$Summary_week6$

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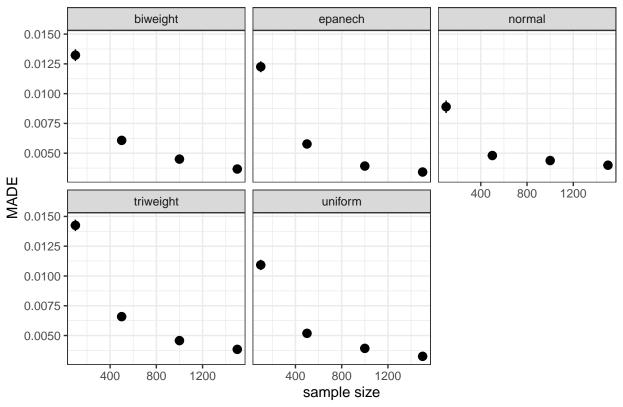
```
library(tidyverse)
library(here)
library(ggplot2)
library(purrr)
devtools::load_all()
```

Different sample sizes & kernels

```
made_comapre <- read_rds(here("results", "week6-sim-plot.rds"))
#png(here("plots", "KDE_small_size.png"),width=1600, height=900)

made_comapre %>%
    ggplot(aes(x = n, y = made_mean)) +
        geom_pointrange(aes(
            ymin = made_mean - 1.96*made_sd/sqrt(100),
            ymax = made_mean + 1.96*made_sd/sqrt(100))) +
        geom_point() +
        facet_wrap(~ kernel_type)+
        theme_bw() +ggtitle("MADEs of different kernels & sample sizes")+
        xlab("sample size")+
        ylab("MADE")+
        theme( legend.title = element_text(size = 14),
            legend.text = element_text(size = 14),
            plot.title = element_text(size=14))
```

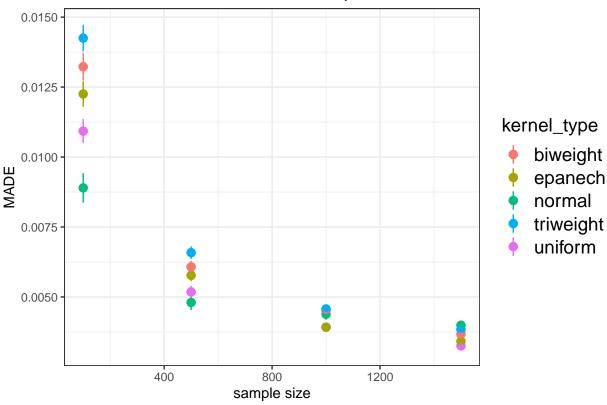
MADEs of different kernels & sample sizes



#dev.off()

Results shows that larger sample size will lead to smaller MADEs (more accurate estimation results).





#dev.off()

And with the same sample size and the same bandwidth, the MADEs are quite similar to each other. Thus, the chioce of kernel functions is not very crucial.