

simulation__week6

Tingyu Zhu

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```
library(tidyverse)
library(ggplot2)
source("kde_est_func.R")
```

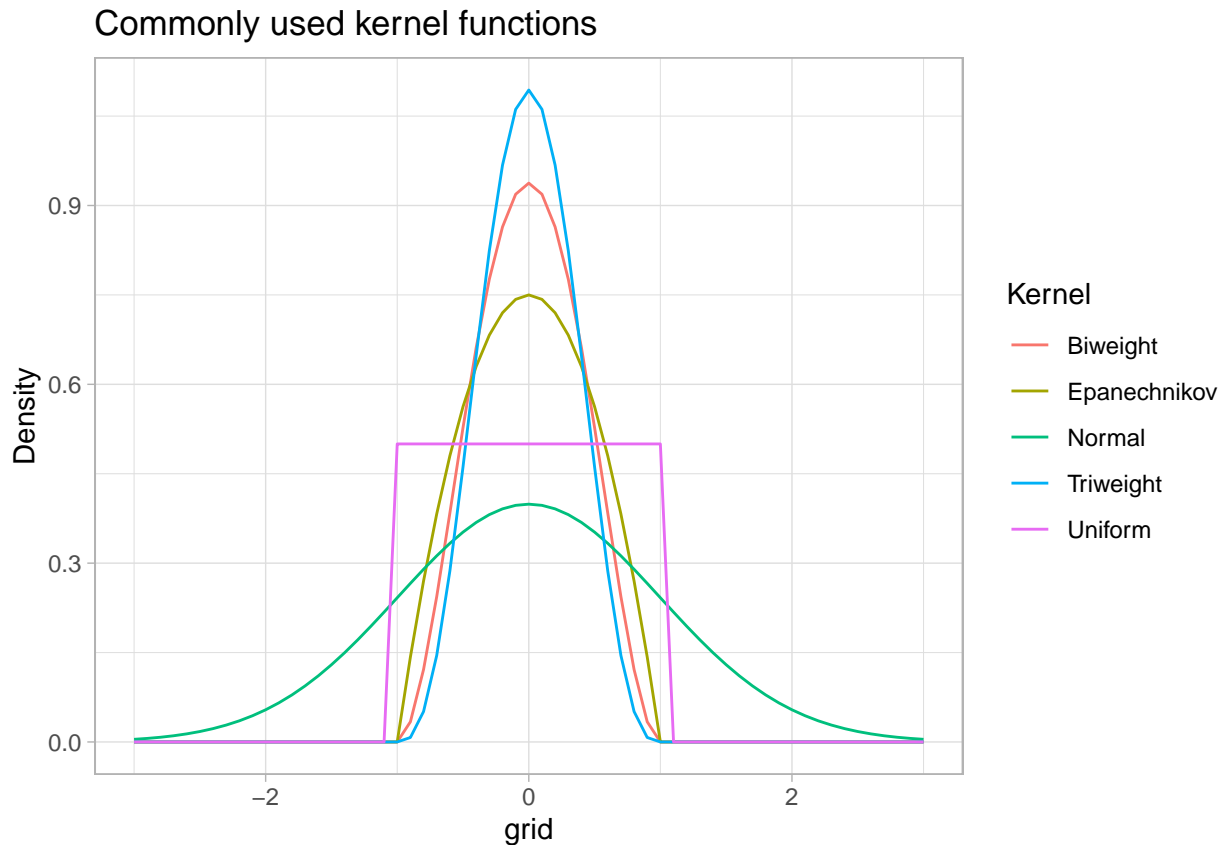
Commonly used kernel functions

In the beta family kernel density, $\lambda=0,1,2,3$ correspond to the uniform, the Epanechnikov, the biweight, and the triweight kernel functions, respectively.

```
# Define kernel functions
kernel_norm <- function(x) dnorm(x)
Kernel_beta <- function(x,lambda){(1-x^2)^lambda*(abs(x)<=1)/beta(lambda+1,0.5)}
kernel_comp <- tibble(
  grid = seq(-3,3,by=0.1),
  Normal = dnorm(grid),
  Uniform = Kernel_beta(grid,0),
  Epanechnikov = Kernel_beta(grid,1),
  Biweight = Kernel_beta(grid,2),
  Triweight = Kernel_beta(grid,3)
)

kernel_comp_long <- kernel_comp %>%
  pivot_longer(c(`Normal`, `Uniform`, `Epanechnikov`,
                `Biweight`, `Triweight`), names_to = "Kernel", values_to = "Density")

ggplot(data=kernel_comp_long) +
  geom_line(aes(x = grid, y = Density, color = Kernel)) +
  labs(title = "Commonly used kernel functions") +
  theme_light()
```



Compute the MADE (Mean Absolute Deviation Errors)

The Mean Absolute Deviation Errors for $\hat{f}(\cdot)$ is defined as

$$MADE = \frac{1}{n} \sum_{k=1}^n |\hat{f}(u_k) - f(u_k)|$$

, where $\hat{f}(u_k)$ is the kernel estimate of $f(u_k)$, and $\{u_k\}$ are the grid points taken to be arbitrary within the range of data.

When the sample is generated from the known distribution like exp, beta, gamma, etc., we can use corresponding built-in functions (`dexp()`, `dgamma`, etc.) to calculate the true densities of the grid points. If the sample is not from the familiar distributions, we can use the built-in function “`density()`” in R to get the value of $f(u_k)$ s.

```
made <- function(f_est, f_true) mean(abs(f_est - f_true))
```

sample from `nrom(mu, s^2)`

```
n <- 200
s <- 0.5
x <- rnorm(n, 1, s)
grid <- seq(-5, 5, by=0.1)
### normal reference bandwidth selector
```

```

h <- 1.06*s*n^{-0.2}
f1 <- KDE_est(x,grid,h,kernel_norm)
f1_true <- dnorm(grid, 1,s)

made(f1 , f1_true)

```

```
## [1] 0.01618103
```

```

###
### plot
data <- cbind(f1_true, f1)
matplot(grid, data, lty=c(1:2),col=c(1:2), type="l")
legend(2,0.6,c("True", "ker_est"),lty=c(1:2),col=c(1:2))

```

