

# Package ‘rbbnp’

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**Type** Package

**Title** A Bias Bound Approach to Non-Parametric Inference

**Version** 0.3.0

**Maintainer** Xinyu DAI <xinyu\_dai@brown.edu>

**Description** A novel bias-bound approach for non-parametric inference is introduced, focusing on both density and conditional expectation estimation.

It constructs valid confidence intervals that account for the presence of a non-negligible bias and thus make it possible to perform inference with optimal mean squared error minimizing bandwidths.

This package is based on Schennach (2020) <[doi:10.1093/restud/rdz065](https://doi.org/10.1093/restud/rdz065)>.

**License** GPL (>= 3)

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**Imports** purrr, pracma, tidyverse, dplyr, ggplot2, gridExtra

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**Author** Xinyu DAI [aut, cre],  
Susanne M Schennach [aut]

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**Index****27****biasBound\_condExpectation***Bias bound approach for conditional expectation estimation***Description**

Estimates the density at a given point or across a range, and provides visualization options for density, bias, and confidence intervals.

**Usage**

```
biasBound_condExpectation(
  Y,
  X,
  x = NULL,
  h = NULL,
  h_method = "cv",
  alpha = 0.05,
  est_Ar = NULL,
  resol = 100,
  xi_lb = NULL,
  xi_ub = NULL,
  methods_get_xi = "Schennach",
  if_plot_ft = FALSE,
  ora_Ar = NULL,
  if_plot_conditional_mean = TRUE,
  kernel.fun = "Schennach2004",
  if_approx_kernel = TRUE,
  kernel.resol = 1000
)
```

**Arguments**

Y	A numerical vector of sample data.
X	A numerical vector of sample data.
x	Optional. A scalar or range of points where the density is estimated. If NULL, a range is automatically generated.
h	A scalar bandwidth parameter. If NULL, the bandwidth is automatically selected using the method specified in 'h_method'.
h_method	Method for automatic bandwidth selection when h is NULL. Options are "cv" (cross-validation) and "silverman" (Silverman's rule of thumb). Default is "cv".
alpha	Confidence level for intervals. Default is 0.05.
est_Ar	Optional list of estimates for A and r. If NULL, they are computed using get_est_Ar().
resol	Resolution for the estimation range. Default is 100.
xi_lb	Optional. Lower bound for the interval of Fourier Transform frequency xi. Used for determining the range over which A and r is estimated. If NULL, it is automatically determined based on the methods_get_xi.
xi_ub	Optional. Upper bound for the interval of Fourier Transform frequency xi. Similar to xi_lb, it defines the upper range for A and r estimation. If NULL, the upper bound is determined based on the methods_get_xi.
methods_get_xi	A string specifying the method to automatically determine the xi interval if xi_lb and xi_ub are NULL. Options are "Schennach" and "Schennach_loose". If "Schennach" the range is selected based on the Theorem 2 in Schennach2020, if "Schennach_loose", it is defined by the initial interval given in Theorem 2 without selecting the xi_n.

if\_plot\_ft      Logical. If TRUE, plots the Fourier transform.  
 ora\_Ar            Optional list of oracle values for A and r.  
 if\_plot\_conditional\_mean  
                   Logical. If TRUE, plots the conditional mean estimation.  
 kernel.fun        A string specifying the kernel function to be used. Options are "Schennach2004",  
                   "sinc", "normal", "epanechnikov".  
 if\_approx\_kernel  
                   Logical. If TRUE, uses approximations for the kernel function.  
 kernel.resol      The resolution for kernel function approximation. See [fun\\_approx](#).

### Value

A list containing various outputs including estimated values, plots, and intervals.

### Examples

```

# Example 1: point estimation of conditional expectation of Y on X
biasBound_condExpectation(
  Y = sample_data$Y,
  X = sample_data$X,
  x = 1,
  h = 0.09,
  kernel.fun = "Schennach2004"
)

# Example 2: conditional expectation with automatic bandwidth selection using cross-validation
# biasBound_condExpectation(
#   Y = sample_data$Y,
#   X = sample_data$X,
#   h = NULL,
#   h_method = "cv",
#   xi_lb = 1,
#   xi_ub = 12,
#   kernel.fun = "Schennach2004"
# )

# Example 3: conditional expectation with automatic bandwidth selection using Silverman's rule
# biasBound_condExpectation(
#   Y = sample_data$Y,
#   X = sample_data$X,
#   h = NULL,
#   h_method = "silverman",
#   methods_get_xi = "Schennach",
#   if_plot_ft = TRUE,
#   kernel.fun = "Schennach2004"
# )
  
```

---

<code>biasBound_density</code>	<i>Bias bound approach for density estimation</i>
--------------------------------	---

---

## Description

Estimates the density at a given point or across a range, and provides visualization options for density, bias, and confidence intervals.

## Usage

```
biasBound_density(
  X,
  x = NULL,
  h = NULL,
  h_method = "cv",
  alpha = 0.05,
  resol = 100,
  xi_lb = NULL,
  xi_ub = NULL,
  methods_get_xi = "Schennach",
  if_plot_density = TRUE,
  if_plot_ft = FALSE,
  ora_Ar = NULL,
  kernel.fun = "Schennach2004",
  if_approx_kernel = TRUE,
  kernel.resol = 1000
)
```

## Arguments

X	A numerical vector of sample data.
x	Optional. A scalar or range of points where the density is estimated. If NULL, a range is automatically generated.
h	A scalar bandwidth parameter. If NULL, the bandwidth is automatically selected using the method specified in 'h_method'.
h_method	Method for automatic bandwidth selection when h is NULL. Options are "cv" (cross-validation) and "silverman" (Silverman's rule of thumb). Default is "cv".
alpha	Confidence level for intervals. Default is 0.05.
resol	Resolution for the estimation range. Default is 100.
xi_lb	Optional. Lower bound for the interval of Fourier Transform frequency xi. Used for determining the range over which A and r is estimated. If NULL, it is automatically determined based on the methods_get_xi.
xi_ub	Optional. Upper bound for the interval of Fourier Transform frequency xi. Similar to xi_lb, it defines the upper range for A and r estimation. If NULL, the upper bound is determined based on the methods_get_xi.

**methods\_get\_xi** A string specifying the method to automatically determine the xi interval if xi\_lb and xi\_ub are NULL. Options are "Schennach" and "Schennach\_loose". If "Schennach" the range is selected based on the Theorem 2 in Schennach2020, if "Schennach\_loose", it is defined by the initial interval given in Theorem 2 without selecting the xi\_n.

**if\_plot\_density** Logical. If TRUE, plots the density estimation.

**if\_plot\_ft** Logical. If TRUE, plots the Fourier transform.

**ora\_Ar** Optional list of oracle values for A and r.

**kernel.fun** A string specifying the kernel function to be used. Options are "Schennach2004", "sinc", "normal", "epanechnikov".

**if\_approx\_kernel** Logical. If TRUE, uses approximations for the kernel function.

**kernel.resol** The resolution for kernel function approximation. See [fun\\_approx](#).

## Value

A list containing various outputs including estimated values, plots, and intervals.

## Examples

```
# Example 1: Specifying x for point estimation with manually selected xi range
# from a fixed bandwidth
biasBound_density(
  X = sample_data$X,
  x = 1,
  h = 0.09,
  xi_lb = 1,
  xi_ub = 12,
  if_plot_ft = TRUE,
  kernel.fun = "Schennach2004"
)

# Example 2: Density estimation with automatic bandwidth selection using cross-validation
# biasBound_density(
#   X = sample_data$X,
#   h = NULL,
#   h_method = "cv",
#   xi_lb = 1,
#   xi_ub = 12,
#   if_plot_ft = FALSE,
#   kernel.fun = "Schennach2004"
# )

# Example 3: Density estimation with automatic bandwidth selection using Silverman's rule
# biasBound_density(
#   X = sample_data$X,
#   h = NULL,
#   h_method = "silverman",
#   methods_get_xi = "Schennach",
```

```
#   if_plot_ft = TRUE,
#   kernel.fun = "Schennach2004"
# )
```

**create\_biasBound\_config***Create a configuration object for bias bound estimations***Description**

Create a configuration object for bias bound estimations

**Usage**

```
create_biasBound_config(
  X,
  Y = NULL,
  h = NULL,
  h_method = "cv",
  use_fft = TRUE,
  alpha = 0.05,
  resol = 100,
  xi_lb = NULL,
  xi_ub = NULL,
  methods_get_xi = "Schennach",
  kernel.fun = "Schennach2004",
  if_approx_kernel = TRUE,
  kernel.resol = 1000
)
```

**Arguments**

X	A numerical vector of sample data.
Y	Optional. A numerical vector of sample data for conditional expectation.
h	A scalar bandwidth parameter. If NULL, the bandwidth is automatically selected using the method specified in 'h_method'.
h_method	Method for automatic bandwidth selection when h is NULL. Options are "cv" (cross-validation) and "silverman" (Silverman's rule of thumb). Default is "cv".
use_fft	Ignored. Maintained for backward compatibility.
alpha	Confidence level for intervals.
resol	Resolution for the estimation range.
xi_lb	Lower bound for the interval of Fourier Transform frequency.
xi_ub	Upper bound for the interval of Fourier Transform frequency.
methods_get_xi	Method to determine xi interval.

kernel.fun Kernel function to be used. Options include "normal", "epanechnikov", "Schennach2004", and "sinc".  
 if\_approx\_kernel Use approximations for the kernel function.  
 kernel.resol Resolution for kernel approximation.

### Value

A configuration object (list) with all parameters

## create\_kernel\_functions

*Create kernel functions based on configuration*

### Description

Create kernel functions based on configuration

### Usage

```
create_kernel_functions(
  kernel.fun = "Schennach2004",
  if_approx_kernel = TRUE,
  kernel.resol = 1000
)
```

### Arguments

kernel.fun A string specifying the kernel function to be used.  
 if\_approx\_kernel Logical. If TRUE, uses approximations for the kernel function.  
 kernel.resol The resolution for kernel function approximation.

### Value

A list containing kernel function, its Fourier transform, and the kernel type

---

**cv\_bandwidth***Cross-Validation for Bandwidth Selection*

---

**Description**

Implements least-squares cross-validation for bandwidth selection with any kernel function. Uses the self-convolution approach for accurate estimation of the integral term.

**Usage**

```
cv_bandwidth(  
  X,  
  h_grid = NULL,  
  kernel_func,  
  kernel_type = "normal",  
  grid_size = 512  
)
```

**Arguments**

X	A numerical vector of sample data.
h_grid	A numerical vector of bandwidth values to evaluate. If NULL (default), a grid is automatically generated based on the range and distribution of the data.
kernel_func	The kernel function to use for cross-validation.
kernel_type	A string identifying the kernel type, used only for reference bandwidth.
grid_size	Number of grid points for evaluation. Default is 512.

**Value**

A scalar representing the optimal bandwidth that minimizes the cross-validation score.

**Examples**

```
# Generate sample data  
X <- rnorm(100)  
# Get optimal bandwidth using cross-validation with a normal kernel  
kernel_functions <- create_kernel_functions("normal")  
h_opt <- cv_bandwidth(X, kernel_func = kernel_functions$kernel,  
                      kernel_type = kernel_functions$kernel_type)
```

DATA\_PATH

*The Path to the Data Folder***Description**

This variable provides the path to the data folder within the package.

**Value**

The path to the package's internal data folder as a character string.

epanechnikov\_kernel

*Epanechnikov Kernel***Description**

Epanechnikov Kernel

**Usage**

```
epanechnikov_kernel(u)
```

**Arguments**

u

A numerical value or vector representing the input to the kernel function.

**Value**

Returns the value of the Epanechnikov kernel function at the given input.

epanechnikov\_kernel\_ft

*Fourier Transform Epanechnikov Kernel***Description**

Fourier Transform Epanechnikov Kernel

**Usage**

```
epanechnikov_kernel_ft(xi)
```

**Arguments**

xi

A numerical value or vector representing the frequency domain.

**Value**

Returns the value of the Fourier transform of the Epanechnikov kernel at the given frequency/frequencies.

EXT\_DATA\_PATH

*The Path to the External Data Folder for Non-R Data Files***Description**

This variable provides the path to the extdata folder within the package, where non-standard R data files are stored.

**Value**

The path to the package's external data folder (for non-standard R data files) as a character string.

fun\_approx

*Approximation Function for Intensive Calculations***Description**

This function provides a lookup-based approximation for calculations that are computationally intensive. Once computed, it stores the results in an environment and uses linear interpolation for new data points to speed up subsequent computations.

**Usage**

```
fun_approx(u, u_lb = -100, u_ub = 100, resol = 10000, fun = W_kernel)
```

**Arguments**

u	A vector of values where the function should be evaluated.
u_lb	Lower bound for the precomputed range. Defaults to -10.
u_ub	Upper bound for the precomputed range. Defaults to 10.
resol	The resolution or number of sample points in the precomputed range. Defaults to 1000.
fun	A function for which the approximation is computed. Defaults to the W function.

## Details

The `fun_approx` function works by initially creating a lookup table of function values based on the range specified by `u_lb` and `u_ub` and the resolution `resol`. This precomputation only happens once for a given set of parameters (`u_lb`, `u_ub`, `resol`, and `fun`). Subsequent calls to `fun_approx` with the same parameters use the lookup table to find the closest precomputed points to the requested `u` values and then return an interpolated result.

Linear interpolation is used between the two closest precomputed points in the lookup table. This ensures a smooth approximation for values in between sample points.

This function is especially useful for computationally intensive functions where recalculating function values is expensive or time-consuming. By using a combination of precomputation and interpolation, `fun_approx` provides a balance between accuracy and speed.

## Value

A vector of approximated function values corresponding to `u`.

`gen_sample_data`      *Generate Sample Data*

## Description

This function used for generate some sample data for experiment

## Usage

```
gen_sample_data(size, dgp, seed = NULL)
```

## Arguments

<code>size</code>	control the sample size.
<code>dgp</code>	data generating process, have options "normal", "chisq", "mixed", "poly", "2_fold_uniform".
<code>seed</code>	random seed number.

## Value

A numeric vector of length `size`. The elements of the vector are generated according to the specified `dgp`:

**normal** Normally distributed values with mean 0 and standard deviation 2.

**chisq** Chi-squared distributed values with `df` = 10.

**mixed** Half normally distributed (mean 0, `sd` = 2) and half chi-squared distributed (`df` = 10) values.

**poly** Values from a polynomial cumulative distribution function on [0, 1].

**2\_fold\_uniform** Sum of two uniformly distributed random numbers.

---

get\_avg\_f1x                    *Kernel point estimation*

---

**Description**

Computes the point estimate using the specified kernel function.

**Usage**

```
get_avg_f1x(X, x, h, inf_k)
```

**Arguments**

X	A numerical vector of sample data.
x	A scalar representing the point where the density is estimated.
h	A scalar bandwidth parameter.
inf_k	Kernel function used for the computation.

**Value**

A scalar representing the kernel density estimate at point x.

---

get\_avg\_fyx                    *Kernel point estimation*

---

**Description**

Computes the point estimate using the specified kernel function.

**Usage**

```
get_avg_fyx(Y, X, x, h, inf_k)
```

**Arguments**

Y	A numerical vector representing the sample data of variable Y.
X	A numerical vector representing the sample data of variable X.
x	A scalar representing the point where the density is estimated.
h	A scalar bandwidth parameter.
inf_k	Kernel function used for the computation.

**Value**

A scalar representing the kernel density estimate at point x.

`get_avg_phi`*Compute Sample Average of Fourier Transform Magnitude***Description**

Compute Sample Average of Fourier Transform Magnitude

**Usage**

```
get_avg_phi(Y = 1, X, xi)
```

**Arguments**

- |                 |   |
|-----------------|---|
| <code>Y</code>  | A numerical vector representing the sample data of variable Y.                                  |
| <code>X</code>  | A numerical vector representing the sample data of variable X.                                  |
| <code>xi</code> | A single numerical value representing the frequency at which the Fourier transform is computed. |

**Value**

Returns the sample estimation of expected Fourier transform at frequency `xi`.

`get_avg_phi_log`*Compute log sample average of fourier transform and get mod***Description**

Compute log sample average of fourier transform and get mod

**Usage**

```
get_avg_phi_log(Y = 1, X, ln_xi)
```

**Arguments**

- |                    |   |
|--------------------|---|
| <code>Y</code>     | A numerical vector representing the sample data of variable Y.                                      |
| <code>X</code>     | A numerical vector representing the sample data of variable X.                                      |
| <code>ln_xi</code> | A single numerical value representing the log frequency at which the Fourier transform is computed. |

**Value**

Returns the log sample estimation of expected Fourier transform at frequency `xi`.

`get_conditional_var`    *get the conditional variance of Y on X for given x*

### Description

get the conditional variance of Y on X for given x

### Usage

```
get_conditional_var(X, Y, x, h, kernel_func)
```

### Arguments

X	A numerical vector representing the sample data of variable X.
Y	A numerical vector representing the sample data of variable Y.
x	The specific point at which the conditional variance is to be calculated.
h	A bandwidth parameter used in the kernel function for smoothing.
kernel_func	A kernel function used to weigh observations in the neighborhood of point x.

### Value

Returns a non-negative scalar representing the estimated conditional variance of Y given X at the point x. Returns 0 if the computed variance is negative.

`get_est_Ar`    *get the estimation of A and r*

### Description

This function estimates the parameters A and r by optimizing an objective function over a specified range of frequency values and r values.

### Usage

```
get_est_Ar(Y = 1, X, xi_interval, r_stepsize = 150)
```

### Arguments

Y	A numerical vector representing the sample data of variable Y.
X	A numerical vector representing the sample data of variable X.
xi_interval	A list with elements xi_lb and xi_ub representing the lower and upper bounds of the frequency interval.
r_stepsize	An integer value representing the number of steps in the r range. This controls the granularity of the estimation. Higher values lead to finer granularity but increase computation time.

**Details**

The function internally defines a range for the natural logarithm of frequency values (`ln_xi_range`) and a range for the parameter  $r$  (`r_range`). It then defines an optimization function `optim_ln_A` to minimize the integral of a given function over the `ln_xi_range`. The actual estimation is done by finding the  $r$  and  $A$  value that minimizes the area of the line  $\ln A - r \ln \xi$  under the constraint that the line should not go below the Fourier transform curve.

**Value**

A named vector with elements `est_A` and `est_r` representing the estimated values of  $A$  and  $r$ , respectively.

`get_est_B`*get the estimation of B***Description**

get the estimation of  $B$

**Usage**

```
get_est_B(Y)
```

**Arguments**

<code>Y</code>	A numerical vector representing the sample data of variable $Y$ .
----------------	---

**Value**

The mean of the absolute values of the elements in  $Y$ , representing the estimated value of  $B$ .

`get_est_b1x`*Estimation of bias b1x***Description**

Computes the bias estimate for given parameters.

**Usage**

```
get_est_b1x(X, h, est_Ar, inf_k_ft, ...)
```

**Arguments**

X	A numerical vector representing the sample data of variable X.
h	A scalar bandwidth parameter.
est_Ar	A vector containing the estimated A and r parameters.
inf_k_ft	A kernel Fourier transform function.
...	Additional arguments passed to the quadgk integration function.

**Value**

A scalar representing the bias b1x estimate.

get\_est\_byx

*Estimation of bias byx***Description**

Estimation of bias byx

**Usage**

```
get_est_byx(Y, X, ...)
```

**Arguments**

Y	A numerical vector representing the sample data of variable Y.
X	A numerical vector representing the sample data of variable X.
...	Additional arguments passed to other methods.

**Value**

A scalar representing the bias byx estimate.

get\_est\_vy

*get the estimation of Vy***Description**

get the estimation of Vy

**Usage**

```
get_est_vy(Y)
```

**Arguments**

Y	A numerical vector representing the sample data of variable Y.
---	--

**get\_sigma***Estimation of sigma***Description**

Computes the sigma estimate for given parameters.

**Usage**

```
get_sigma(X, x, h, inf_k)
```

**Arguments**

X	A numerical vector of sample data.
x	A scalar representing the point where the density is estimated.
h	A scalar bandwidth parameter.
inf_k	Kernel function used for the computation.

**Value**

A scalar representing the sigma estimate at point x. Returns 0 if the density estimate is negative.

**get\_sigma\_yx***Estimation of sigma\_yx***Description**

Estimation of sigma\_yx

**Usage**

```
get_sigma_yx(Y, X, x, h, inf_k)
```

**Arguments**

Y	A numerical vector representing the sample data of variable Y.
X	A numerical vector representing the sample data of variable X.
x	The specific point at which sigma_yx is to be estimated.
h	A bandwidth parameter used in the kernel function for smoothing.
inf_k	A kernel function used to weigh observations in the neighborhood of point x.

**Value**

Returns a scalar representing the estimated value of sigma\_yx at the point x. Returns 0 if either fyx or conditional variance is negative.

---

get_xi_interval	<i>get xi interval</i>
-----------------	------------------------

---

### Description

get xi interval

### Usage

```
get_xi_interval(Y = 1, X, methods = "Schennach")
```

### Arguments

Y	A numerical vector representing the sample data of variable Y.
X	A numerical vector representing the sample data of variable X.
methods	A character string indicating the method to use for calculating the xi interval. Supported methods are "Schennach" and "Schennach_loose". Defaults to "Schennach".

### Details

The "Schennach" method computes the xi interval by performing a test based on the Schennach's theorem, adjusting the upper bound  $\xi_{ub}$  if the test condition is met. The "Schennach\_loose" method provides a looser calculation of the xi interval without performing the Schennach's test.

### Value

A list containing the lower ( $\xi_{lb}$ ) and upper ( $\xi_{ub}$ ) bounds of the xi interval.

---

kernel_reg	<i>Kernel Regression function</i>
------------	-----------------------------------

---

### Description

Kernel Regression function

### Usage

```
kernel_reg(X, Y, x, h, kernel_func)
```

**Arguments**

X	A numerical vector representing the sample data of variable X.
Y	A numerical vector representing the sample data of variable Y.
x	The point at which the regression function is to be estimated.
h	A bandwidth parameter that determines the weight assigned to each observation in X.
kernel_func	A function that computes the weight of each observation based on its distance to x.

**Value**

Returns a scalar representing the estimated value of the regression function at the point x.

normal_kernel	<i>Normal Kernel Function</i>
---------------	-------------------------------

**Description**

Normal Kernel Function

**Usage**

```
normal_kernel(u)
```

**Arguments**

u	A numerical value or vector representing the input to the kernel function.
---	--

**Value**

Returns the value of the Normal kernel function at the given input.

normal_kernel_ft	<i>Fourier Transform of Normal Kernel</i>
------------------	---

**Description**

Fourier Transform of Normal Kernel

**Usage**

```
normal_kernel_ft(xi)
```

**Arguments**

- `xi` A numerical value or vector representing the frequency domain.

**Value**

Returns the value of the Fourier transform of the Normal kernel at the given frequency/frequencies.

<code>plot_ft</code>	<i>Plot the Fourier Transform</i>
----------------------	-----------------------------------

**Description**

Plot the Fourier Transform of the

**Usage**

```
plot_ft(X, xi_interval, ft_plot.resol = 500)
```

**Arguments**

- `X` A numerical vector of sample data.  
`xi_interval` A list containing the lower (`xi_lb`) and upper (`xi_ub`) bounds of the `xi` interval.  
`ft_plot.resol` An integer representing the resolution of the plot, specifically the number of points used to represent the Fourier transform. Defaults to 500.

**Details**

$C = 1$ , the parameter in  $O(1/n^{0.25})$ , see more details in Schennach (2020).

**Value**

A ggplot object representing the plot of the Fourier transform.

**Examples**

```
plot_ft(
  sample_data$X,
  xi_interval = list(xi_lb = 1, xi_ub = 50),
  ft_plot.resol = 1000
)
```

**rpoly01***Generate n samples from the distribution***Description**

Generate n samples from the distribution

**Usage**

```
rpoly01(n, k = 5)
```

**Arguments**

- n               The number of samples to generate.
- k               The exponent in the distribution function, defaults to 5.

**Value**

A vector of n samples from the specified polynomial distribution.

CDF:  $f(x) = (x-1)^k + 1$

**sample\_data***Sample Data***Description**

Sample Data

**Usage**

```
sample_data
```

**Format**

A data frame with 1000 rows and 2 variables:

**X** Numeric vector, generated from 2 fold uniform distribution.

**Y** Numeric vector,  $Y = -X^2 + 3*X + rnorm(1000)*X$ .

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<code>select_bandwidth</code>	<i>Select Optimal Bandwidth</i>
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## Description

Selects an optimal bandwidth using the specified method.

## Usage

```
select_bandwidth(
  X,
  Y = NULL,
  method = "cv",
  kernel.fun = "normal",
  if_approx_kernel = TRUE,
  kernel.resol = 1000
)
```

## Arguments

<code>X</code>	A numerical vector of sample data.
<code>Y</code>	Optional. A numerical vector of sample data for conditional expectation estimation.
<code>method</code>	A string specifying the bandwidth selection method. Options are "cv" for cross-validation and "silverman" for Silverman's rule of thumb. Defaults to "cv".
<code>kernel.fun</code>	A string specifying the kernel type. Options include "normal", "epanechnikov", "Schennach2004", and "sinc".
<code>if_approx_kernel</code>	Logical. If TRUE, uses approximations for the kernel function.
<code>kernel.resol</code>	The resolution for kernel function approximation.

## Value

A scalar representing the optimal bandwidth.

## Examples

```
# Generate sample data
X <- rnorm(100)
# Get optimal bandwidth using cross-validation with normal kernel
h_opt <- select_bandwidth(X, method = "cv", kernel.fun = "normal")
# Get optimal bandwidth using Silverman's rule with Schennach kernel
h_opt <- select_bandwidth(X, method = "silverman", kernel.fun = "Schennach2004")
```

`silverman_bandwidth`    *Silverman's Rule of Thumb for Bandwidth Selection*

### Description

Implements Silverman's rule of thumb for selecting an optimal bandwidth in kernel density estimation.

### Usage

```
silverman_bandwidth(X, kernel_type = "normal")
```

### Arguments

<code>X</code>	A numerical vector of sample data.
<code>kernel_type</code>	A string identifying the kernel type.

### Value

A scalar representing the optimal bandwidth.

### Examples

```
# Generate sample data
X <- rnorm(100)
# Get optimal bandwidth using Silverman's rule
h_opt <- silverman_bandwidth(X, kernel_type = "normal")
```

`sinc`                          *Infinite Kernel Function*

### Description

Infinite Kernel Function

### Usage

```
sinc(u)
```

### Arguments

<code>u</code>	A numerical value or vector where the sinc function is evaluated.
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### Value

The value of the sinc function at each point in `u`.

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**sinc\_ft***Define the closed form FT of the infinite order kernel sin(x)/(pi\*x)*

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**Description**

Define the closed form FT of the infinite order kernel  $\sin(x)/(pi*x)$

**Usage**

`sinc_ft(x)`

**Arguments**

`x` A numerical value or vector where the Fourier Transform is evaluated.

**Value**

The value of the Fourier Transform of the sinc function at each point in `x`.

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**true\_density\_2fold***True density of 2-fold uniform distribution*

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**Description**

True density of 2-fold uniform distribution

**Usage**

`true_density_2fold(x)`

**Arguments**

`x` A numerical value or vector where the true density function is evaluated.

**Value**

The value of the true density of the 2-fold uniform distribution at each point in `x`.

**W\_kernel1***Define the inverse Fourier transform function of W***Description**

Define the inverse Fourier transform function of W

**Usage**

```
W_kernel1(u, L = 10)
```

**Arguments**

- |   |  |
|---|--|
| u | A numerical value or vector representing the time or space domain.                                   |
| L | The limit for numerical integration, defines the range of integration as $[-L, L]$ . Defaults to 10. |

**Value**

A numerical value or vector representing the inverse Fourier transform of the infinite order kernel at the given time or space point(s).

**W\_kernel\_ft***Define the Fourier transform of a infinite kernel proposed in Schennach 2004***Description**

Define the Fourier transform of a infinite kernel proposed in Schennach 2004

**Usage**

```
W_kernel_ft(xi, xi_lb = 0.5, xi_ub = 1.5)
```

**Arguments**

- |       |  |
|-------|--|
| xi    | A numerical value or vector representing the frequency domain. |
| xi_lb | The lower bound for the frequency domain. Defaults to 0.5.     |
| xi_ub | The upper bound for the frequency domain. Defaults to 1.5.     |

**Value**

A numerical value or vector representing the Fourier transform of the infinite order kernel at the given frequency/frequencies.

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