STA3007_hw10_codes

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2025-04-20

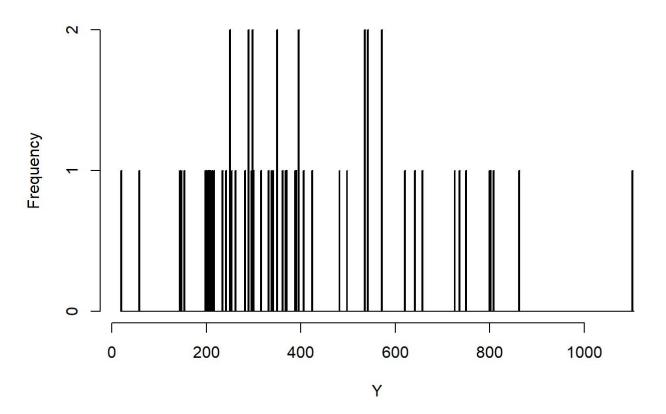
```
library(np)
## Nonparametric Kernel Methods for Mixed Datatypes (version 0.60-18)
## [vignette("np_faq", package="np") provides answers to frequently asked questions]
## [vignette("np", package="np") an overview]
## [vignette("entropy np", package="np") an overview of entropy-based methods]
library (ggplot2)
ceodat <- read.table("C:\\Users\\Penguin\\Desktop\\STA3007\\ceodat.txt")</pre>
ceodat <- ceodat[-31,]
ceodat <- ceodat[-1,]
colnames(ceodat) <- c("AGE", "SAL")</pre>
rownames(ceodat) <- 1:nrow(ceodat)</pre>
Y <- ceodat$SAL
X <- ceodat$AGE
Y <- as.numeric(Y)
X \leftarrow as.numeric(X)
```

#Histogram

```
LSCV_hist <- function(Y, h) {
  n \leftarrow 1ength(Y)
  if (n < 2) stop("Need at least two data points")
  range Y <- range(Y)
  # Create breaks covering the data range with bin width h
  breaks \langle - \text{ seq}(\text{from = floor}(\text{range } Y[1]/h)*h - h,
                 to = ceiling(range Y[2]/h)*h + h,
                by = h
  hist counts <- hist(Y, breaks = breaks, plot = FALSE)$counts
  term1 \langle - sum(hist counts^2) / (n^2 * h)
  sum term2 <- sum(hist counts * (hist counts - 1))</pre>
  term2 < -2 * sum term2 / (n * h * (n - 1))
  return(term1 - term2)
h values \langle - \text{ seq}(0.1, 2, \text{ by } = 0.1) + \text{Adjust based on data spread} \rangle
1scores <- sapply (h values, function (h) LSCV hist (Y, h))
optimal h <- h values[which.min(lscores)]</pre>
hist(Y, breaks = seq(min(Y) - optimal_h, max(Y) + optimal_h, by = optimal_h),
     main = paste ("Histogram of Y with Optimal Bin Width", round (optimal h, 2)),
     xlab = "Y", col = "lightblue", border = "black")
```

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Histogram of Y with Optimal Bin Width 2



Kernel Method

```
data <- data.frame(X = X, Y = Y)
n <- nrow(ceodat)

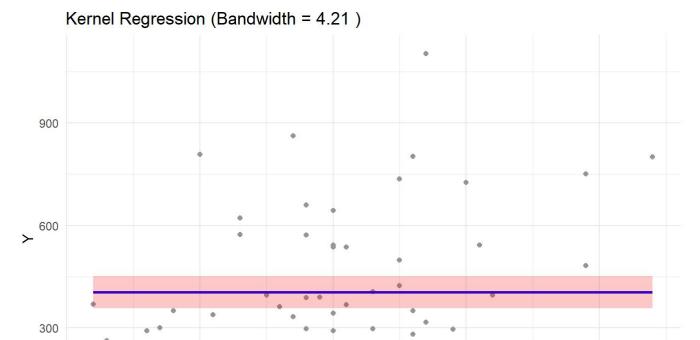
h_normal <- 1.06 * sd(X) * n^(-1/5)

# Fit kernel regression model with the computed bandwidth
bw <- npregbw(
    formula = Y ^ X,
    data = data,
    bws = h_normal,  # Use the precomputed bandwidth
    bwtype = "fixed",  # Fix the bandwidth (no cross-validation)
    ckertype = "gaussian" # Gaussian kernel
)</pre>
```

```
##
Multistart 1 of 1 |
Multistart 1 of 1 |
Multistart 1 of 1 |
Multistart 1 of 1 /
Multistart 1 of 1 |
Multistart 1 of 1 |
```

```
# Fit the model
mode1 <- npreg(bw)</pre>
# Generate predictions and confidence intervals
x grid <- seq(min(X), max(X), length.out = 100) # Grid of X values
pred <- predict(model, newdata = data.frame(X = x grid), se.fit = TRUE) # Predictions with SEs
# Compute 95% confidence bands
conf lower <- pred$fit - 1.96 * pred$se.fit</pre>
conf upper <- pred$fit + 1.96 * pred$se.fit</pre>
# Plot results with confidence bands
ggplot() +
  geom point (data = data, aes (x = X, y = Y), color = "gray60") +
  geom line(aes(x = x grid, y = pred$fit), color = "blue", linewidth = 1) +
  geom ribbon(
   aes(x = x grid, ymin = conf lower, ymax = conf upper),
   fill = "red", alpha = 0.2
  ) +
  labs(
   title = paste("Kernel Regression (Bandwidth =", round(h normal, 2), ")"),
   x = "X", y = "Y"
  theme minimal()
```

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X

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0

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```
data <- data.frame(X = X, Y = Y)
n <- nrow(ceodat)

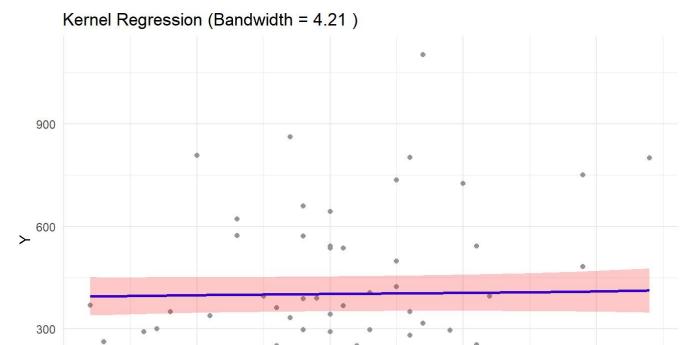
h_normal <- 1.06 * sd(X) * n^(-1/5)

# Fit kernel regression model with the computed bandwidth
bw <- npregbw(
    formula = Y ^ X,
    data = data,
    bws = h_normal,  # Use the precomputed bandwidth
    bwtype = "fixed",  # Fix the bandwidth (no cross-validation)
    ckertype = "epanechnikov"
)</pre>
```

```
##
Multistart 1 of 1 |
Multistart 1 of 1 |
Multistart 1 of 1 |
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Multistart 1 of 1 |
Multistart 1 of 1 |
```

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