

simulation__week6

Tingyu Zhu

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```
library(tidyverse)
library(purrr)
library(ggplot2)
library(here)
devtools::load_all()
set.seed(1222)
```

Commonly used kernel functions

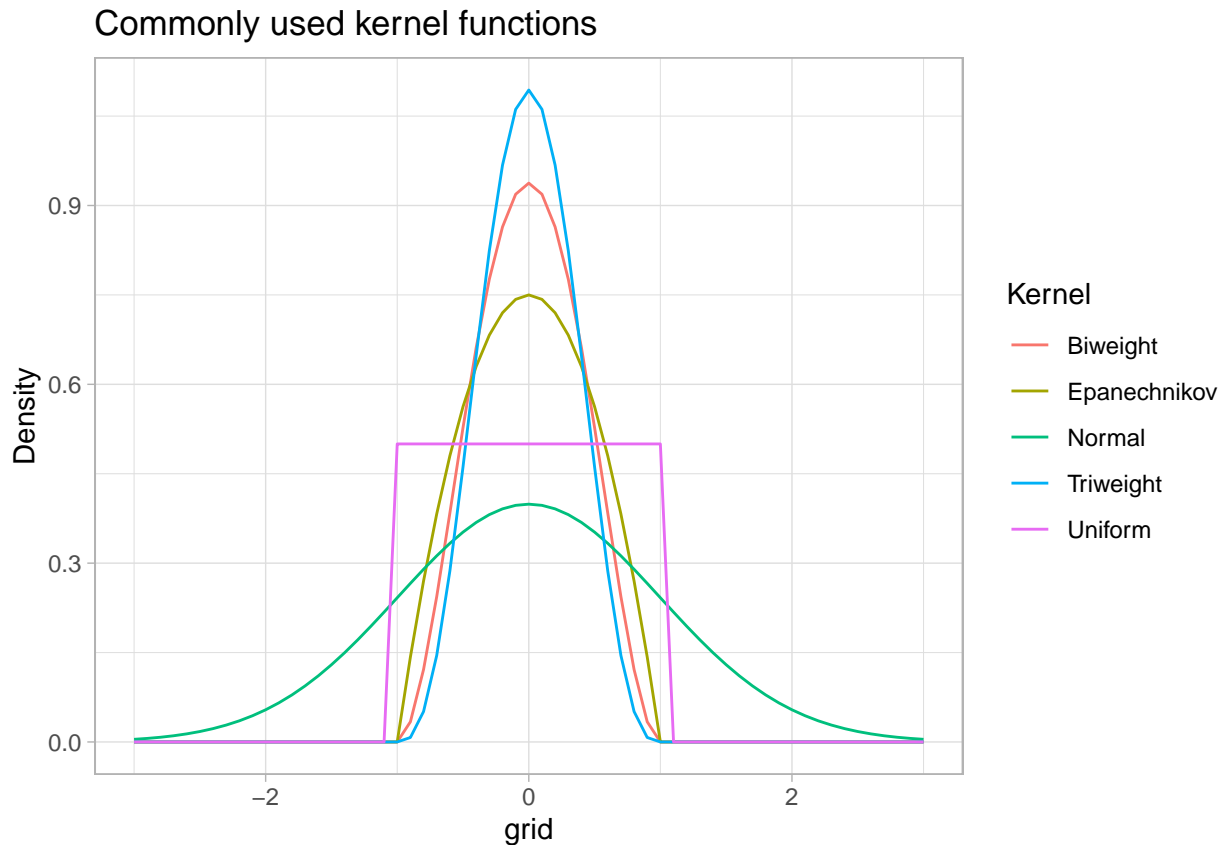
In the beta family kernel density, $\lambda=0,1,2,3$ correspond to the uniform, the Epanechnikov, the biweight, and the triweight kernel functions, respectively.

```
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))
```

Different sample sizes & kernels

```
# Define some functions for given h=bandwidth,
# simulate over different sample size and kernels

kde_n_ker_est <- function(x, n, ker, grid){
  # x large population
```

```

# n sample sizes
# ker name of kernels
# grid grid points
map2(.x = n, .y = ker,
      ~KDE_est(sample(x, .x, replace = FALSE),
                grid, h= 1.06*100(-0.2), ker = .y))
}

sim_made <- function(ns, kes, x, grid, true_f){
  # generates every combination of parameters (ns:kernels)
  simulation_params_big <- list(
    n = ns,
    kernel_type = kes)
  n_ker_est_big <- cross_df(simulation_params_big)

  n_ker_est_big <- n_ker_est_big %>%
  mutate(
    f_est = kde_n_ker_est(x, n, kernel_type, grid),
    f_true = map(1:length(n), ~true_f)
  ) %>%
  mutate(
    MADE = map2_dbl(.x = f_est, .y = f_true, ~made(.x,.y))
  )
  return(n_ker_est_big$MADE)
}

```

Simulation

Generate samples from $t(3)$ distribution.

```

x <- rt(50000, df=15)
ns <- c(100, 500, 1000, 1500)
kers <- c("normal", "uniform", "epanechnikov", "biweight", "triweight")
grid <- seq(-5,15, 0.1)
true_f <- dt(grid, df=15)

n.sim <- 100

sim_rlt <- map(1:n.sim, ~sim_made(ns, kers, x, grid,true_f))

made_mat <- matrix(unlist(sim_rlt), nrow = length(ns)*length(kers) )

made_comapre <- tibble(
  n =rep(ns, each=length(kers)),
  kernel_type = rep(kers, time=length(ns)),
  made_mean = apply(made_mat, 1, mean),
  made_sd = apply(made_mat, 1, sd)
)
write_rds(made_mat , here("results", "week6-sim.rds"))
made_comapre

## # A tibble: 20 x 4

```

##		n	kernel_type	made_mean	made_sd
##	<dbl>	<chr>		<dbl>	<dbl>
##	1	100	normal	0.00731	0.00241
##	2	100	uniform	0.00472	0.00134
##	3	100	epanechnikov	0.00414	0.00108
##	4	100	biweight	0.00386	0.000753
##	5	100	triweight	0.00913	0.00215
##	6	500	normal	0.00441	0.00109
##	7	500	uniform	0.00316	0.000774
##	8	500	epanechnikov	0.00265	0.000692
##	9	500	biweight	0.0104	0.00230
##	10	500	triweight	0.00480	0.00105
##	11	1000	normal	0.00340	0.000832
##	12	1000	uniform	0.00284	0.000667
##	13	1000	epanechnikov	0.0118	0.00313
##	14	1000	biweight	0.00513	0.00101
##	15	1000	triweight	0.00363	0.000771
##	16	1500	normal	0.00294	0.000664
##	17	1500	uniform	0.0118	0.00265
##	18	1500	epanechnikov	0.00546	0.00116
##	19	1500	biweight	0.00376	0.000669
##	20	1500	triweight	0.00313	0.000620