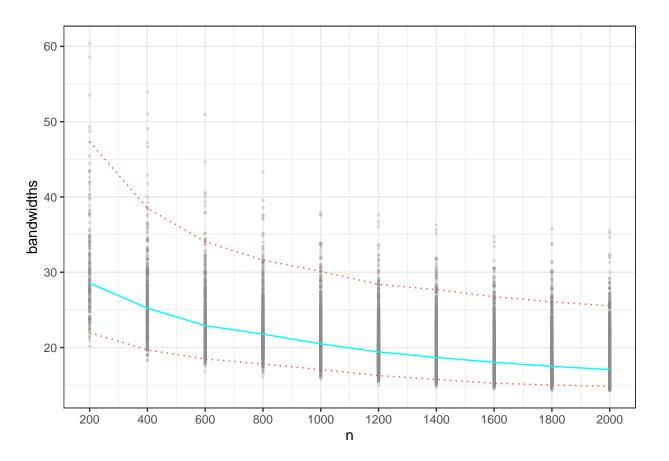
KDE-debugging-demo

```
# takes a n x 1 vector of data calculates silverman's rule of
# thumb for sigma
# https://qithub.com/statsmodels/statsmodels/blob/main/statsmodels/nonparametric/bandwidths.py
silverman <- function(x) {</pre>
    # apply and return silverman's rule of thumb
    sigma \leftarrow (0.9 * min(sd(x), IQR(x)/1.35) * length(x)^(-0.2))
    return(sigma)
}
# takes a point z, a vector of n means, and scalar sigma returns
# the density of z under the density of a mixture of Gaussians,
# each centered at mu with standard deviation sigma
phi0_k.z <- function(z, mu, sigma) {</pre>
    # calculate and return the density of z
    return((1/length(mu)) * sum(dnorm(z, mu, sigma)))
}
# vectorized version of phio_k - takes a vector Z of length N,
\# vector of n means and a scalar sigma Returns a vector of N
# densities
phi0_k <- function(Z, mu, sigma) {</pre>
    return(sapply(Z, phi0_k.z, mu = mu, sigma = sigma))
# function that takes a data matrix X and returns a bandwidth
# for each of the subjects (vector of length n)
get_bandwidths <- function(X) {</pre>
    \# get dimensions of X
    n \leftarrow nrow(X)
    p \leftarrow ncol(X)
    # find the component-wise density for each of the
    # individuals also find silverman's rule of thumb for each
    \# of the columns of X
    densities <- matrix(NA, n, p)</pre>
    sigma <- rep(NA, p)
    for (j in 1:p) {
        # find the value of sigma corresponding to the j-th
        # predictor
        sigma[j] <- silverman(X[, j])</pre>
        # calculate the resulting densities
```

```
densities[, j] <- phi0_k(X[, j], X[, j], sigma[j])</pre>
    }
    # calculate the harmonic mean for the sigma^2
    H <- 1/mean(1/sigma)</pre>
    # calculate the square root of the row-wise product of the
    # densities
    rowProds_sqrt <- rep(NA, n)</pre>
    for (1 in 1:n) {
        rowProds_sqrt[1] <- prod(sqrt(densities[1, ]))</pre>
    }
    # return the final bandwidths
    return(H/rowProds_sqrt)
}
# generate some data and try it out
set.seed(1)
n <- 2000
p <- 5
limits \leftarrow matrix(1:p, p, 2) * matrix(c(-1, 1), p, 2, T)
X <- matrix(runif(n * p, limits[, 1], limits[, 2]), n, p, T)</pre>
summary(X)
                              ٧2
                                                 VЗ
                                                                      ۷4
##
          ۷1
          :-0.997946
                               :-1.99727
                                                 :-2.999276
                                                                       :-3.99840
## Min.
                        Min.
                                           Min.
                                                               Min.
## 1st Qu.:-0.486186
                        1st Qu.:-1.00985
                                           1st Qu.:-1.488389
                                                               1st Qu.:-2.02589
## Median :-0.016691
                        Median :-0.01738
                                           Median :-0.006958
                                                               Median :-0.11502
## Mean
         :-0.005323
                        Mean :-0.01006
                                           Mean : 0.001604
                                                               Mean :-0.04986
## 3rd Qu.: 0.507396
                        3rd Qu.: 1.02458
                                           3rd Qu.: 1.553405
                                                                3rd Qu.: 1.94929
## Max. : 0.997820
                        Max. : 1.99305
                                           Max. : 2.995168
                                                               Max. : 3.99945
##
          ۷5
## Min.
         :-4.99894
## 1st Qu.:-2.37765
## Median : 0.09663
## Mean : 0.11980
## 3rd Qu.: 2.70045
## Max. : 4.99455
# demo the functions:
library(ggplot2)
# silverman rule of thumb for each column of X:
apply(X, 2, silverman)
## [1] 0.1138329 0.2312477 0.3442222 0.4513766 0.5785286
# different n to try
(n_j \leftarrow seq(200, n, 200))
```

[1] 200 400 600 800 1000 1200 1400 1600 1800 2000

```
# matrices for storing quantiles and bandwidths
bandwidth.dist <- matrix(NA, 3, length(n_j))</pre>
colnames(bandwidth.dist) <- paste("n =", n_j)</pre>
rownames(bandwidth.dist) <- c("0.025", "0.5", "0.975")
bandwidths <- vector("list", length(n_j))</pre>
# plot the distribution of the bandwidths for varying p
bandwidth_dist_graph <- ggplot()</pre>
for (j in 1:length(n_j)) {
    bandwidths[[j]] \leftarrow get\_bandwidths(X[1:n_j[j], ])
    bandwidth.dist[, j] <- quantile(bandwidths[[j]], c(0.025, 0.5,
    bandwidth_dist_graph <- bandwidth_dist_graph + geom_point(data = cbind.data.frame(n = n_j[j],
        bandwidths = bandwidths[[j]]), aes(n, bandwidths), size = 0.5,
        color = "gray55", alpha = 0.3)
}
bandwidth_dist_graph + theme_bw() + geom_line(data = data.frame(n = n_j,
    bandwidths = bandwidth.dist[1, ]), aes(n, bandwidths), color = "tomato2",
    linetype = "dotted") + geom_line(data = data.frame(n = n_j, bandwidths = bandwidth.dist[3,
    ]), aes(n, bandwidths), color = "tomato2", linetype = "dotted") +
    geom_line(data = data.frame(n = n_j, bandwidths = bandwidth.dist[2,
        ]), aes(n, bandwidths), color = "cyan") + scale_x_continuous(labels = n_j,
    breaks = n_j)
```



bandwidth.dist