

Summary of recent updates to `spatstat`

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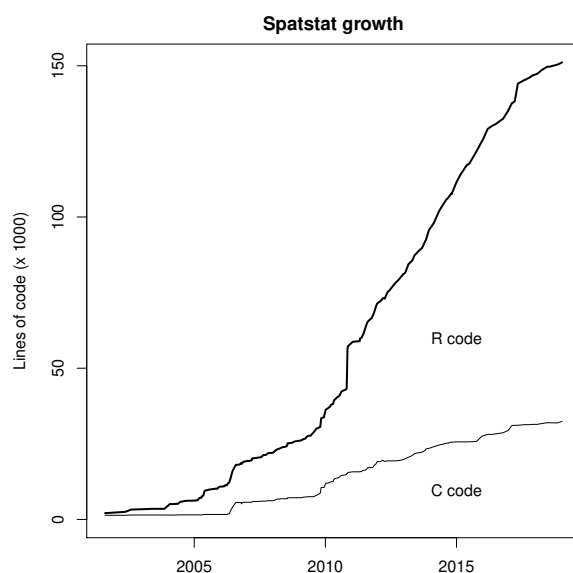
For `spatstat` version 1.58-2

This is a summary of changes that have been made to the `spatstat` package since the publication of the accompanying book [2].

The book [2], published in December 2015, covers everything in `spatstat` up to version 1.42-0, released in May 2015.

The `spatstat` package has grown by 29% since the book was published. This document summarises the most important changes.

The current version of `spatstat` is 1.58-2. It contains 618 new functions and 3 new datasets introduced after May 2015.



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1 spatstat is splitting into parts

`spatstat` is being split into sub-packages, to satisfy the requirements of CRAN. This should not affect the user: existing code will continue to work in the same way.

Typing `library(spatstat)` will load the familiar `spatstat` package which can be used as before.

1.1 The parts of spatstat

Currently there are three sub-packages, called `spatstat.utils`, `spatstat.data` and `spatstat`.

- The `spatstat` package contains the main code.
- The `spatstat.data` package now contains all the datasets for `spatstat`.
- The `spatstat.utils` package contains utility functions for `spatstat`.

Typing `library(spatstat)` will automatically load `spatstat.data` and silently “import” `spatstat.utils`.

To access the functions in `spatstat.utils` directly, you would need to type `library(spatstat.utils)`.

1.2 Extension packages

There are also extension packages which provide additional capabilities and must be loaded explicitly when you need them. Currently there are two extension packages:

- `spatstat.local` for local model-fitting,
- `spatstat.sphere` for analysing point patterns on a sphere.

2 Precise of all changes

Here is the text from the ‘overview’ sections of the News and Release Notes for each update.

- `spatstat` now Imports the package `spatstat.utils`.
- `spatstat` now requires the package `spatstat.data` which contains the datasets.
- `spatstat` now suggests the package `fftwtools`.
- Standard error calculation for inverse-distance weighting.
- Kernel estimate of intensity as a `function(x,y)`.
- Extract discrete and continuous components of a measure.
- Improvements and extensions to leverage and influence code.
- Plot a line segment pattern using line widths.
- Find connected components of each tile in a tessellation.
- Geometrical operations on `distfun` objects.
- Join vertices in a linear network.
- Distance map and contact distribution for rectangular structuring element.
- Lurking variable plot for models fitted to several point patterns.
- New dataset `cetaceans`.
- Gamma correction for colour maps and image plots.
- Class `units` has been renamed `unitname` to avoid package collision.
- More support for tessellations.
- Fixed longstanding bug in leverage and influence diagnostics.
- Improvements and bug fixes for leverage and influence diagnostics.
- Tighter bounding box for `psp`, `lpp`, `linnet` objects.
- Improved layout in `plot.solist`
- Tools to increase colour saturation.
- Connected components of a 3D point pattern.
- Accelerated computations on linear networks.
- Accelerated simulation of determinantal point processes.
- Improved printing of 3D point patterns.
- Minor corrections to handling of unitnames.
- Improvements to `ppm` and `update.ppm`.

- Correction to `lohboot`
- Numerous bug fixes for linear networks code.
- Now handles disconnected linear networks.
- Effect function is now available for all types of fitted model.
- Geometric-mean smoothing.
- A model can be fitted or re-fitted to a sub-region of data.
- New fast algorithm for kernel smoothing on a linear network.
- Leverage and influence diagnostics extended to Poisson/Gibbs models fitted by logistic composite likelihood.
- Two-stage Monte Carlo test.
- Dirichlet/Voronoi tessellation on a linear network.
- Thinning of point patterns on a linear network.
- More support for functions and tessellations on a linear network.
- Bandwidth selection for pair correlation function.
- Pooling operations improved.
- Operations on signed measures.
- Operations on lists of pixel images.
- Improved pixellation of point patterns.
- Stieltjes integral extended.
- Subset operators extended.
- Greatly accelerated `rmh` when using `nsave`
- Sufficient Dimension Reduction for point processes.
- Alternating Gibbs Sampler for point process simulation.
- New class of spatially sampled functions.
- ROC and AUC extended to other types of point patterns and models.
- More support for linear networks.
- More support for infinite straight lines.
- `spatstat` now depends on the packages `nlme` and `rpart`.
- Important bug fix in `linearK`, `linearpcf`
- Changed internal format of `linnet` and `lpp` objects.

- Faster computation in linear networks.
- Bias correction techniques.
- Bounding circle of a spatial object.
- Option to plot marked points as arrows.
- Kernel smoothing accelerated.
- Workaround for bug in some graphics drivers affecting image orientation.
- Non-Gaussian smoothing kernels.
- Improvements to inhomogeneous multitype K and L functions.
- Variance approximation for pair correlation function.
- Leverage and influence for multitype point process models.
- Functions for extracting components of vector-valued objects.
- Recursive-partition point process models.
- Minkowski sum, morphological dilation and erosion with any shape.
- Minkowski sum also applicable to point patterns and line segment patterns.
- Important bug fix in Smooth.ppp
- Important bug fix in spatial CDF tests.
- More bug fixes for replicated patterns.
- Simulate a model fitted to replicated point patterns.
- Inhomogeneous multitype F and G functions.
- Summary functions recognise `correction="all"`
- Leverage and influence code handles bigger datasets.
- More support for pixel images.
- Improved progress reports.
- New dataset `redwood3`
- Fixed namespace problems arising when spatstat is not loaded.
- Important bug fix in leverage/influence diagnostics for Gibbs models.
- Surgery with linear networks.
- Tessellations on a linear network.
- Laslett's Transform.
- Colour maps for point patterns with continuous marks are easier to define.

- Pair correlation function estimates can be pooled.
- Stipulate a particular version of a package.
- More support for replicated point patterns.
- More support for tessellations.
- More support for multidimensional point patterns and point processes.
- More options for one-sided envelopes.
- More support for model comparison.
- Convexifying operation.
- Subdivide a linear network.
- Penttinen process can be simulated (by Metropolis-Hastings or CFTP).
- Calculate the predicted variance of number of points.
- Accelerated algorithms for linear networks.
- Quadrat counting accelerated, in some cases.
- Simulation algorithms have been accelerated; simulation outcomes are *not* identical to those obtained from previous versions of **spatstat**.
- Determinantal point process models.
- Random-effects and mixed-effects models for replicated patterns.
- Dao-Genton test, and corresponding simulation envelopes.
- Simulated annealing and simulated tempering.
- spatstat colour tools now handle transparent colours.
- Improvements to `[]` and `subset` methods
- Extensions to kernel smoothing on a linear network.
- Support for one-dimensional smoothing kernels.
- Mark correlation function may include weights.
- Cross-correlation version of the mark correlation function.
- Penttinen pairwise interaction model.
- Improvements to simulation of Neyman-Scott processes.
- Improvements to fitting of Neyman-Scott models.
- Extended functionality for pixel images.
- Fitted intensity on linear network
- Triangulation of windows.
- Corrected an edge correction.

3 New datasets

The following datasets have been added to the package.

- `austates`: The states and large mainland territories of Australia represented as polygonal regions forming a tessellation.
- `redwood3`: a more accurate version of the `redwood` data.
- `cetaceans`: point patterns of whale and dolphin sightings.

4 New classes

- `ssf`: Class of spatially sampled functions.

5 New Functions

Following is a list of all the functions that have been added.

- `markmarkscatter`: Mark–mark scatterplot.
- `bw.CvL`: Cronie-van Lieshout bandwidth selection for density estimation.
- `subset.psp`: subset method for line segment patterns.
- `densityfun`, `densityfun.ppp`: Compute a kernel estimate of intensity of a point pattern and return it as a function of spatial location.
- `as.im.densityfun`: Convert `function(x,y)` to a pixel image.
- `measureDiscrete`, `measureContinuous`: Extract the discrete and continuous components of a measure.
- `connected.tess`: Find connected components of each tile in a tessellation and make a new tessellation composed of these pieces.
- `dffit.ppm`: Effect change diagnostic DFFIT for spatial point process models.
- `shift.distfun`, `rotate.distfun`, `reflect.distfun`, `flipxy.distfun`, `affine.distfun`, `scalardilate.distfun`: Methods for geometrical operations on `distfun` objects.
- `rescale.distfun`: Change the unit of length in a `distfun` object.
- `plot.indicfun`: Plot method for indicator functions created by `as.function.owin`.
- `Smooth.leverage.ppm`, `Smooth.influence.ppm`: Smooth a leverage function or an influence measure.
- `integral.leverage.ppm`, `integral.influence.ppm`: Compute the integral of a leverage function or an influence measure.
- `mean.leverage.ppm`: Compute the mean value of a leverage function.
- `rectdistmap`: Distance map using rectangular metric.

- `rectcontact`: Contact distribution function using rectangular structuring element.
- `joinVertices`: Join specified vertices in a linear network.
- `summary.ssf`: Summary method for a spatially sampled function (class `ssf`).
- `unstack.tess`: Given a tessellation with multiple columns of marks, take the columns one at a time, and return a list of tessellations, each carrying only one of the original columns of marks.
- `contour.leverage.ppm`: Method for `contour` for leverage functions of class `leverage.ppm`
- `lurking`: New generic function for lurking variable plots.
- `lurking.ppp`, `lurking.ppm`: These are equivalent to the original function `lurking`. They are now methods for the new generic `lurking`.
- `lurking.mppm`: New method for class `mppm`. Lurking variable plot for models fitted to several point patterns.
- `print.lurk`: Prints information about the object returned by the function `lurking` representing a lurking variable plot.
- `model.matrix.mppm`: Method for `model.matrix` for models of class `mppm`.
- `test.crossing.psp`, `test.selfcrossing.psp`: Previously undocumented functions for testing whether segments cross.
- `to.saturated`: Convert a colour value to the corresponding fully-saturated colour.
- `intensity.psp`: Compute the average total length of segments per unit area.
- `boundingbox.psp`: Bounding box for line segment patterns. This produces a tighter bounding box than the previous default behaviour.
- `boundingbox.lpp`: Bounding box for point patterns on a linear network. This produces a tighter bounding box than the previous default behaviour.
- `boundingbox.linnet`: Bounding box for a linear network. This produces a tighter bounding box than the previous default behaviour.
- `"Frame<-".default`: New default method for assigning bounding frame to a spatial object.
- `connected.pp3`: Connected components of a 3D point pattern.
- `colouroutputs`, `"colouroutputs<-"`: Extract or assign colour values in a colour map. (Documented a previously-existing function)
- `fitin.profilepl`: Extract the fitted interaction from a model fitted by profile likelihood.
- `[<-".linim`: Subset assignment method for pixel images on a linear network.
- `nnfromvertex`: Given a point pattern on a linear network, find the nearest data point from each vertex of the network.
- `tile.lengths`: Calculate the length of each tile in a tessellation on a network.
- `text.ppp`, `text.lpp`, `text.psp`: Methods for `text` for spatial patterns.

- `as.data.frame.envelope`: Extract function data from an envelope object, including the functions for the simulated data ('simfuns') if they were saved.
- `is.connected`, `is.connected.default`, `is.connected.linnet`: Determines whether a spatial object consists of one topologically connected piece, or several pieces.
- `is.connected.ppp`: Determines whether a point pattern is connected after all pairs of points closer than distance R are joined.
- `hist.funxy`: Histogram of values of a spatial function.
- `model.matrix.ippm`: Method for `model.matrix` which allows computation of regular and irregular score components.
- `harmonise.msr`: Convert several measures (objects of class `msr`) to a common quadrature scheme.
- `bits.test`: Balanced Independent Two-Stage Monte Carlo test, an improvement on the Dao-Genton test.
- `lineardirichlet`: Computes the Dirichlet-Voronoi tessellation associated with a point pattern on a linear network.
- `domain.lintess`, `domain.linfun`: Extract the linear network from a `lintess` or `linfun` object.
- `summary.lintess`: Summary of a tessellation on a linear network.
- `clicklpp`: Interactively add points on a linear network.
- `envelopeArray`: Generate an array of envelopes using a function that returns `fasp` objects.
- `bw.pcf`: Bandwidth selection for pair correlation function.
- `grow.box3`: Expand a three-dimensional box.
- `hexagon`, `regularpolygon`: Create regular polygons.
- `Ops.msr`: Arithmetic operations for measures.
- `Math.imlist`, `Ops.imlist`, `Summary.imlist`, `Complex.imlist`: Arithmetic operations for lists of pixel images.
- `measurePositive`, `measureNegative`, `measureVariation`, `totalVariation`: Positive and negative parts of a measure, and variation of a measure.
- `as.function.owin`: Convert a spatial window to a `function(x,y)`, the indicator function.
- `as.function.ssf`: Convert an object of class `ssf` to a `function(x,y)`
- `as.function.leverage.ppm`: Convert an object of class `leverage.ppm` to a `function(x,y)`
- `sdr`, `dimhat`: Sufficient Dimension Reduction for point processes.
- `simulate.rhohat`: Simulate a Poisson point process with the intensity estimated by `rhohat`.
- `rlpp`: Random points on a linear network with a specified probability density.

- `cut.lpp`: Method for `cut` for point patterns on a linear network.
- `has.close`: Faster way to check whether a point has a close neighbour.
- `psib`: Sibling probability (index of clustering strength in a cluster process).
- `rags`, `ragsAreaInter`, `ragsMultiHard`: Alternating Gibbs Sampler for point processes.
- `bugfixes`: List all bug fixes in recent versions of a package.
- `ssf`: Create a spatially sampled function
- `print.ssf`, `plot.ssf`, `contour.ssf`, `image.ssf`: Display a spatially sampled function
- `as.im.ssf`, `as.ppp.ssf`, `marks.ssf`, `marks<-ssf`, `unmark.ssf`, `[.ssf`, `with.ssf`: Manipulate data in a spatially sampled function
- `Smooth.ssf`: Smooth a spatially sampled function
- `integral.ssf`: Approximate integral of spatially sampled function
- `roc.kppm`, `roc.lppm`, `roc.lpp`: Methods for `roc` for fitted models of class `"kppm"` and `"lppm"` and point patterns of class `"lpp"`
- `auc.kppm`, `auc.lppm`, `auc.lpp`: Methods for `auc` for fitted models of class `"kppm"` and `"lppm"` and point patterns of class `"lpp"`
- `timeTaken`: Extract the timing data from a `"timed"` object or objects.
- `rotate.inflin`, `shift.inflin`, `reflect.inflin`, `flipxy.inflin`: Geometrical transformations for infinite straight lines.
- `whichhalfplane`: Determine which side of an infinite line a point lies on.
- `matrixpower`, `matrixsqrt`, `matrixinvsqrt`: Raise a matrix to any power.
- `points.lpp`: Method for `points` for point patterns on a linear network.
- `pairs.linim`: Pairs plot for images on a linear network.
- `closetriples`: Find close triples of points.
- `anyNA.im`: Method for `anyNA` for pixel images.
- `bc`: Bias correction (Newton-Raphson) for fitted model parameters.
- `rex`: Richardson extrapolation for numerical integrals and statistical model parameter estimates.
- `boundingcircle`, `boundingcentre`: Find the smallest circle enclosing a window or point pattern.
- `[.linim`: Subset operator for pixel images on a linear network.
- `mean.linim`, `median.linim`, `quantile.linim`: The mean, median, or quantiles of pixel values in a pixel image on a linear network.
- `weighted.median`, `weighted.quantile`: Median or quantile of numerical data with associated weights.

- "`.linim`": Subset operator for pixel images on a linear network.
- `mean.linim`, `median.linim`, `quantile.linim`: The mean, median, or quantiles of pixel values in a pixel image on a linear network.
- `boundingcircle`, `boundingcentre`: Smallest circle enclosing a spatial object.
- `split.msr`: Decompose a measure into parts.
- `unstack.msr`: Decompose a vector-valued measure into its component measures.
- `unstack.ppp`, `unstack.psp`, `unstack.lpp`: Given a spatial pattern with several columns of marks, separate the columns and return a list of spatial patterns, each having only one column of marks.
- `kernel.squint`: Integral of squared kernel, for the kernels used in density estimation.
- `as.im.data.frame`: Build a pixel image from a data frame of coordinates and pixel values.
- `covering`: Cover a window using discs of a given radius.
- `dilationAny`, `erosionAny`, `%(-)%`: Morphological dilation and erosion by any shape.
- `FmultiInhom`, `GmultiInhom` Inhomogeneous multitype/marked versions of the summary functions `Fest`, `Gest`.
- `kernel.moment` Moment or incomplete moment of smoothing kernel.
- `MinkowskiSum`, `%(+)%`: Minkowski sum of two windows: `A %(+)% B`, or `MinkowskiSum(A,B)`
- `nobjects`: New generic function for counting the number of 'things' in a dataset. There are methods for `ppp`, `ppx`, `psp`, `tess`.
- `parameters.interact`, `parameters.fii`: Extract parameters from interpoint interactions. (These existing functions are now documented.)
- `ppmInfluence`: Calculate leverage `ppm`, influence `ppm` and `dfbetas.ppm` efficiently.
- `rppm`, `plot.rppm`, `predict.rppm`, `prune.rppm`: Recursive-partition point process models.
- `simulate.mppm` Simulate a point process model fitted to replicated point patterns.
- `update.interact`: Update the parameters of an interpoint interaction. [This existing function is now documented.]
- `where.max`, `where.min` Find the spatial location(s) where a pixel image achieves its maximum or minimum value.
- `compileK`, `compilepcf`: make a K function or pair correlation function given the pairwise distances and their weights. [These existing internal functions are now documented.]
- `laslett`: Laslett's Transform.
- `lintess`: Tessellation on a linear network.
- `divide.linnet`: Divide a linear network into pieces demarcated by a point pattern.

- `insertVertices`: Insert new vertices in a linear network.
- `thinNetwork`: Remove vertices and/or segments from a linear network etc.
- `connected.linnet`: Find connected components of a linear network.
- `nvertices`, `nvertices.linnet`, `nvertices.owin`: Count the number of vertices in a linear network or vertices of the boundary of a window.
- `as.data.frame.linim`, `as.data.frame.linfun`: Extract a data frame of spatial locations and function values from an object of class `linim` or `linfun`.
- `as.linfun`, `as.linfun.linim`, `as.linfun.lintess`: Convert other kinds of data to a `linfun` object.
- `requireversion`: Require a particular version of a package (for use in stand-alone R scripts).
- `as.function.tess`: Convert a tessellation to a `function(x,y)`. The function value indicates which tile of the tessellation contains the point (x,y) .
- `tileindex`: Determine which tile of a tessellation contains a given point (x,y) .
- `persp.leverage.ppm`: Method for persp plots for objects of class `leverage.ppm`
- `AIC.mppm`, `extractAIC.mppm`: AIC for point process models fitted to replicated point patterns.
- `nobs.mppm`, `terms.mppm`, `getCall.mppm`: Methods for point process models fitted to replicated point patterns.
- `rPenttinen`: Simulate the Penttinen process using perfect simulation.
- `varcount`: Given a point process model, compute the predicted variance of the number of points falling in a window.
- `inside.boxx`: Test whether multidimensional points lie inside a specified multidimensional box.
- `lixellate`: Divide each segment of a linear network into smaller segments.
- `nsegments.linnet`, `nsegments.lpp`: Count the number of line segments in a linear network.
- `grow.boxx`: Expand a multidimensional box.
- `deviance.ppm`, `deviance.lppm`: Deviance for a fitted point process model.
- `pseudoR2`: Pseudo-R-squared for a fitted point process model.
- `tiles.empty` Checks whether each tile of a tessellation is empty or nonempty.
- `summary.linim`: Summary for a pixel image on a linear network.
- Determinantal Point Process models:
 - `dppm`: Fit a determinantal point process model.
 - `fitted.dppm`, `predict.dppm`, `intensity.dppm`: prediction for a fitted determinantal point process model.
 - `Kmodel.dppm`, `pcfmodel.dppm`: Second moments of a determinantal point process model.

- `rdpp`, `simulate.dppm`: Simulation of a determinantal point process model.
 - `logLik.dppm`, `AIC.dppm`, `extractAIC.dppm`, `nobs.dppm`: Likelihood and AIC for a fitted determinantal point process model.
 - `print.dppm`, `reach.dppm`, `valid.dppm`: Basic information about a `dpp` model.
 - `coef.dppm`, `formula.dppm`, `print.dppm`, `terms.dppm`, `labels.dppm`, `model.frame.dppm`, `model.matrix.dppm`, `model.images.dppm`, `is.stationary.dppm`, `reach.dppm`, `unitname.dppm`, `unitname<- .dppm`, `Window.dppm`: Various methods for `dppm` objects.
 - `parameters.dppm`: Extract meaningful list of model parameters.
 - `objsurf.dppm`: Objective function surface of a `dppm` object.
 - `residuals.dppm`: Residual measure for a `dppm` object.
- Determinantal Point Process model families:
 - `dppBessel`, `dppCauchy`, `dppGauss`, `dppMatern`, `dppPowerExp`: Determinantal Point Process family functions.
 - `detpointprocfamilyfun`: Create a family function.
 - `update.detpointprocfamily`: Set parameter values in a determinantal point process model family.
 - `simulate.dppm`: Simulation.
 - `is.stationary.detpointprocfamily`, `intensity.detpointprocfamily`, `Kmodel.detpointprocfamily`, `pcfmodel.detpointprocfamily`: Moments.
 - `dim.detpointprocfamily`, `dppapproxkernel`, `dppapproxpcf`, `dppeigen`, `dppkernel`, `dppparbounds`, `dppspecdenrange`, `dppspecden`: Helper functions.
 - `dg.envelope`: Simulation envelopes corresponding to Dao-Genton test.
 - `dg.progress`: Progress plot (envelope representation) for the Dao-Genton test.
 - `dg.sigtrace`: significance trace for the Dao-Genton test.
 - `markcrosscorr`: Mark cross-correlation function for point patterns with several columns of marks.
 - `rtemper`: Simulated annealing or simulated tempering.
 - `rgb2hsva`: Convert RGB to HSV data, like `rgb2hsv`, but preserving transparency.
 - `superimpose.ppplist`, `superimpose.splitppp`: New methods for 'superimpose' for lists of point patterns.
 - `dkernel`, `pkernel`, `qkernel`, `rkernel`: Probability density, cumulative probability, quantiles and random generation from distributions used in basic one-dimensional kernel smoothing.
 - `kernel.factor`: Auxiliary calculations for one-dimensional kernel smoothing.
 - `spatdim`: Spatial dimension of any object in the `spatstat` package.
 - `as.boxx`: Convert data to a multi-dimensional box.
 - `intensity.ppx`: Method for `intensity` for multi-dimensional space-time point patterns.

- `fourierbasis`: Evaluate Fourier basis functions in any number of dimensions.
- `valid`: New generic function, with methods `valid.ppm`, `valid.lppm`, `valid.dppm`.
- `emend`, `emend.ppm`, `emend.lppm`: New generic function with methods for `ppm` and `lppm`. `emend.ppm` is equivalent to `project.ppm`.
- `Penttinen`: New pairwise interaction model.
- `quantile.density`: Calculates quantiles from kernel density estimates.
- `CDF.density`: Calculates cumulative distribution function from kernel density estimates.
- `triangulate.owin`: decompose a spatial window into triangles.
- `fitted.lppm`: fitted intensity values for a point process on a linear network.
- `parameters`: Extract all parameters from a fitted model.

6 Alphabetical list of changes

Here is a list of all changes made to existing functions, listed alphabetically.

- `affine.owin`: Allows transformation matrix to be singular, if the window is polygonal.
- `anova.mppm`: Now handles Gibbs models, and performs the adjusted composite likelihood ratio test. New argument `fine`.
- `as.function.tess`: New argument `values` specifies the function values.
- `as.im.distfun`: New argument `approx` specifies the choice of algorithm.
- `as.im.function`: New argument `strict`.
- `as.im.leverage.ppm`: New argument `what`.
- `as.im.nnfun`: New argument `approx` chooses between a fast, approximate algorithm and a slow, exact algorithm.
- `as.im.smoothfun`: New argument `approx` chooses between a fast, approximate algorithm and a slow, exact algorithm.
- `as.layered`: Default method now handles a (vanilla) list of spatial objects.
- `as.linfun.lintess`:
 - New argument `values` specifies the function value for each tile.
 - New argument `navalue`.
- `as.linim.default`: New argument `delta` controls spacing of sample points in internal data.
- `as.linnet.psp`: If the line segment pattern has marks, then the resulting linear network also carries these marks in the `$lines` component.
- `as.owin.default`:

- Now refuses to convert a `box3` to a two-dimensional window.
- Now accepts a structure with entries named `xmin,xmax, ymin, ymax` in any order. This handles objects of class `bbox` in the `sf` package.
- Now detects objects of class `SpatialPolygons` and issues a more helpful error message.
- `as.owin.data.frame`: New argument `step`
- `as.polygonal`:
 - Can now repair errors in polygon data, if `repair=TRUE`.
 - Accelerated when `w` is a pixel mask.
- `as.solist`: The argument `x` can now be a spatial object; `as.solist(cells)` is the same as `solist(cells)`.
- `bdist.pixels`: Accelerated for polygonal windows. New argument `method`.
- `bdist.points`: Accelerated for polygonal windows.
- `bind.fv`: New argument `clip`.
- `bw.diggle`, `bw.ppl`, `bw.relrisk`, `bw.smoothppp`: These functions now extract and store the name of the unit of length from the point pattern dataset. When the bandwidth selection criterion is plotted, the name of the unit of length is shown on the x-axis.
- `bw.ppl`: New arguments `weights` and `sigma`.
- `bw.scott`: the two bandwidth values in the result now have names `sigma.x` and `sigma.y`.
- `cdf.test`:
 - Calculations are more robust against numerical rounding effects.
 - The methods for classes `ppp`, `ppm`, `lpp`, `lppm`, `slrm` have a new argument `interpolate`.
 - Monte Carlo test runs much faster.
 - More jittering is applied when `jitter=TRUE`. Warnings about tied values should not occur any more.
- `cdf.test.mppm`:
 - Now handles Gibbs models.
 - Now recognises `covariate="x"` or `"y"`.
- `clarkevans`: The argument `correction="all"` is now recognised: it selects all the available options. [This is also the default.]
- `clickpoly`: The polygon is now drawn progressively as the user clicks new vertices.
- `closepairs.ppp`: New argument `periodic`.
- `closepairs.ppp`, `closepairs.pp3`:
 - New arguments `distinct` and `neat` allow more options.
 - Argument `ordered` has been replaced by `twice` (but `ordered` is still accepted, with a warning).

- Performance improved (computation time and memory requirements reduced.) This should improve the performance of many functions in `spatstat`.
- `closepairs.pp3`: Argument `what` can take the value "ijd"
- `clusterset`: Improved behaviour.
- `clusterfit`:
 - New argument `algorithm` specifies the choice of optimisation algorithm.
 - Changed precedence rule for handling the algorithm parameters in the minimum contrast algorithm. Individually-named arguments `q`, `p`, `rmax`, `rmin` now take precedence over entries with the same names in the list `ctrl`.
- `collapse.fv`: This is now treated as a method for the `nlme` generic `collapse`. Its syntax has been adjusted slightly.
- `connected.im`: Now handles a logical-valued image properly. Arguments `...` now determine pixel resolution.
- `connected.owin`: Arguments `...` now determine pixel resolution.
- `contour.im`: New argument `col` specifies the colour of the contour lines. If `col` is a colour map, then the contours are drawn in different colours.
- `convolve.im`: the name of the unit of length is preserved.
- `crossing.psp`: New argument `details` gives more information about the intersections between the segments.
- `crosspairs.pp3`: Argument `what` can take the value "ijd"
- `cut.ppp`: Argument `z` can be "x" or "y" indicating one of the spatial coordinates.
- `dclf.test`, `mad.test`, `dclf.progress`, `mad.progress`, `dclf.sigtrace`, `mad.sigtrace`, `dg.progress`, `dg.sigtrace`:
 - New argument `clamp` determines the test statistic for one-sided tests.
 - New argument `rmin` determines the left endpoint of the test interval.
 - New argument `leaveout` specifies how to calculate discrepancy between observed and simulated function values.
 - New argument `scale` allows summary function values to be rescaled before the comparison is performed.
 - New argument `interpolate` supports interpolation of p -value.
 - New argument `interpolate` supports interpolation of critical value of test.
- `default.rmhcontrol`, `default.rmhexpand`: New argument `w`.
- `density.lpp`:
 - New fast algorithm (up to 1000 times faster) for the default case where `kernel="gaussian"` and `continuous=TRUE`. Generously contributed by Greg McSwiggan.
 - Fast algorithm has been further accelerated.

- New argument `kernel` specifies the smoothing kernel. Any of the standard one-dimensional smoothing kernels can be used.
 - Now supports both the ‘equal-split continuous’ and ‘equal-split discontinuous’ smoothers. New argument `continuous` determines the choice of smoother.
 - New arguments `weights` and `old`.
- `density.ppp`:
 - A non-Gaussian kernel can now be specified using the argument `kernel`.
 - Argument `weights` can now be a pixel image.
 - Infinite bandwidth `sigma=Inf` is supported.
 - Accelerated by about 30% when `at="pixels"`.
 - Accelerated by about 15% in the case where `at="points"` and `kernel="gaussian"`.
 - Accelerated in the cases where weights are given or `diggle=TRUE`.
 - New argument `verbose`.
 - `density.psp`:
 - New argument `method`.
 - Accelerated by 1 to 2 orders of magnitude.
 - `dfbetas.ppm`:
 - For Gibbs models, memory usage has been dramatically reduced, so the code can handle larger datasets and finer quadrature schemes.
 - Increased the default resolution of the pixel images. Spatial resolution can now be controlled by the arguments `dimyx`, `eps`.
 - `diagnose.ppm`: Infinite values of `rbord` are now ignored and treated as zero. This ensures that `diagnose.ppm` has a sensible default when the fitted model has infinite reach.
 - `diagnose.ppm`, `plot.diagppm`: New arguments `col.neg`, `col.smooth` control the colour maps.
 - `dilation.ppp`: Improved geometrical accuracy. Now accepts arguments to control resolution of polygonal approximation.
 - `discs`:
 - Now accepts a single numeric value for `radii`.
 - New argument `npoly`.
 - Accelerated in some cases.
 - `distfun`: When the user calls a distance function that was created by `distfun`, the user may now give a `ppp` or `lpp` object for the argument `x`, instead of giving two coordinate vectors `x` and `y`.
 - `dppm`: Changed precedence rule for handling the algorithm parameters in the minimum contrast algorithm. Individually-named arguments `q`, `p`, `rmax`, `rmin` now take precedence over entries with the same names in the list `ctrl`.

- `edge.Trans`: New argument `gW` for efficiency.
- `effectfun`: Now works for `ppm`, `kppm`, `lppm`, `dppm`, `rppm` and `profilepl` objects.
- `envelope`:
 - New argument `clamp` gives greater control over one-sided envelopes.
 - New argument `funargs`
 - New argument `scale` allows global envelopes to have width proportional to a specified function of r , rather than constant width.
 - New argument `funYargs` contains arguments to the summary function when applied to the data pattern only.
- `envelope.lpp`, `envelope.lppm`: New arguments `fix.n` and `fix.marks` allow envelopes to be computed using simulations conditional on the observed number of points.
- `eval.im`: New argument `warn`.
- `eval.linim`: New argument `warn`.
- `ewcdf`: Argument `weights` can now be `NULL`.
- `Fest`: Additional checks for errors in input data.
- `fitted.lppm`: New argument `leaveoneout` allows leave-one-out computation of fitted value.
- `fitted.ppm`:
 - New option, `type="link"`.
 - New argument `ignore.hardcore`.
- `funxy`: When the user calls a function that was created by `funxy`, the user may now give a `ppp` or `lpp` object for the argument `x`, instead of giving two coordinate vectors `x` and `y`.
- `Geyer`: The saturation parameter `sat` can now be less than 1.
- `grow.rectangle`: New argument `fraction`.
- `Hest`:
 - Argument `X` can now be a pixel image with logical values.
 - New argument `W`. [Based on code by Kassel Hingee.]
 - Additional checks for errors in input data.
- `hist.im`: New argument `xname`.
- `identify.psp`: Improved placement of labels. Arguments can be passed to `text.default` to control the plotting of labels.
- `idw`: Standard errors can now be calculated by setting `se=TRUE`.
- `imcov`: the name of the unit of length is preserved.
- `im.apply`:

- Computation accelerated
- New argument `fun.handles.na`
- New argument `check`
- `influence.ppm`: For Gibbs models, memory usage has been dramatically reduced, so the code can handle larger datasets and finer quadrature schemes.
- `integral.linfun`:
 - New argument `delta` controls step length of approximation to integral.
 - Argument `domain` can be a tessellation.
- `integral.linim`: Argument `domain` can be a tessellation.
- `integral.ssf`: Argument `domain` can be a tessellation.
- `intensity.ppm`: Intensity approximation is now implemented for area-interaction model, and Geyer saturation model.
- `interp.im`: New argument `bilinear`.
- `ippm`:
 - Accelerated.
 - The internal format of the result has been extended slightly.
 - Improved defaults for numerical algorithm parameters.
- `Kcross.inhom`, `Kdot.inhom`, `Kmulti.inhom`: These functions now allow intensity values to be given by a fitted point process model. New arguments `update`, `leaveoneout`, `lambdaX`.
- `Kest`: Accelerated computation (for translation and rigid corrections) when window is an irregular shape.
- `Kest.fft`: Now has ... arguments allowing control of spatial resolution.
- `Kinhom`:
 - New argument `ratio`.
 - Stops gracefully if `lambda` contains any zero values.
- `kppm`:
 - Fitting a model with `clusters="LGCP"` no longer requires the package `RandomFields` to be loaded explicitly.
 - New argument `algorithm` specifies the choice of optimisation algorithm.
 - Left hand side of formula can now involve entries in the list `data`.
 - refuses to fit a log-Gaussian Cox model with anisotropic covariance.
 - A warning about infinite values of the summary function no longer occurs when the default settings are used. Also affects `mincontrast`, `cauchy.estpcf`, `lgcp.estpcf`, `mat-clust.estpcf`, `thomas.estpcf`, `vargamma.estpcf`.

- Changed precedence rule for handling the algorithm parameters in the minimum contrast algorithm. Individually-named arguments `q`, `p`, `rmax`, `rmin` now take precedence over entries with the same names in the list `ctrl`.
- Improved printed output.
- `Lcross.inhom`, `Ldot.inhom`: These functions now allow intensity values to be given by a fitted point process model. New arguments `update`, `leaveoneout`, `lambdaX`.
- `lengths.psp`: New argument `squared`.
- `leverage.ppm`:
 - For Gibbs models, memory usage has been dramatically reduced, so the code can handle larger datasets and finer quadrature schemes.
 - Increased the default resolution of the pixel images. Spatial resolution can now be controlled by the arguments `dimyx`, `eps`.
- `leverage.ppm`, `influence.ppm`, `dfbetas.ppm`:
 - These methods now work for models that were fitted by logistic composite likelihood (`method='logi'`).
 - Computation has been vastly accelerated for models with Geyer interaction fitted using isotropic or translation edge corrections.
 - Faster computation in many cases.
 - Virtually all models and edge corrections are now supported, using a “brute force” algorithm. This can be slow in some cases.
- `linearK`, `linearpcf` and relatives:
 - substantially accelerated.
 - ratio calculations are now supported.
 - new argument `ratio`.
- `linearKinhom`: new argument `normpower`.
- `linearKinhom`, `linearpcfinhom`:
 - Changed behaviour when `lambda` is a fitted model.
 - New arguments `update` and `leaveoneout`.
- `linearpcf`: new argument `normpower`.
- `linim`:
 - The image `Z` is now automatically restricted to the network.
 - New argument `restrict`.
- `linnet`:
 - The internal format of a `linnet` (linear network) object has been changed. Existing datasets of class `linnet` are still supported. However, computation will be faster if they are converted to the new format. To convert a `linnet` object `L` to the new format, use `L <- as.linnet(L)`.

- If the argument `edges` is given, then this argument now determines the ordering of the sequence of line segments. For example, the *i*-th row of `edges` specifies the *i*-th line segment in `as.psp(L)`.
 - New argument `warn`.
 - When argument `edges` is specified, the code now checks whether any edges are duplicated.
- `lintess`: Argument `df` can be missing or `NULL`, resulting in a tessellation with only one tile.
- `logLik.ppm`:
 - New argument `absolute`.
 - The warning about pseudolikelihood ('log likelihood not available') is given only once, and is not repeated in subsequent calls, within a spatstat session.
- `logLik.mppm`: new argument `warn`.
- `lohboot`: Algorithm has been corrected and extended thanks to Christophe Biscio and Rasmus Waagepetersen. New arguments `block`, `basicboot`, `Vcorrection`.
- `lpp`:
 - The internal format of an `lpp` object has been changed. Existing datasets of class `lpp` are still supported. However, computation will be faster if they are converted to the new format. To convert an `lpp` object `X` to the new format, use `X <- as.lpp(X)`.
 - `X` can be missing or `NULL`, resulting in an empty point pattern.
- `lpp`, `as.lpp`: These functions now handle the case where coordinates `seg` and `tp` are given but `x` and `y` are missing.
- `lppm`:
 - New argument `random` controls placement of dummy points.
 - Computation accelerated.
- `lurking.ppm`: accelerated.
- `markcorr`: New argument `weights` allows computation of the weighted version of the mark correlation function.
- `mppm`:
 - Now handles models with a random effect component. (This is covered in [2, Chap. 16].)
 - New argument `random` is a formula specifying the random effect. (This is covered in [2, Chap. 16].)
 - Performs more checks for consistency of the input data.
 - New arguments `gcontrol` and `reltol.pql` control the fitting algorithm.
- `msr`: Infinite and `NA` values are now detected (if `check=TRUE`) and are reset to zero, with a warning.
- `nbfires`:
 - the unit of length for the coordinates is now specified in this dataset.

- This dataset now includes information about the different land and sea borders of New Brunswick.
- `nndist.lpp`, `nnwhich.lpp`, `nncross.lpp`, `distfun.lpp`: New argument `k` allows computation of k -th nearest point. Computation accelerated.
- `nnfun.lpp`: New argument `k`.
- `padimage`: New argument `W` allows an image to be padded out to fill any window.
- `pcf.ppp`:
 - New argument `close` for advanced use.
 - New argument `ratio` allows several estimates of `pcf` to be pooled.
 - Now calculates an analytic approximation to the variance of the estimate of the pair correlation function (when `var.approx=TRUE`).
 - Now returns the smoothing bandwidth used, as an attribute of the result.
 - New argument `close` for advanced use.
- `pcfinhom`:
 - New argument `close` for advanced use.
 - Default behaviour is changed when `lambda` is a fitted model. The default is now to re-fit the model to the data before computing `pcf`. New arguments `update` and `leaveoneout` control this.
 - New argument `close` for advanced use.
 - Now handles `correction="good"`
- `persp.funxy`: Improved z -axis label.
- `pixellate.ppp`:
 - If the pattern is empty, the result is an integer-valued image (by default) for consistency with the results for non-empty patterns.
 - Accelerated in the case where weights are given.
 - New arguments `fractional` and `preserve` for more accurate discretisation.
 - New argument `savemap`.
- `plot.anylist`:
 - If a list entry `x[[i]]` belongs to class `"anylist"`, it will be expanded so that each entry `x[[i]][[j]]` will be plotted as a separate panel.
 - New arguments `panel.begin.args`, `panel.end.args`
 - Result is now an (invisible) list containing the result from executing the plot of each panel.
- `plot.colourmap`: New argument `increasing` specifies whether the colours are displayed in order left-to-right/bottom-to-top. Changed default behaviour for discrete colour maps when `vertical=FALSE`.
- `plot.im`:

- Now handles complex-valued images.
 - New argument `workaround` to avoid a bug in some MacOS device drivers that causes the image to be displayed in the wrong spatial orientation.
 - The number of tick marks in the colour ribbon can now be controlled using the argument `nint` in `ribargs`.
 - Improved behaviour when all pixel values are `NA`.
 - Improved handling of tickmarks on colour ribbon.
- `plot.imlist`: Result is now an (invisible) list containing the result from executing the plot of each panel.
 - `plot.influence.ppm`: New argument `multiplot`.
 - `plot.kppm`:
 - New arguments `pause` and `xname`.
 - The argument `what="all"` is now recognised: it selects all the available options. [This is also the default.]
 - `plot.leverage.ppm`:
 - New arguments `multiplot` and `what`.
 - A contour line showing the average value of leverage is now drawn on the colour ribbon, as well as on the main image. New argument `args.contour`.
 - `plot.linfun`:
 - Now passes arguments to the function being plotted.
 - A scale bar is now plotted when `style="width"`.
 - New argument `legend`.
 - The return value has a different format.
 - `plot.linim`:
 - The return value has a different format.
 - A scale bar is now plotted when `style="width"`.
 - When `style="width"`, negative values are plotted in red (by default). New argument `negative.args` controls this.
 - New argument `zlim` specifies the range of values to be mapped.
 - `plot.lintess`: Improved plot method, with more options.
 - `plot.lpp`:
 - New argument `show.network`.
 - For a point pattern with continuous marks (“real numbers”) the colour arguments `cols`, `fg`, `bg` can now be vectors of colour values, and will be used to determine the default colour map for the marks.
 - `plot.mppm`: New argument `se`.

- `plot.msr`:
 - Now handles multitype measures.
 - New argument `multiplot`.
 - New argument `massthresh`.
 - New arguments `equal.markscale` and `equal.ribbon`.
- `plot.onearrow`: Graphical parameters, specified when the object was created, are now taken as the defaults for graphical parameters to the plot.
- `plot.psp`:
 - Segments can be plotted with widths proportional to their mark values.
 - New argument `style`.
- `plot.pp3`: New arguments `box.front`, `box.back` control plotting of the box.
- `plot.ppp`:
 - The default colour for the points is now a transparent grey, if this is supported by the plot device.
 - For a point pattern with continuous marks (“real numbers”) the colour arguments `cols`, `fg`, `bg` can now be vectors of colour values, and will be used to determine the default colour map for the marks.
 - Now recognises graphics parameters for text, such as `family` and `srt`
 - When `clipwin` is given, any parts of the boundary of the window of `x` that lie inside `clipwin` will also be plotted.
 - Improved placement of symbol map legend when argument `symap` is given.
- `plot.tess`:
 - This plot method can now fill each tile with a different colour.
 - New arguments `do.col`, `values`, `col` and `ribargs`. Old argument `col` has been renamed `border` for consistency.
 - Now generates a separate plot panel for each column of marks, if `do.col=TRUE`.
 - New argument `multiplot`.
- `plot.profilepl`, `plot.quadratcount`, `plot.quadratcount`, `plot.tess`: Now recognise graphics parameters for text, such as `family` and `srt`
- `plot.solist`:
 - New arguments `panel.begin.args`, `panel.end.args`
 - Result is now an (invisible) list containing the result from executing the plot of each panel.
- `plot.studpermutest`: This existing function now has a help file.
- `plot.symbolmap`: New argument `nsymbols` controls the number of symbols plotted.
- `ponderosa`: In this installed dataset, the function `ponderosa.extra$plotit` has changed slightly (to accommodate the dependence on the package `spatstat.utils`).

- `polynom`: This function now has a help file.
- `pool.fv`:
 - The default plot of the pooled function no longer includes the variance curves.
 - New arguments `relabel` and `variance`.
- `pool.rat`: New arguments `weights`, `relabel` and `variance`.
- `ppm`:
 - Argument `interaction` can now be a function that makes an interaction, such as `Poisson`, `Hardcore`, `MultiHard`.
 - Argument `subset` can now be a window (class `"owin"`) specifying the sub-region of data to which the model should be fitted.
- `ppm.ppp`, `ppm.quad`:
 - New argument `emend`, equivalent to `project`.
 - New arguments `subset` and `clipwin`.
- `ppmInfluence`: The result now belongs to class `ppmInfluence`, for which there are methods for `leverage`, `influence`, `dfbetas` which extract the desired component.
- `ppp`:
 - New argument `checkdup`.
 - If the coordinate vectors `x` and `y` contain `NA`, `NaN` or infinite values, these points are deleted with a warning, instead of causing a fatal error.
- `predict.kppm`, `residuals.kppm`: Now issues a warning when the calculation ignores the cluster/Cox component and treats the model as if it were Poisson. (This currently happens in `predict.kppm` when `se=TRUE` or `interval != "none"`, and in `residuals.kppm` when `type != "raw"`).
- `predict.lppm`: Argument `locations` can now be an `lpp` object.
- `predict.mppm`: The argument `type="all"` is now recognised: it selects all the available options. [This is also the default.]
- `predict.ppm`:
 - Now recognises the arguments `dimyx` and `eps` for specifying the resolution of the grid of prediction points.
 - New argument `ignore.hardcore`.
- `predict.rhohat`: New argument `what` determines which value should be calculated: the function estimate, the upper/lower confidence limits, or the standard error.
- `print.linim`: More information is printed.
- `print.quad`: More information is printed.
- `progressreport`

- Behaviour improved.
 - New arguments `state`, `tick`, `showtime`.
 - New option: `style="tk"`
 - `pseudoR2.ppm`, `pseudoR2.lppm`:
 - The null model now includes any offset terms, by default.
 - New argument `keepoffset`.
 - `quadratcount.ppp`: Computation accelerated in some cases.
 - `quadrat.test.ppm`: Computation accelerated in some cases.
 - `quantile.ewcdf`: The function is now normalised to the range `[0,1]` before the quantiles are computed. This can be suppressed by setting `normalise=FALSE`.
 - `qqplot.ppm`: Argument `expr` can now be a list of point patterns, or an envelope object containing a list of point patterns.
 - `rcellnumber`: New argument `mu`.
 - `rgbim`, `hsvim`: New argument `A` controls the alpha (transparency) channel.
 - `rgb2hex`, `col2hex`, `paletteindex`, `is.colour`, `samecolour`, `complementarycolour`, `is.grey`, `to.grey`: These colour tools now handle transparent colours.
 - `rgb2hex`: New argument `maxColorValue`
 - `relrisk.ppp`:
 - If `se=TRUE` and `at="pixels"`, the result belongs to class `solist`.
 - The arguments `adjust`, `edge`, `diggle` are now explicit formal arguments.
- `rhohat`:
- New argument `subset` allows computation for a subset of the data.
 - New argument `positiveCI` specifies whether confidence limits should always be positive.
- `rhohat.lpp`: New argument `random` controls placement of dummy points.
- `rLGCP`: This function no longer requires the package `RandomFields` to be loaded explicitly.
 - `rMaternI`, `rMaternII`: These functions can now generate random patterns in three dimensions and higher dimensions, when the argument `win` is of class `box3` or `boxx`.
 - `rmh`: Accelerated, in the case where multiple patterns are saved using `nsave`.
 - `rmh.ppm`, `rmhmodel.ppm`, `simulate.ppm`: A model fitted using the `Penttinen` interaction can now be simulated.
 - `rmh.default`, `rmhmodel.default`:
 - These functions now recognise `cif='penttinen'` for the `Penttinen` interaction.
 - New arguments `nsim`, `saveinfo`.

- `rmhcontrol`: New parameter `pstage` determines when to generate random proposal points.
- `rose.default` New argument `weights`.
- `rose` New arguments `start` and `clockwise` specify the convention for measuring and plotting angles.
- `rotmean`:
 - New argument `padzero`.
 - Default behaviour has changed.
 - Improved algorithm stability.
- `rpoispp`: Accelerated, when `lambda` is a pixel image.
- `rpoisppx`: New argument `drop`.
- `rpoisline`: Also returns information about the original infinite random lines.
- `rStrauss`, `rHardcore`, `rStraussHard`, `rDiggleGratton`, `rDGS`, `rPenttinen`: New argument `drop`.
- `rtemper`: new argument `track`.
- `rthin`
 - Accelerated, when `P` is a single number.
 - `X` can now be a point pattern on a linear network (class `lpp`).
- `rThomas`, `rMatClust`, `rCauchy`, `rVarGamma`:
 - When the model is approximately Poisson, it is simulated using `rpoispp`. This avoids computations which would require huge amounts of memory. New argument `poisthresh` controls this behaviour.
 - New argument `saveparents`.
- `runifpointOnLines`, `rpoisppOnLines`: New argument `drop`.
- `runifpointx`: New argument `drop`.
- `setcov`: the name of the unit of length is preserved.
- Simulation: Several basic simulation algorithms have been accelerated. Consequently, simulation outcomes are not identical to those obtained with previous versions of `spatstat`, even when the same random seed is used. To ensure compatibility with previous versions of `spatstat`, revert to the slower code by setting `spatstat.options(fastthin=FALSE, fastpois=FALSE)`.
- `shapley`: In this installed dataset, the function `shapley.extra$plotit` has changed slightly (to accommodate the dependence on the package `spatstat.utils`).
- `simulate.ppm` New argument `w` controls the window of the simulated patterns. New argument `verbose`.
- `Smooth.ppp`:

- A non-Gaussian kernel can now be specified using the argument `kernel`.
 - Argument `weights` can now be a pixel image.
 - Infinite bandwidth `sigma=Inf` is supported.
 - Accelerated by about 30% in the case where `at="pixels"`.
 - Accelerated by about 15% in the case where `at="points"` and `kernel="gaussian"`.
 - Now exits gracefully if any mark values are `NA`, `NaN` or `Inf`.
 - New argument `geometric` supports geometric-mean smoothing.
 - The arguments `adjust`, `edge`, `diggle` are now explicit formal arguments.
- `solist`: New argument `.NameBase`
 - `spatstat.options` New options `fastthin` and `fastpois` enable fast simulation algorithms. Set these options to `FALSE` to reproduce results obtained with previous versions of `spatstat`.
 - `split.ppp`, `split.ppx`: The splitting variable `f` can now be a logical vector.
 - `square`: Handles a common error in the format of the arguments.
 - `step`: now works for models of class `"mppm"`.
 - `stieltjes`: Argument `M` can be a stepfun object (such as an empirical CDF).
 - `subset.ppp`, `subset.lpp`, `subset.pp3`, `subset.ppx`: The argument `subset` can now be any argument acceptable to the `"["` method.
 - summary functions The argument `correction="all"` is now recognised: it selects all the available options.

This applies to `Fest`, `F3est`, `Gest`, `Gcross`, `Gdot`, `Gmulti`, `G3est`, `Gfox`, `Gcom`, `Gres`, `Hest`, `Jest`, `Jmulti`, `Jcross`, `Jdot`, `Jfox`, `Kest`, `Kinhom`, `Kmulti`, `Kcross`, `Kdot`, `Kcom`, `Kres`, `Kmulti.inhom`, `Kcross.inhom`, `Kdot.inhom`, `Kscaled`, `Ksector`, `Kmark`, `K3est`, `Lscaled`, `markcorr`, `markcrosscorr`, `norient`, `pairorient`, `pcfinhom`, `pcfcross.inhom`, `pcfcross`, `pcf`, `Tstat`.
 - `Summary.linim` family supporting `range`, `max`, `min` etc: Recognises the argument `finite` so that `range(x, finite=TRUE)` works for a `linim` object `x`.
 - `summary.distfun`, `summary.funxy`:
 - More information is printed.
 - Pixel resolution can now be controlled.
 - `summary.kppm`: prints more information about algorithm convergence.
 - `summary.ppm`: New argument `fine` selects the algorithm for variance estimation.
 - `summary.owin`, `summary.im`: The fraction of frame area that is occupied by the window/image is now reported.
 - `sumouter`: New argument `y` allows computation of asymmetric outer products.
 - `symbolmap`:

- Now accepts a vector of colour values for the arguments `col`, `cols`, `fg`, `bg` if the argument `range` is given.
 - New option: `shape="arrows"`.
- `tess`: Argument `window` is ignored when `xgrid`, `ygrid` are given.
- `texturemap`: Argument `textures` can be missing or `NULL`.
- `textureplot`: Argument `x` can now be something acceptable to `as.im`.
- `to.grey` New argument `transparent`.
- `union.owin`: Improved behaviour when there are more than 2 windows.
- `update`: now works for models of class `"mppm"`.
- `update.kppm`:
 - New argument `evaluate`.
 - Now handles additional arguments in any order, with or without names.
 - Changed arguments.
 - Improved behaviour.
- `update.ppm`: For the case `update(model, X)` where `X` is a point pattern, if the window of `X` is different from the original window, then the model is re-fitted from scratch (i.e. `use.internal=FALSE`).
- `valid.ppm` This is now a method for the generic function `valid`.
- `vcov.mppm`: Now handles models with Gibbs interactions.
- `vcov.ppm`: Performance slightly improved, for Gibbs models.
- `[<-.im` Accepts an array for `value`.
- `[.im` The subset index `i` can now be a linear network. Then the result of `x[i, drop=FALSE]` is a pixel image of class `linim`.
- `[.layered`:
 - Subset index `i` can now be an `owin` object.
 - Additional arguments `...` are now passed to other methods.
- `[.leverage.ppm`: New argument `update`.
- `[.linnet`:
 - New argument `snip` determines what to do with segments of the network that cross the boundary of the window. Default behaviour has changed.
 - More robust against artefacts when the subset index is a pixel mask.
- `[.linim`: More robust against artefacts.
- `[.lpp`: New argument `snip` determines what to do with segments of the network that cross the boundary of the window. Default behaviour has changed.

- [.ppx: The subset index `i` may now be a spatial domain of class `boxx` or `box3`.
- [.ppp New argument `clip` determines whether the window is clipped.
- [.ppp The previously-unused argument `drop` now determines whether to remove unused levels of a factor.
- [.pp3, [.lpp, [.ppx, subset.ppp, subset.pp3, subset.lpp, subset.ppx: These methods now have an argument `drop` which determines whether to remove unused levels of a factor.
- [.psp:
 - accelerated.
 - New argument `fragments` specifies whether to keep fragments of line segments that are cut by the new window, or only to retain segments that lie entirely inside the window.
- [.solist: Subset index `i` can now be an `owin` object.

References

- [1] A. Baddeley. Analysing spatial point patterns in R. Technical report, CSIRO, 2010. Version 4. URL <https://research.csiro.au/software/r-workshop-notes/>
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