

Package ‘kdensity’

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Type Package

Title Kernel Density Estimation with Parametric Starts and Asymmetric
Kernels

Version 1.2.0

Description Handles univariate non-parametric density estimation with parametric starts and asymmetric kernels in a simple and flexible way. Kernel density estimation with parametric starts involves fitting a parametric density to the data before making a correction with kernel density estimation, see Hjort & Glad (1995) <[doi:10.1214/aos/1176324627](https://doi.org/10.1214/aos/1176324627)>. Asymmetric kernels make kernel density estimation more efficient on bounded intervals such as (0, 1) and the positive half-line. Supported asymmetric kernels are the gamma kernel of Chen (2000) <[doi:10.1023/A:1004165218295](https://doi.org/10.1023/A:1004165218295)>, the beta kernel of Chen (1999) <[doi:10.1016/S0167-9473\(99\)00010-9](https://doi.org/10.1016/S0167-9473(99)00010-9)>, and the copula kernel of Jones & Henderson (2007) <[doi:10.1093/biomet/asm068](https://doi.org/10.1093/biomet/asm068)>. User-supplied kernels, parametric starts, and bandwidths are supported.

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URL <https://github.com/JonasMoss/kdensity>

BugReports <https://github.com/JonasMoss/kdensity/issues>

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bandwidths	<i>Bandwidth Selectors</i>
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Description

The available options for bandwidth selectors, passed as the `bw` argument to `kdensity`.

Arguments

<code>x</code>	The input data.
<code>kernel_str</code>	A string specifying the kernel, e.g. "gaussian."
<code>start_str</code>	A string specifying the parametric start, e.g. "normal".
<code>support</code>	The domain of definition for the kernel. (-Inf, Inf) for symmetric kernels.

Details

The bandwidth functions are not exported. They are members of the environment `bw_environments`, and can be accessed by `kdensity:::bw_environments`.

Bandwidth selectors

"nrd0", "nrd", "bcv", "SJ": Bandwidth selectors from `stats`. They are documented in `[bandwidth][stats:bandwidth] stats:bandwidth`. "nrd0" is the standard bandwidth selector for symmetric kernels with constant parametric starts.

"ucv": Unbiased cross validation. The standard option for asymmetric kernels.

"HS": Hallberg Szabadváry's closed-form reference rule for the beta kernel on the unit interval. This is the default for `kernel = "beta"` with a uniform or constant start.

"RHE": Selector for parametric starts with a symmetric kernel, based on a reference rule with Hermite polynomials. Described in Hjort & Glad (1995). The default method in `kdensity` when a parametric start is supplied and the kernel is symmetric.

"JH": Selector for the Gaussian copula kernel, based on normal reference rule. Described in Jones & Henderson. The default method when the `gcopula` kernel is used in `kdensity`.

Structure

The bandwidth selector is a function of four arguments: The data `x`, a kernel string `kernel`, a start string `start`, and a support vector `support`. To obtain the functions associated with these strings, use `get_kernel` and `get_start`. The function should return a double.

References

Jones, M. C., and D. A. Henderson. "Kernel-type density estimation on the unit interval." *Biometrika* 94.4 (2007): 977-984. Hjort, Nils Lid, and Ingrid K. Glad. "Nonparametric density estimation with a parametric start." *The Annals of Statistics* (1995): 882-904. Hallberg Szabadváry, Johan. "A Fast, Closed-Form Bandwidth Selector for the Beta Kernel Density Estimator." arXiv preprint arXiv:2601.19553 (2026).

See Also

`kdensity()`, `stats::bandwidth.kernel()` for the bandwidth selectors of `stats::density()`. In addition, `kernels()`; `parametric_starts()`

Examples

```
## Not a serious bandwidth function.
silly_width <- function(x, kernel = NULL, start = NULL, support = NULL) {
  rexp(1)
}
kdensity(mtcars$mpg, start = "gumbel", bw = silly_width)
```

kdensity

Parametrically guided kernel density estimation

Description

`kdensity` computes a parametrically guided kernel density estimate for univariate data. It supports asymmetric kernels and parametric starts through the `kernel` and `start` arguments.

Usage

```
kdensity(
  x,
  bw = NULL,
  adjust = 1,
  kernel = NULL,
  start = NULL,
  support = NULL,
  na.rm = FALSE,
  normalized = TRUE,
  tolerance = 0.01
)
```

Arguments

`x` Numeric vector containing the data.

`bw` A bandwidth function. Can be either a string, a custom-made function, or a double. The supported bandwidth functions are documented in `bandwidths()`.

adjust	An adjustment constant, so that $h = \text{adjust} \cdot \text{bw} \cdot \text{sd}$, where sd varies with the chosen kernel.
kernel	The kernel function. Can be chosen from the list of built-in kernels or be custom-made. See <code>kernels()</code> for details.
start	Parametric start. Can be chosen from the list of built-in parametric starts or be custom-made. See <code>parametric_starts()</code> for details.
support	The support of the data. Must be compatible with the supplied x and the supplied start and kernel. Is used to find the normalization constant, see <code>normalized</code> .
na.rm	Logical; if TRUE, NAs will be removed from x .
normalized	Logical; if TRUE, the density is normalized.
tolerance	Numeric; the relative error to tolerate in normalization.

Details

The default values for `bw`, `kernel`, `start`, and `support` are interdependent, and are chosen to make sense. E.g., the default value for `support` when `start = beta` is `c(0, 1)`.

The `start` argument defaults to `uniform`, which corresponds to ordinary kernel density estimation. The typical default value for `kernel` is `gaussian`.

If `normalized` is FALSE and `start != "uniform"`, the resulting density will not integrate to 1 in general.

Value

`kdensity` returns an S3 function object of `base::class()` "kdensity". This is a callable function with the following elements, accessible by '\$':

`x` The data supplied in x .

`bw_str`, `bw`, `adjust`, `h` The bandwidth function, the resulting bandwidth, the `adjust` argument, and the adjusted bandwidth.

`kernel_str`, `kernel`, `start`, `start_str`, `support` Name of the kernel, the kernel object, name of the parametric start, the start object, and the support of the density.

`data.name`, `n`, `range`, `has.na`, `na.rm`, `normalized` Name of the data, number of observations, the range of the data, whether the data x contained NA values, whether `na.rm` is TRUE or not, and whether the density is normalized.

`call` The call to `kdensity`.

`estimates` Named numeric vector containing the parameter estimates from the parametric start.

`logLik` The log-likelihood of the parametric starts. Is NA for the uniform start.

References

Hjort, Nils Lid, and Ingrid K. Glad. "Nonparametric density estimation with a parametric start." *The Annals of Statistics* (1995): 882-904.

Jones, M. C., and D. A. Henderson. "Miscellanea kernel-type density estimation on the unit interval." *Biometrika* 94.4 (2007): 977-984.

Chen, Song Xi. "Probability density function estimation using gamma kernels." *Annals of the Institute of Statistical Mathematics* 52.3 (2000): 471-480.

Silverman, Bernard W. *Density estimation for statistics and data analysis*. Vol. 26. CRC press, 1986.

See Also

The stats package function `stats::density()`.

Examples

```
## Use gamma kernels to model positive data, the concentration of
## theophylline

concentration <- Theoph$conc + 0.001
plot(kdensity(concentration, start = "gamma", kernel = "gamma", adjust = 1 / 3),
     ylim = c(0, 0.15), lwd = 2, main = "Concentration of theophylline"
)
lines(kdensity(concentration, start = "gamma", kernel = "gaussian"),
      lty = 2, col = "grey", lwd = 2
)
lines(kdensity(concentration, start = "gaussian", kernel = "gaussian"),
      lty = 3, col = "blue", lwd = 2
)
lines(kdensity(concentration, start = "gaussian", kernel = "gamma", adjust = 1 / 3),
      lty = 4, col = "red", lwd = 2
)
rug(concentration)
```

kernels

Kernel functions

Description

Kernel functions are an important part of `kdensity`. This document lists the available built-in functions and the structure of them. Any kernel in the list can be used in `kdensity` by using `kernel = "kernel"` for the intended kernel.

Details

Be careful combining kernels with compact support with parametric starts, as the normalizing integral typically fails to converge. Use `gaussian` instead.

Symmetric kernels

`gaussian`, `normal`: The Gaussian kernel. The default argument when starts is supported on R. `epanechnikov`, `rectangular` (uniform), `triangular`, `biweight`, `cosine`, `optcosine`: Standard symmetric kernels, also used in `stats::density()`. `tricube`, `triweight`: Standard symmetric kernels. Not supported by `stats::density()`. `laplace`: Uses the Laplace density, also known as the double exponential density.

Asymmetric kernels

`gamma`, `gamma_biased`: The gamma kernel of Chen (2000). For use on the positive half-line. `gamma` is the recommended biased-corrected kernel. `gcopula`: The Gaussian copula kernel of Jones & Henderson (2007). For use on the unit interval. `beta`, `beta_biased`: The beta kernel of Chen (1999). For use on the unit interval. `beta` is the recommended bias-corrected kernel.

Structure

A kernel is a list containing two mandatory elements and one optional element. The mandatory element 'kernel' is the kernel function. It takes arguments `y`, `x`, `h`, where `x` is the data supplied to `kdensity` and `y` is the point of evaluation. `h` is the bandwidth. Internally, the kernel function is evaluated as $1/h * \text{kernel}(y, x, h)$. It should be vectorized in `x`, but vectorization in `y` is not needed.

The second mandatory element is `support`, stating the domain of definition for the kernel. This is used to distinguish kernels on the unit interval / positive half-line from kernels on \mathbb{R} .

`sd` is used for symmetric kernels, and states the standard error of the kernel. This is used to make kernels comparable to the Gaussian kernel when calculating bandwidths.

References

Chen, Song Xi. "Probability density function estimation using gamma kernels." *Annals of the Institute of Statistical Mathematics* 52.3 (2000): 471-480. Jones, M. C., and D. A. Henderson. "Kernel-type density estimation on the unit interval." *Biometrika* 94.4 (2007): 977-984. Chen, Song Xi. "Beta kernel estimators for density functions." *Computational Statistics & Data Analysis* 31.2 (1999): 131-145.

See Also

[kdensity\(\)](#); [parametric_starts\(\)](#); [bandwidths\(\)](#).

Examples

```
gaussian <- list(
  kernel = function(y, x, h) stats::dnorm((y - x) / h),
  sd = 1,
  support = c(-Inf, Inf)
)

gcopula <- list(
  kernel = function(y, x, h) {
    rho <- 1 - h^2
    inside <- rho^2 * (qnorm(y)^2 + qnorm(x)^2) - 2 * rho * qnorm(y) * qnorm(x)
    exp(-inside / (2 * (1 - rho^2)))
  },
  support = c(0, 1)
)
```

parametric_starts *Parametric starts*

Description

A parametric start is a density function with an associated estimator which is used as a starting point in `kdensity`. Several parametric starts are implemented, all with maximum likelihood estimation. Custom-made parametric starts are possible, see the Structure section.

Structure

The parametric start contains three elements: The density function, an estimation function, and the support of the density. The parameters of the density function must partially match the parameters of the estimator function. The estimator function takes one argument, a numeric vector, which is passed from `kdensity`.

Supported parametric starts

`kdensity` supports more than 20 built-in starts from the [univariateML](#) package, see `univariateML::univariateML_models` for a list. Densities with variable support, power, are not supported. The pareto density has its support fixed to $(1, \text{Inf})$. The options `uniform`, `constant` makes `kdensity` estimate a kernel density without parametric starts.

See Also

[kdensity\(\)](#); [kernels\(\)](#); [bandwidths\(\)](#)

Examples

```
start_exponential <- list(
  density = stats::dexp,
  estimator = function(data) {
    c(rate = 1 / mean(data))
  },
  support = c(0, Inf)
)

start_inverse_gaussian <- list(
  density = extraDistr::dwald,
  estimator = function(data) {
    c(
      mu = mean(data),
      lambda = mean(1 / data - 1 / mean(data))
    )
  },
  support = c(0, Inf)
)
```

`plot.kdensity`*Plot, Lines and Points Methods for Kernel Density Estimation*

Description

The plot method for `kdensity` objects.

Usage

```
## S3 method for class 'kdensity'  
plot(x, range = NULL, plot_start = FALSE, zero_line = TRUE, ...)  
  
## S3 method for class 'kdensity'  
lines(x, range = NULL, plot_start = FALSE, zero_line = TRUE, ...)  
  
## S3 method for class 'kdensity'  
points(x, range = NULL, plot_start = FALSE, zero_line = TRUE, ...)
```

Arguments

<code>x</code>	a <code>kdensity</code> object.
<code>range</code>	range of <code>x</code> values.
<code>plot_start</code>	logical; if <code>TRUE</code> , plots the parametric start instead of the kernel density estimate.
<code>zero_line</code>	logical; if <code>TRUE</code> , add a base line at $y = 0$.
<code>...</code>	further plotting parameters.

Value

None.

See Also

[kdensity\(\)](#)

Examples

```
## Using the data set "precip" to eye-ball the similarity between  
## a kernel fit, a parametric fit, and a kernel with parametric start fit.  
kde_gamma <- kdensity(precip, kernel = "gaussian", start = "gamma")  
kde <- kdensity(precip, kernel = "gaussian", start = "uniform")  
  
plot(kde_gamma, main = "Annual Precipitation in US Cities")  
lines(kde_gamma, plot_start = TRUE, lty = 2)  
lines(kde, lty = 3)  
rug(precip)
```

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