# stove - clustering

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# 1 Introduction

- 1) 본 문서는 stove 패키지를 Shiny app에서 사용하는 것을 상정해 작성했습니다.
- 2) 본 문서의 케이스 스타일은 Camel case와 Snake case가 혼용되어 있습니다.
- Camel case: stove의 함수명 및 파라미터명
- Snake case: 유저로부터 받는 입력, shiny app의 server에서 사용(될 것이라고 예상)하는 object명, snake case로 작성된 dependencies의 함수명 등

# 2 Import sample data

- 1) 전처리가 완료된 샘플데이터를 불러옵니다.
- NA가 없어야 함
- string value가 있는 열은 factor로 변환

- 한 열이 모두 같은 값으로 채워져 있을 경우 제외해야 함
- Date type column이 없어야 함
- Outcome 변수는 classification의 경우 factor, regression의 경우 numeric이어야 함 (clustering은 outcome변수를 사용하지 않음)

```
# remotes::install_github("statgarten/datatoys")
library(stove)
library(datatoys)
library(dplyr)

set.seed(1234)

cleaned_data <- datatoys::bloodTest

cleaned_data <- cleaned_data %>%
    sample_n(1000) %>%
    subset(select = -c(TG))
```

# 3 K-means clustering

#### km model\$result

K-means clustering with 2 clusters of sizes 426, 574

#### Cluster means:

SEX AGE\_G HGB TCHOL HDL ANE IHD STK
1 1.497653 14.98592 14.18991 231.4671 56.5493 0.05868545 0.06338028 0.05399061
2 1.480836 14.17596 13.91446 170.3484 52.4007 0.09059233 0.06620209 0.09407666

#### Clustering vector:

 $\begin{smallmatrix} 1112 \end{smallmatrix} \end{smallmatrix} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 2 \hspace*{0.1cm} 1 \hspace*{0.1cm} 2 \hspace*$ [186] 2 2 2 2 1 2 1 1 1 1 1 1 1 2 1 2 2 2 1 2 1 1 1 1 2 2 2 2 2 2 2 1 2 1 1 2 2 1 1 [445] 2 1 1 2 2 2 2 2 2 1 1 2 2 2 2 2 2 1 1 2 2 1 1 2 2 2 1 1 2 2 2 2 1 1 2 2 2 2 1 2 2  $[630] \ 2\ 2\ 1\ 2\ 1\ 2\ 1\ 2\ 2\ 2\ 1\ 1\ 1\ 2\ 1\ 1\ 1\ 2\ 1\ 1\ 2\ 2\ 2\ 2\ 2\ 1\ 1\ 2\ 1\ 2\ 1\ 2\ 2\ 2\ 2$  $[667] \ 1\ 2\ 1\ 2\ 1\ 1\ 1\ 2\ 2\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 2\ 2\ 2\ 1\ 2\ 2\ 1\ 1\ 1\ 2\ 2\ 2\ 2$ [778] 2 2 1 1 2 1 1 1 1 2 2 1 2 2 2 1 2 1 1 1 1 2 1 1 2 2 1 2 1 2 1 2 1 2 1 1 1 2 [1000] 1

Within cluster sum of squares by cluster:

[1] 391527.4 394370.1

```
(between_SS / total_SS = 53.9 %)
```

Available components:

```
[1] "cluster" "centers" "totss" "withinss" "tot.withinss"
```

[6] "betweenss" "size" "iter" "ifault"

# 4 K-means clustering without hyperparameters

```
# K-means clustering
km_model <- stove::kMeansClustering(data = cleaned_data)
km_model$result</pre>
```

K-means clustering with 2 clusters of sizes 426, 574

Cluster means:

 SEX
 AGE\_G
 HGB
 TCHOL
 HDL
 ANE
 IHD
 STK

 1
 1.497653
 14.98592
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 231.4671
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 0.05868545
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 0.05399061

 2
 1.480836
 14.17596
 13.91446
 170.3484
 52.4007
 0.09059233
 0.06620209
 0.09407666

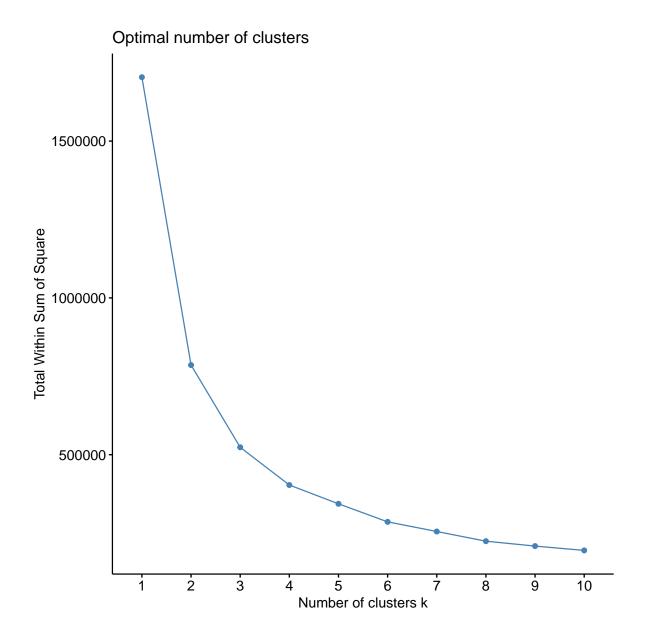
Clustering vector:

 $\begin{smallmatrix} 1112 \end{smallmatrix} \end{smallmatrix} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 1 \hspace*{0.1cm} 2 \hspace*{0.1cm} 1 \hspace*{0.1cm} 2 \hspace*$ [186] 2 2 2 2 1 2 1 1 1 1 1 1 1 2 1 2 2 2 1 2 1 1 1 1 2 2 2 2 2 2 2 1 2 1 1 2 2 1 1  $[334] \ 2\ 1\ 1\ 1\ 1\ 2\ 1\ 1\ 2\ 1\ 1\ 2\ 2\ 2\ 1\ 2\ 2\ 2\ 1\ 1\ 2\ 1\ 2\ 2\ 2\ 2\ 1\ 1\ 2\ 1$  $[445] \ 2\ 1\ 1\ 2\ 2\ 2\ 2\ 2\ 1\ 1\ 2\ 2\ 2\ 1\ 1\ 2\ 2\ 2\ 1\ 1\ 2\ 2\ 2\ 2\ 1\ 1\ 2\ 2\ 2\ 2\ 1\ 2\ 2$ [482] 2 2 2 2 2 2 2 1 2 2 2 2 1 1 2 2 2 2 1 1 2 2 2 2 1 1 1 2 2 2 1 1 1 1 2 2 2 1 2 1 2 2 

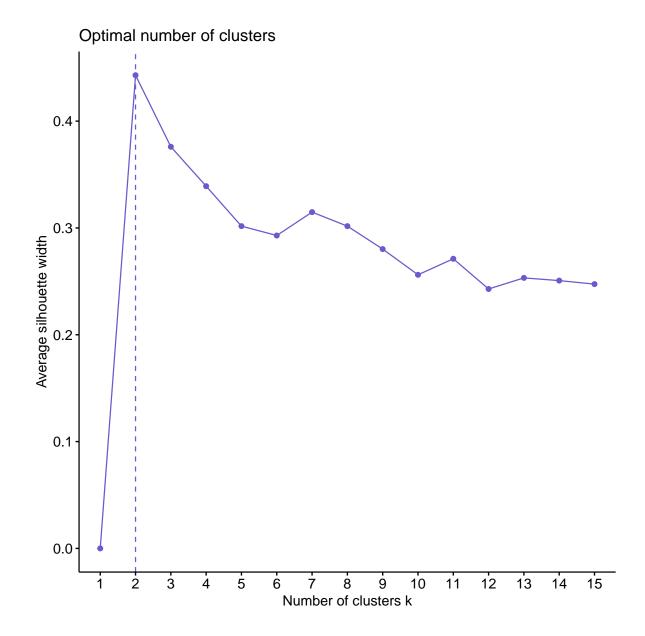
```
 [630] \ 2\ 2\ 1\ 2\ 1\ 2\ 1\ 2\ 2\ 2\ 1\ 1\ 1\ 2\ 1\ 1\ 1\ 2\ 1\ 1\ 2\ 2\ 2\ 2\ 1\ 1\ 2\ 1\ 2\ 1\ 2\ 2\ 2\ 2
[667] \ 1\ 2\ 1\ 2\ 1\ 1\ 1\ 2\ 2\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 2\ 2\ 2\ 1\ 2\ 1\ 1\ 1\ 2\ 2\ 2\ 2
[778] \ 2\ 2\ 1\ 1\ 2\ 1\ 1\ 1\ 1\ 2\ 2\ 1\ 2\ 1\ 1\ 1\ 1\ 2\ 1\ 1\ 2\ 2\ 1\ 2\ 1\ 2\ 1\ 1\ 1\ 2
[1000] 1
Within cluster sum of squares by cluster:
[1] 391527.4 394370.1
(between_SS / total_SS = 53.9 %)
Available components:
[1] "cluster"
                                 "tot.withinss"
          "centers"
                 "totss"
                         "withinss"
[6] "betweenss"
         "size"
                 "iter"
                         "ifault"
```

#### 5 Visualize

km\_model\$elbowPlot



km\_model\$optimalK



km\_model\$clustVis

