4523370
type            effect.power 0.6546176
temp            effect.power 0.6546176
(Intercept)    parameter.power 0.6546176
size1           parameter.power 0.4962631
size2           parameter.power 0.4570906
type1           parameter.power 0.6546176
temp            parameter.power 0.6546176
```

Import and evaluate external designs

Simply import
csv/excel file and skpr
can evaluate it.

skpr automatically
detects and converts
JMP-generated split-
plot designs.

```
> jmp_design <- data.frame(read_csv("U:/R/jmp_design.csv"))
Parsed with column specification:
cols(
  x1 = col_integer(),
  x2 = col_integer(),
  x3 = col_integer(),
  x4 = col_integer(),
  x5 = col_integer(),
  x6 = col_integer(),
  Y = col_character()
)
> jmp_design$Y = NULL
> jmp_design
  x1 x2 x3 x4 x5 x6
1 -1 -1  1  1  1  1
2  1  1 -1 -1 -1 -1
3 -1  1 -1  1  1 -1
4  1  1 -1  1  1  1
5 -1 -1 -1  1 -1  1
6  1 -1  1  1 -1 -1
7 -1 -1  1 -1 -1 -1
8  1 -1  1 -1 -1  1
9 -1  1 -1 -1  1  1
10  1 -1 -1 -1  1 -1
11 -1  1  1  1 -1  1
12  1  1  1 -1  1 -1
> eval_design(jmp_design, ~., 0.2)
  parameters      type      power
1 (Intercept) effect.power 0.9689267
2          x1 effect.power 0.9445757
3          x2 effect.power 0.9445757
4          x3 effect.power 0.9445757
5          x4 effect.power 0.9445757
6          x5 effect.power 0.9445757
7          x6 effect.power 0.9445757
```

Main functions in skpr

- **gen_design**: Optimal design generation
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No need to approximate when you can simulate

Calculating power for anything more complex than a non-blocked design with a normal response requires making approximations.

Those approximations tend to fail at low numbers of runs (where power matters most).

In many cases, no approximations exist.

eval_design_mc: Simple Monte Carlo power

Parametric

```
> expand.grid(size=as.factor(c("Large", "Medium", "Small")),
+           type=as.factor(c("Kona", "Java")),
+           temp=c(80,100)) %>%
+   gen_design(~size+type+temp, 18) %>%
+   eval_design(~size+type+temp, 0.2)
```

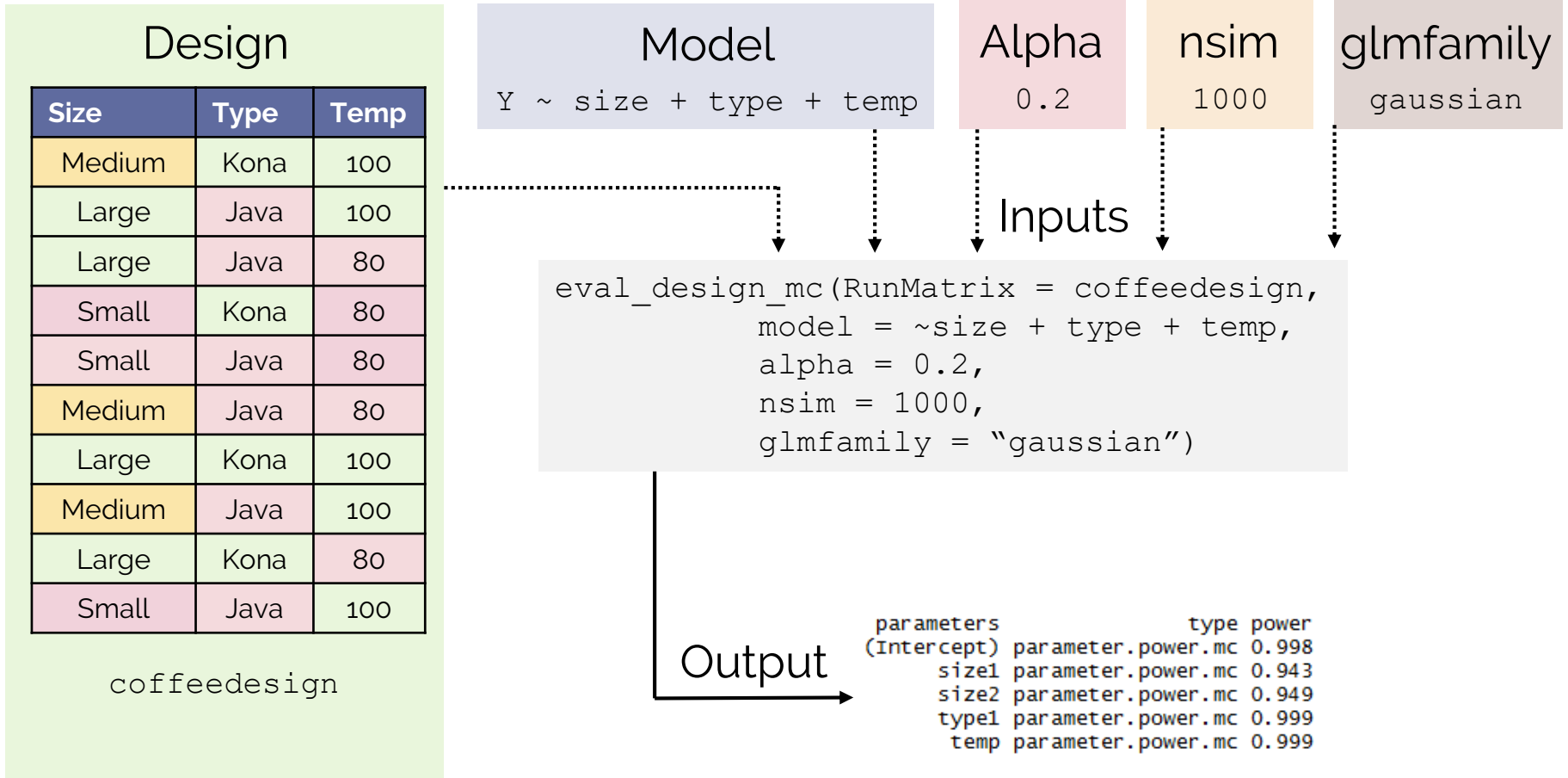
	parameters	type	power
1	(Intercept)	effect.power	0.9975907
2	size	effect.power	0.9521264
3	type	effect.power	0.9973936
4	temp	effect.power	0.9973936
5	(Intercept)	parameter.power	0.9975907
6	size1	parameter.power	0.9473817
7	size2	parameter.power	0.9473817
8	type1	parameter.power	0.9973936
9	temp	parameter.power	0.9973936

Monte Carlo

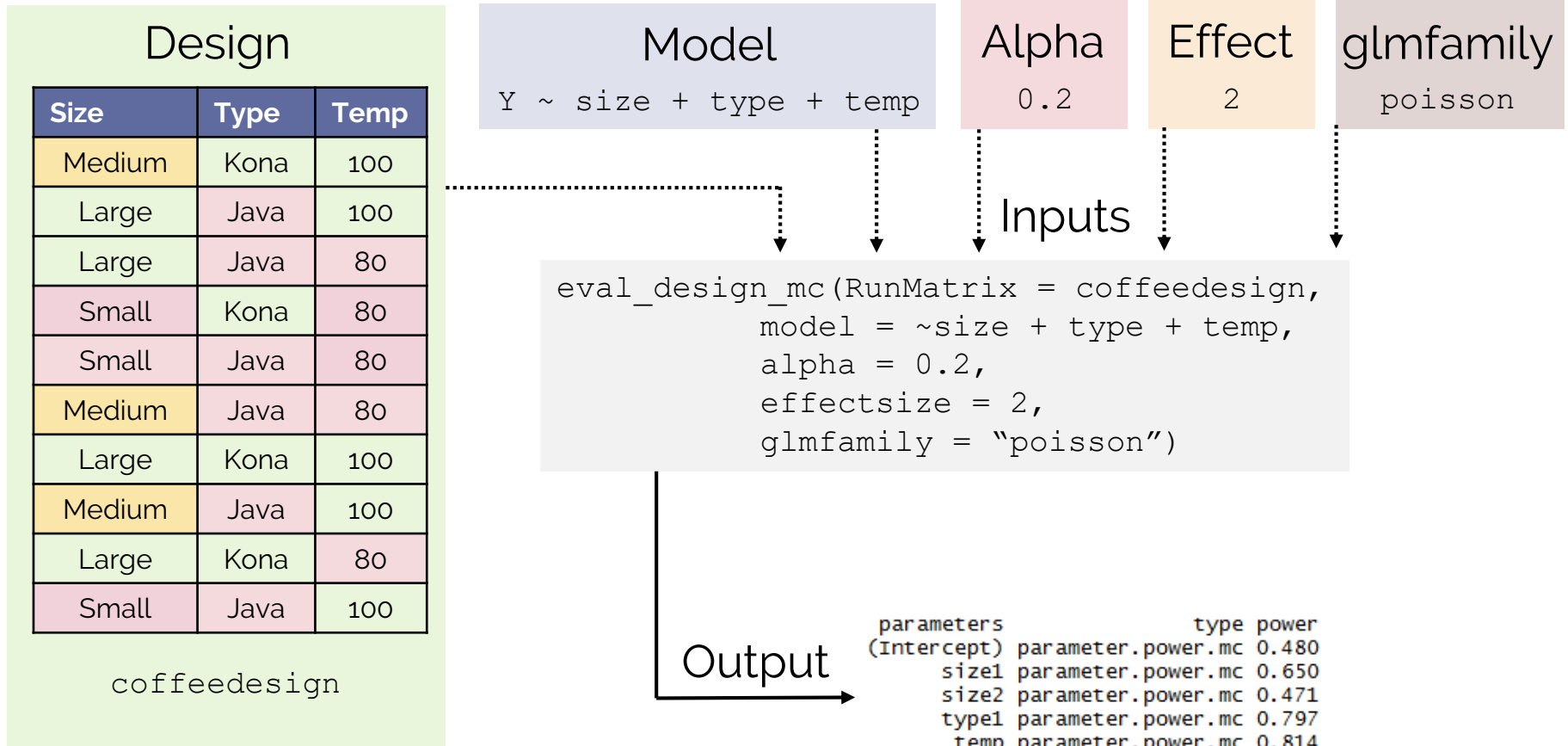
```
> expand.grid(size=as.factor(c("Large", "Medium", "Small")),
+           type=as.factor(c("Kona", "Java")),
+           temp=c(80,100)) %>%
+   gen_design(~size+type+temp, 18) %>%
+   eval_design_mc(~size+type+temp, 0.2)
```

	parameters	type	power
1	(Intercept)	parameter.power.mc	0.998
2	size1	parameter.power.mc	0.943
3	size2	parameter.power.mc	0.949
4	type1	parameter.power.mc	0.999
5	temp	parameter.power.mc	0.999

eval_design_mc – Basic Functionality

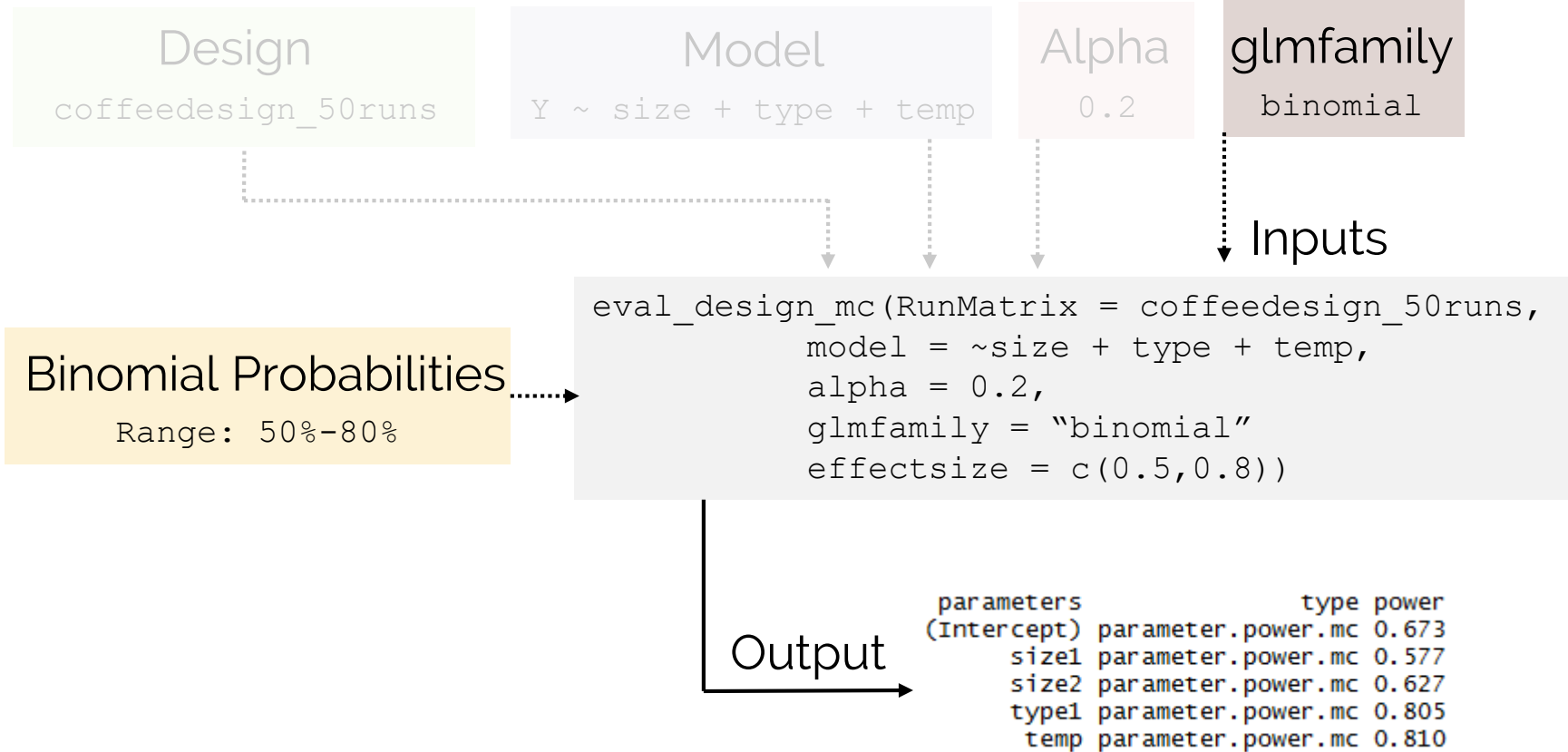


eval_design_mc – Four built-in GLM families



gaussian, binomial, poisson, exponential

eval_design_mc – Easy binomial power calculations



eval_design_mc – Split-plot power w/ REML

Design
splitplot_coffee

Model
Y ~ size + type + temp + rostdarkness

Alpha
0.2

Blocking
TRUE

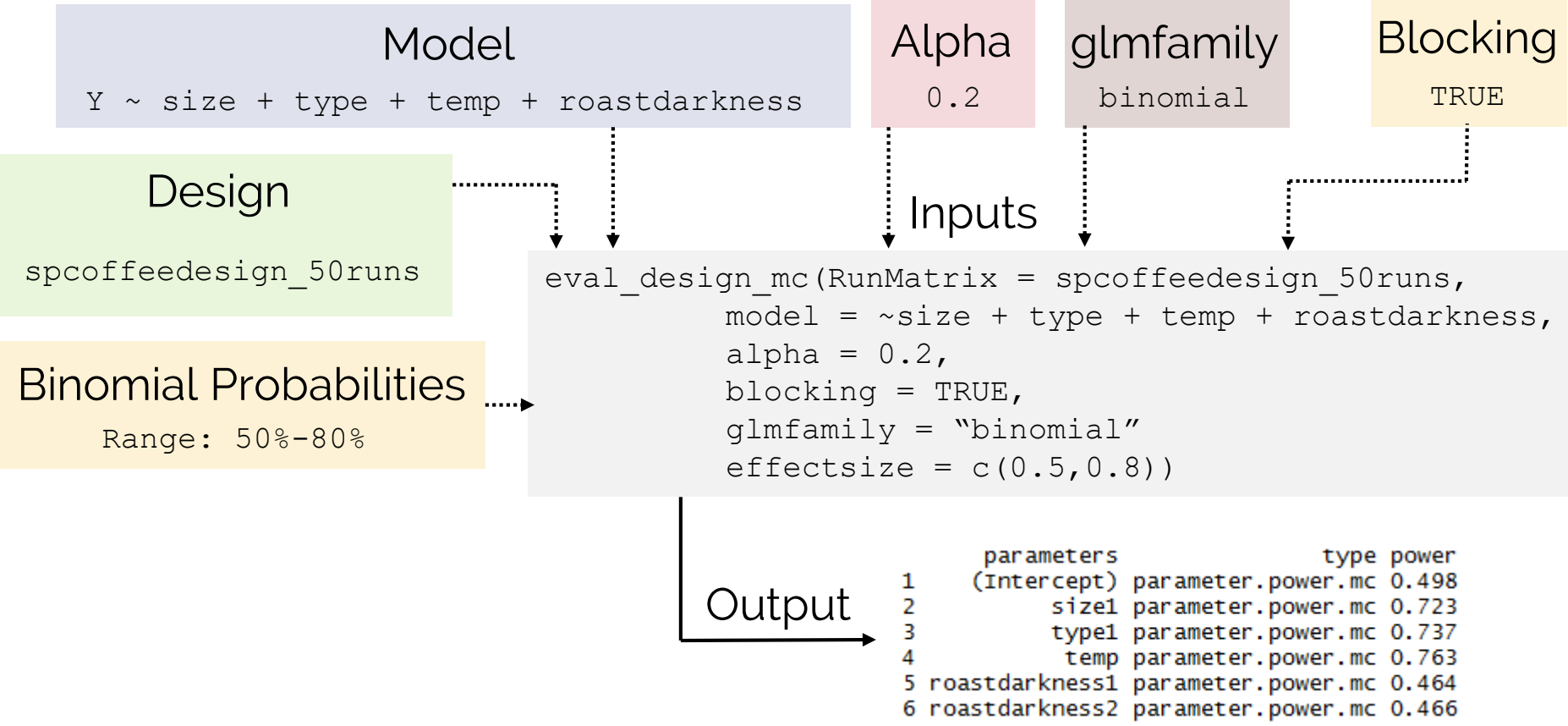
Inputs

```
eval_design_mc(RunMatrix = splitplot_coffee,  
              model = ~size + type + temp + rostdarkness,  
              alpha = 0.2,  
              blocking=TRUE)
```

Output

	parameters	type	power
1	(Intercept)	parameter.power.mc	0.781
2	size1	parameter.power.mc	0.990
3	type1	parameter.power.mc	0.999
4	temp	parameter.power.mc	0.995
5	rostdarkness1	parameter.power.mc	0.612
6	rostdarkness2	parameter.power.mc	0.616

eval_design_mc - Non-normal split-plot power



No approximate way to calculate this.

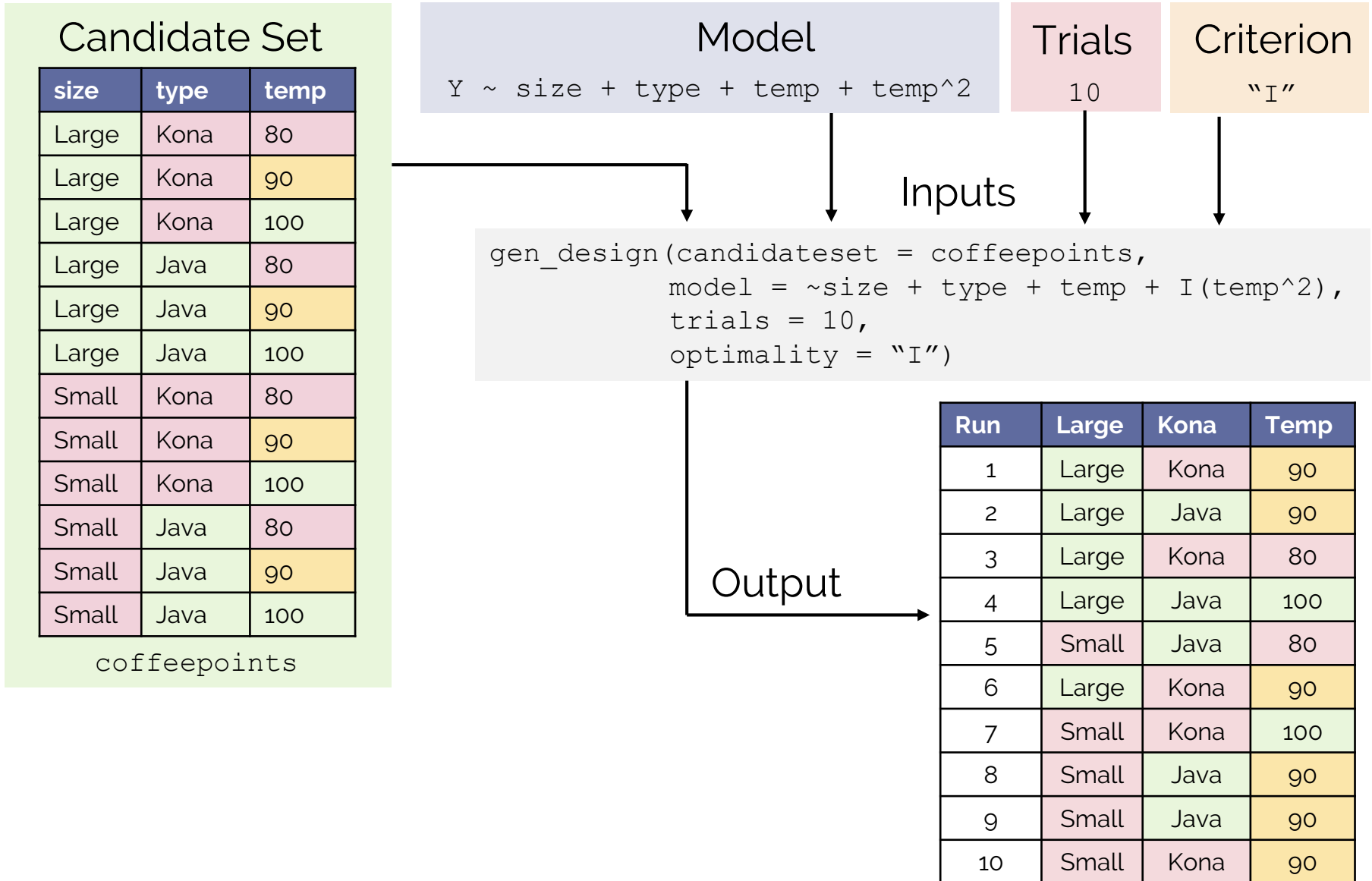
skprGUI

Summary & Questions

Quickly execute your entire design of experiments workflow in one short script. Seamlessly integrate advanced statistical techniques into your planning process. Know exactly how your results were calculated.

Extra Slides

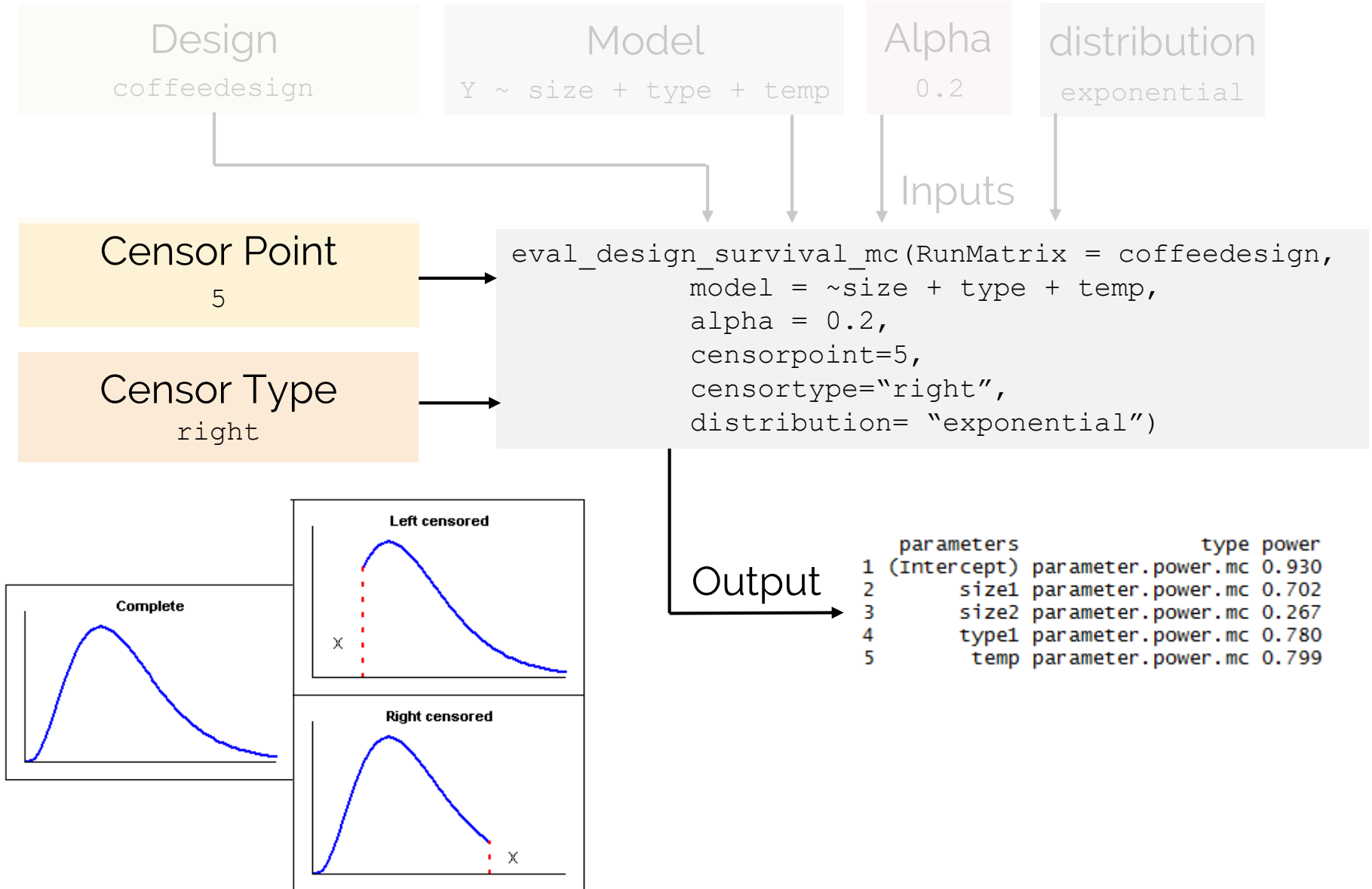
gen_design - Change optimality criterion



Five main functions

- **gen_design**: Optimal design generation
- **eval_design**: Parametric power evaluation
- **eval_design_mc**: Monte Carlo power evaluation
- **eval_design_survival_mc**: Censored Monte Carlo power evaluation
- **eval_design_custom_mc**: Custom library Monte Carlo power evaluation

eval_design_survival_mc: Censored power calculations



eval_design_custom_mc: Extensible framework

Recreate eval_design_survival_mc

```
rsurvival = function(x,b) {
  Y = rexp(n=nrow(x),rate=exp(-(X %**% b)))
  censored = Y > 1
  Y[censored] = 1
  return(Surv(time=Y,event=!censored,type="right"))
}

fitsurv = function(formula, x, contrastlist=NULL) {
  return(survreg(formula, data=x,dist="exponential"))
}

pvalsurv = function(fit) {
  return(summary(fit)$table[,4])
}

eval_design_custom_mc(RunMatrix=design,model=~a,alpha=0.05,nsim=100,
  fitfunction=fitsurv, pvalfunction=pvalsurv, rfunction=rsurvival, delta=1)
```

Same output

```
> eval_design_custom_mc(RunMatrix=design,model=~a,alpha=0.05,nsim=10000,
+   fitfunction=fitsurv, pvalfunction=pvalsurv, rfunction=rsurvival, delta=1)
  parameters          type power
1 (Intercept) parameter.power.mc 0.9388
2           a parameter.power.mc 0.9222
> eval_design_survival_mc(RunMatrix=design,model=~a,alpha=0.05,nsim=10000,
+   censortype="right",distribution="exponential",censorpoint=1,delta=1)
  parameters          type power
1 (Intercept) parameter.power.mc 0.9351
2           a parameter.power.mc 0.9146
```

Type-I error increase in non-normal split-plot designs

