Spectral Clustering

Much of this uses information from A Tutorial on Spectral Clustering by Ulrike von Luxburg and Proximity in Statistical Machine Learning by Michael Trosset.

The Ratio Cut Problem

Ratio Cut for k=2

It can be shown that in the relaxed case for k=2, minimizing:

$$W(k) = \sum_{i=1}^{k} (x_i - m_1)^2 + \sum_{i=k+1}^{n} (x_i - m_2)^2$$

where m_1 and m_2 are k-means centers, as perscribed by Luxburg results in the same clustering as by assigning clusters by minimizing

$$R(k) = \sum_{i=1}^{k} \left(x_i + \sqrt{\frac{n-k}{k}} \right)^2 + \sum_{i=k+1}^{n} \left(x_i - \sqrt{\frac{k}{n-k}} \right)^2$$

in \mathbb{R}^1 and where x_i s are ordered, i.e., $x_i \leq \cdots \leq x_n$. We also constrain the problem to $\sum_{i=1}^{n} x_i = 0$ and $\sum_{i=1}^{n} x_i^2 = n$.

The k-means centers in $\mathbb R$ are

$$m_1 = \frac{1}{k} \sum_{i=1}^k x_i$$

$$m_2 = \frac{1}{n-k} \sum_{i=k+1}^n x_i$$

Note that m_1 and m_2 are functions of k.

Numerical results

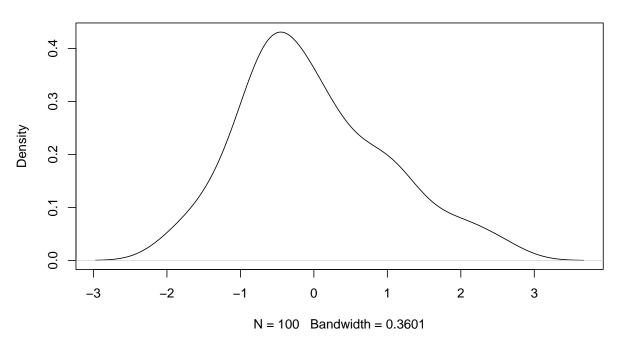
Using an arbitrary vector $\overrightarrow{x} \in \mathbb{R}^n$ such that $|\overrightarrow{x}|_1 = 0$ and $|\overrightarrow{x}|_2^2 = n$:

```
library(ggplot2)
import::from(magrittr, `%>%`)
theme_set(theme_bw())

normalize <- function(x) {
   y <- x - mean(x)
   z <- y / sqrt(mean(y ** 2))
   return(z)
}</pre>
```

```
k.means <- function(x) {
  x <- sort(x)
  n <- length(x)</pre>
  sapply(seq(n - 1), function(k) {
    m1 \leftarrow 1 / k * sum(x[seq(k)])
    m2 \leftarrow 1 / (n - k) * sum(x[seq(k + 1, n)])
    W \leftarrow sum((x[seq(k)] - m1)^2) + sum((x[seq(k + 1, n)] - m2)^2)
    return(W)
 })
}
ratio.cut <- function(x) {</pre>
  x <- sort(x)
  n <- length(x)</pre>
  sapply(seq(n - 1), function(k) {
    Ac \leftarrow sqrt((n - k) / k)
    A <- sqrt(k / (n - k))
    R \leftarrow sum((x[seq(k)] + Ac)^2) + sum((x[seq(k + 1, n)] - A)^2)
    return(R)
  })
}
# generate random data
n <- 100
k <- 4
x \leftarrow c(rnorm(n / k, -1),
        rnorm(n / k, 0),
        rnorm(n / k, 1),
        rnorm(n / k, 3))
z <- normalize(x)</pre>
plot(density(z))
```

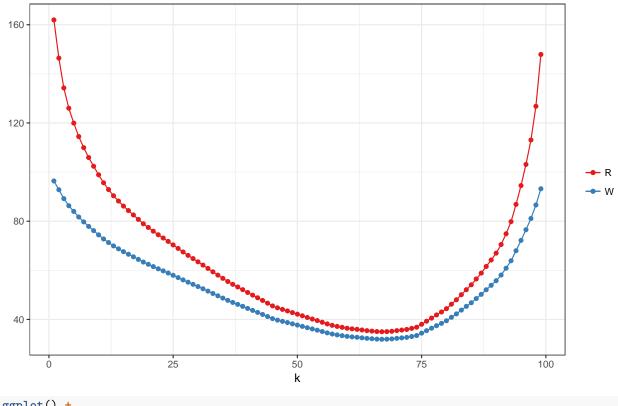
density.default(x = z)

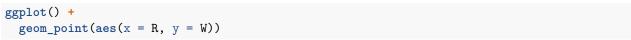


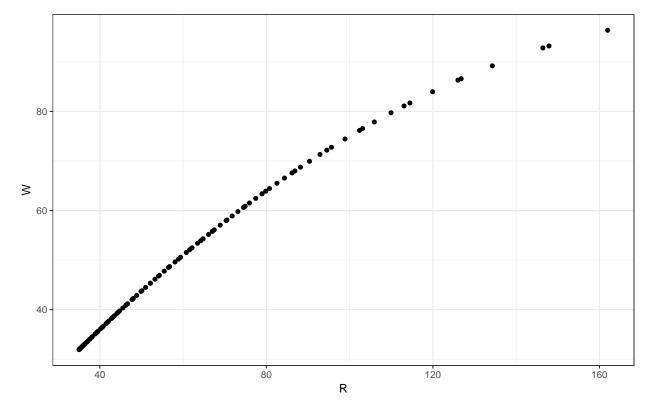
```
# compute W and R
W <- k.means(z)
R <- ratio.cut(z)

# visualizations
k <- seq(n - 1)

ggplot() +
    geom_point(aes(x = k, y = W, colour = 'W')) +
    geom_line(aes(x = k, y = W, colour = 'W')) +
    geom_point(aes(x = k, y = R, colour = 'R')) +
    geom_line(aes(x = k, y = R, colour = 'R')) +
    geom_line(aes(x = k, y = R, colour = 'R')) +
    scale_colour_brewer(palette = 'Set1') +
    labs(x = 'k', y = NULL, colour = NULL)</pre>
```

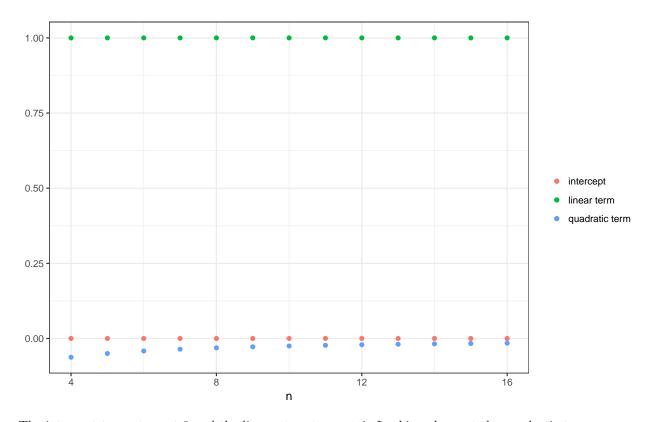






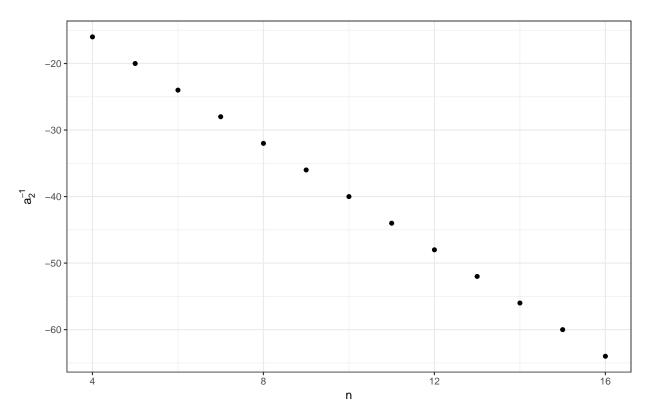
It appears that there is a definite relationship between W and R. Using quadratic regression:

```
summary(lm(W \sim R + I(R ** 2)))
Call:
lm(formula = W \sim R + I(R^2))
Residuals:
       Min
                   1Q
                           Median
                                          ЗQ
                                                     Max
-3.250e-14 -4.566e-15 2.140e-16 4.333e-15 5.514e-14
Coefficients:
              Estimate Std. Error
                                      t value Pr(>|t|)
(Intercept) 1.143e-14 6.116e-15 1.868e+00
                                               0.0648 .
             1.000e+00 1.679e-16 5.958e+15
                                               <2e-16 ***
            -2.500e-03 9.952e-19 -2.512e+15 <2e-16 ***
I(R^2)
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.035e-14 on 96 degrees of freedom
                         1, Adjusted R-squared:
Multiple R-squared:
F-statistic: 1.416e+32 on 2 and 96 DF, p-value: < 2.2e-16
\dots we get a perfect fit. We can try several values of the data size n:
\mathbb{N} \leftarrow 2 ** 4 # number of obs to try
coefs.df <- lapply(seq(4, N), function(n) {</pre>
 x <- normalize(rnorm(n)) # generate data
  # compute results
  W <- k.means(x)
  R <- ratio.cut(x)</pre>
  # compute coefs for quadratic equation
  quad.coefs <- lm(W \sim R + I(R ** 2))$coefficients
  a0 <- quad.coefs['(Intercept)']</pre>
  a1 <- quad.coefs['R']
  a2 <- quad.coefs['I(R^2)']
  # compile into data frame
  dplyr::data_frame(n, a0, a1, a2)
}) %>%
  dplyr::bind_rows()
ggplot(coefs.df) +
  geom_point(aes(x = n, y = a0, colour = 'intercept')) +
  geom_point(aes(x = n, y = a1, colour = 'linear term')) +
  geom_point(aes(x = n, y = a2, colour = 'quadratic term')) +
  labs(colour = NULL, y = NULL)
```



The intercept term stays at 0 and the linear term stays at 1. Looking closer at the quadratic term:

```
ggplot(coefs.df) +
geom_point(aes(x = n, y = a2 ** -1)) +
labs(y = expression(a[2]^-1))
```



Then we arrive at the result $W = R - \frac{1}{4n}R^2$.

Analytic result

We can show:

$$W(k) = R(k) - \frac{(R(k))^2}{4n}$$

or $W(R) = R - \frac{1}{4n}R^2$. This function is strictly increasing for $R(k) \le 2n$. Expanding R(k), we get:

$$R(k) = 2n + 2\sqrt{\frac{n-k}{k}} \sum_{i=1}^{k} x_i - 2\sqrt{\frac{k}{n-k}} \sum_{i=k+1}^{n} x_i$$

It can be shown that R(k) is maximized at the endpoints.

Note that since $\sum_{i=1}^{n} x_i = 0$, $\sum_{i=1}^{k < n} x_i \le 0$, $x_n \ge 0$, and $x_i \le 0$.

k can range from 1 to n-1. If k=n-1, we get $R(n-1)=2n+2\sqrt{\frac{1}{k}}\sum_{i}^{n-1}x_{i}-2\sqrt{n-1}x_{n}$. The second term is ≤ 0 and the third term is ≥ 0 , so we get $R \leq 2n$. On the other hand, if k=1, $R(1)=2n+2\sqrt{n-1}x_{1}-\frac{2}{\sqrt{n-1}}\sum_{i=2}^{n}x_{i}=2n+2\sqrt{n-1}x_{1}-\frac{2}{\sqrt{n-1}}(-x_{1})=2n+x_{1}\left(2\sqrt{n-1}+\frac{2}{\sqrt{n-1}}\right)\leq 2n$.

Expanding W(k), we get:

$$W(k) = \sum_{i=1}^{k} x_i^2 - 2m_1 \sum_{i=1}^{k} x_i + km_1^2 + \sum_{i=k+1}^{n} x_i^2 - 2m_2 \sum_{i=k+1}^{n} x_i + m_2^2(n-k)$$

$$= n - \frac{2}{k} \left(\sum_{i=1}^{k} x_i\right)^2 + \frac{1}{k} \left(\sum_{i=1}^{k} x_i\right)^2 - \frac{2}{n-k} \left(\sum_{i=k+1}^{n} x_i\right)^2 + \frac{1}{n-k} \left(\sum_{i=k+1}^{n} x_i\right)^2$$

$$= n - \frac{1}{k} \left(\sum_{i=1}^{k} x_i\right)^2 - \frac{1}{n-k} \left(\sum_{i=k+1}^{n} x_i\right)^2$$

$$= n - km_1^2 - (n-k)m_2^2$$

Since $\sum_{i=1}^{n} x_i = 0$, $km_1 + (n-k)m_2 = 0$, or, $-nm_2 = k(m_1 - m_2)$. Then

$$W(k) = n - km_1^2 - (n - k)m_2^2$$

$$= n - km_1^2 - nm_2^2 + km_2^2$$

$$= n - km_1^2 + (nm_2)m_2 + km_2^2$$

$$= n - km_1 + k(m_1 - m_2)m_2 + km_2^2$$

$$= n + k(-m_1^2 + m_1m_2 - m_2^2 + m_2^2)$$

$$= n + km_1(m_2 - m_1)$$

Using the relationship $-nm_2 = k(m_1 - m_2) \implies m_2 - m_1 = \frac{nm_2}{k}$ from before, we can again rewrite W(k):

$$W(k) = n - (n - k)m_2 \frac{nm_2}{k}$$
$$= n \frac{(n - k)n}{k} m_2^2$$

And we use the same relationship again: $m_2 - m_1 = \frac{n}{k} m_2 \implies m_2 = (m_2 - m_1) \frac{k}{n}$.

Then we can finally write W(k) as:

$$W(k) = n - \frac{(n-k)n}{k} \frac{k^2}{n^2} (m_1 - m_2)^2$$
$$W(k) = n - \frac{(n-k)k}{n} (m_1 - m_2)^2$$

On the other hand, if we expand R(k):

$$R(k) = \sum_{1}^{k} x_{i}^{2} + 2\sqrt{\frac{n-k}{k}} \sum_{1}^{k} x_{i} + n - k + \sum_{k+1}^{n} x_{i}^{2} - 2\sqrt{\frac{k}{n-k}} \sum_{k+1}^{n} x_{i} + k$$
$$= 2n + 2\sqrt{k(n-k)}m_{1} - 2\sqrt{k(n-k)}m_{2}$$

If we expand and simplify $-\frac{(R(k))^2}{4n}$, we get:

$$-\frac{(R(k))^2}{4n} = -n - \frac{k(n-k)}{n}m_1^2 - \frac{k(n-k)}{n}m_2^2 - 2m_1\sqrt{k(n-k)} + 2m_2\sqrt{k(n-k)} + \frac{2k(n-k)}{n}m_1m_2$$

Then noting that some terms cancel each other out, $R(k) - \frac{(R(k))^2}{4n}$:

$$n - \frac{k(n-k)}{n}m_1^2 - \frac{k(n-k)}{n}m_2^2 + 2\frac{k(n-k)}{n}m_1m_2$$
$$= n - \frac{k(n-k)}{n}(m_1^2 + m_2^2 - 2m_1m_2)$$
$$= n - \frac{k(n-k)}{n}(m_1 - m_2)^2$$

Which is exactly the same as our expression for W(k).

k > 2

When k > 2, the methods as prescribed by Luxburg involve embedding the graph to \mathbb{R}^k and then performing k-means clustering. In this case, we cannot perform the same sort of analysis since there is no way to "order" the x_i 's.