

# Package ‘ppdiag’

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**Type** Package

**Title** Diagnosis and Visualizations Tools for Temporal Point Processes

**Version** 0.1.1

**Description** A suite of diagnostic tools for univariate point processes.

This includes tools for simulating and fitting both common and more complex temporal point processes. We also include functions to visualise these point processes and collect existing diagnostic tools of Brown et al. (2002) <[doi:10.1162/08997660252741149](https://doi.org/10.1162/08997660252741149)> and Wu et al. (2021) <[doi:10.1002/9781119821588.ch7](https://doi.org/10.1002/9781119821588.ch7)>, which can be used to assess the fit of a chosen point process model.

**License** MIT + file LICENSE

**Encoding** UTF-8

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**URL** <https://owenward.github.io/ppdiag/>

**BugReports** <https://github.com/OwenWard/ppdiag/issues>

**Depends** R (>= 3.5)

**Imports** graphics, stats

**Suggests** testthat (>= 2.1.0), knitr, rmarkdown, covr

**VignetteBuilder** knitr

**NeedsCompilation** no

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drawHPIntensity	<i>Draw the intensity of Hawkes Process</i>
-----------------	---

---

**Description**

Draw the intensity of a Hawkes Process

**Usage**

```
drawHPIntensity(
  hp = NULL,
  events,
  int_title = "Hawkes Intensity",
  start = 0,
  end = max(events),
  history = NULL,
  color = 1,
  i = 1,
  add = FALSE,
  fit = FALSE,
  plot_events = TRUE,
  verbose = FALSE
)
```

**Arguments**

hp	object parameters for Hawkes process.
events	the event times happened in this state
int_title	title of the intensity plot
start	the start time of current state
end	the end time of current state
history	the past event times
color	specify the default plotting color.
i	state number, used only for drawUniMMHPIntensity
add	whether to add the hawkes intensity to an existing plot, used for drawUniMMHPIntensity
fit	a boolean indicating whether to fit a new HP to events
plot_events	indicate whether events will be plotted
verbose	whether to output informative messages as running

**Value**

no return value, intensity plot of Hawkes process

**Examples**

```
set.seed(100)
hp_obj <- pp_hp(lambda0 = 0.5, alpha = 0.45, beta = 0.5)
events <- pp_simulate(hp_obj, start = 0, end = 20)
drawHPPIntensity(hp_obj, events)
```

---

drawHPPIntensity	<i>Draw intensity of homogeneous Poisson process</i>
------------------	--

---

**Description**

Draw the intensity for a homogeneous Poisson process

**Usage**

```
drawHPPIntensity(
  hpp = NULL,
  events,
  int_title = "Homogeneous Poisson Process",
  start = 0,
  end = max(events),
  color = "red",
  plot_events = TRUE,
  fit = FALSE,
  add = FALSE,
  verbose = FALSE
)
```

**Arguments**

hpp	object for homogeneous Poisson process
events	event times input
int_title	the plot title
start	start of events
end	end of events
color	a specification for the default plotting color.
plot_events	a boolean indicating whether input events will be plotted
fit	a boolean indicating whether to fit a hpp or use the passed object
add	whether to add the hpp intensity to an existing plot
verbose	whether to output informative messages as running

**Value**

no return value, intensity plot of homogeneous Poisson process

**Examples**

```
pois_y <- pp_hpp(lambda = 1)
drawHPPIntensity(pois_y, events = pp_simulate(pois_y, end = 10))
```

---

drawUniMMHPIntensity *Draw the intensity of the Markov-modulated Hawkes Process(MMHP)*

---

**Description**

Take a mmhp object and draw its intensity accordingly

**Usage**

```
drawUniMMHPIntensity(
  mmhp,
  simulation,
  int_title = "Intensity of MMHP",
  leg_location = "topright",
  color = 1,
  add = FALSE
)
```

**Arguments**

mmhp	a mmhp object including its state, state_time, events, lambda0, lambda1, beta and alpha.
simulation	the simulated Markov-modulated Hawkes Process(MMHP)
int_title	title of the plot.
leg_location	location of legend, if moving needed
color	A specification for the default plotting color.
add	logical; if TRUE add to an already existing plot; if NA start a new plot taking the defaults for the limits and log-scaling of the x-axis from the previous plot. Taken as FALSE (with a warning if a different value is supplied) if no graphics device is open.

**Value**

no return value, intensity plot of Markov-modulated Hawkes process

**Examples**

```
Q <- matrix(c(-0.4, 0.4, 0.2, -0.2), ncol = 2, byrow = TRUE)
x <- pp_mmhp(Q,
  delta = c(1 / 3, 2 / 3), lambda0 = 0.9, lambda1 = 1.1,
  alpha = 0.8, beta = 1.2
)
y <- pp_simulate(x, n = 25)
drawUniMMHPIntensity(x, y)
```

---

drawUniMMPPIntensity *Draw the intensity of the Markov-modulated Poisson Process(MMPP)*

---

**Description**

Take a mmpp object and draw its intensity accordingly

**Usage**

```
drawUniMMPPIntensity(
  mmpp,
  simulation,
  add = FALSE,
  color = 1,
  fit = FALSE,
  int_title = "Intensity Plot of MMPP"
)
```

**Arguments**

<code>mmp</code>	a mmp object including its transition probability matrix, $\lambda_0$ , $\delta$ , and $c$ .
<code>simulation</code>	the simulated Markov-modulated Poisson Process(MMPP)
<code>add</code>	logical; if TRUE add to an already existing plot; if NA start a new plot taking the defaults for the limits and log-scaling of the x-axis from the previous plot. Taken as FALSE (with a warning if a different value is supplied) if no graphics device is open.
<code>color</code>	A specification for the default plotting color.
<code>fit</code>	a boolean indicating whether to fit the events provided
<code>int_title</code>	title of the plot.

**Value**

no return value, intensity plot of Markov-modulated Poisson process

**Examples**

```
Q <- matrix(c(-0.4, 0.4, 0.2, -0.2), ncol = 2, byrow = TRUE)
x <- pp_mmp(Q, delta = c(1 / 3, 2 / 3), lambda0 = 0.9, c = 1.2)
y <- pp_simulate(x, n = 10)
drawUniMMPPIIntensity(x, y)
```

---

fithp

*Determine the MLE of Hawkes process numerically*

---

**Description**

Determine the MLE of Hawkes process numerically

**Usage**

```
fithp(events, end = max(events), vec = c(0.1, 0.2, 0.3))
```

**Arguments**

<code>events</code>	event times
<code>end</code>	end of observation period starting from 0 (default last event)
<code>vec</code>	vector of initial parameter values

**Value**

a hp object indicating the maximum likelihood parameter values ( $\lambda_0, \alpha, \beta$ ) for Hawkes process. This is a non-convex problem and a (unique) solution is not guaranteed.

**Examples**

```
hp_obj <- pp_hp(lambda0 = 0.1, alpha = 0.45, beta = 0.5)
sims <- pp_simulate(hp_obj, start = 0, n = 10)
fithp(sims)
```

---

**fithpp***Fit a homogeneous poisson process to event data*

---

**Description**

Compute maximum likelihood estimator of the rate of a homogeneous Poisson process for the given events.

**Usage**

```
fithpp(events, end = max(events))
```

**Arguments**

**events**            vector containing the event times.  
**end**                end of observation period, starting from 0 (default is last event)

**Value**

a hpp object containing the events and the estimated parameter

**Examples**

```
pois_y <- pp_hpp(lambda = 1)
events <- pp_simulate(pois_y, end = 10)
fithpp(events)
```

---

**intensityqqplot***Draw intensity of fitted point process and QQ-Plot of rescaled events*

---

**Description**

Draw the intensity and q-q plot for models

**Usage**

```
intensityqqplot(object, events, markov_states)

## Default S3 method:
intensityqqplot(object, events, markov_states)

## S3 method for class 'hp'
intensityqqplot(object, events, markov_states = NULL)

## S3 method for class 'hpp'
intensityqqplot(object, events, markov_states = NULL)

## S3 method for class 'mmp'
intensityqqplot(object, events = markov_states$events, markov_states)

## S3 method for class 'mmhp'
intensityqqplot(object, events = markov_states$events, markov_states)
```

**Arguments**

object            parameters for the models: hp, hpp, and mmhp  
 events            event times  
 markov\_states    only for mmp and mmpp, markov states simulation output

**Value**

no return value, intensity and qq-plot in a single plot

---

pp\_compensator            *Compensators for point processes*

---

**Description**

Computes the compensator for included point processes

**Usage**

```
pp_compensator(object, events)

## Default S3 method:
pp_compensator(object, events)

## S3 method for class 'mmp'
pp_compensator(object, events)

## S3 method for class 'hp'
```



```

pp_compensator(object, events)

## S3 method for class 'mmhp'
pp_compensator(object, events)

## S3 method for class 'hpp'
pp_compensator(object, events)

```

### Arguments

object            a point process model  
events            event times, which can have first value as 0

### Value

compensator vector of rescaled interevent times

### Examples

```

hpp_obj <- pp_hpp(lambda = 1)
events <- pp_simulate(hpp_obj, end = 10)
comp <- pp_compensator(hpp_obj, events)

```

---

pp\_diag

*Summarise diagnostics for point process models*


---

### Description

Generate diagnostic tools for different point process models, including quantile-quantile plot, ks plot, raw residual and pearson residual.

### Usage

```

pp_diag(object, events)

## Default S3 method:
pp_diag(object, events)

## S3 method for class 'hp'
pp_diag(object, events)

## S3 method for class 'mmhp'
pp_diag(object, events)

## S3 method for class 'mmp'
pp_diag(object, events)

## S3 method for class 'hpp'
pp_diag(object, events)

```

**Arguments**

object	a point process model
events	event times

**Value**

Invisibly returns NULL. Outputs plots and summary of diagnostics to console

**Examples**

```
hpp_obj <- pp_hpp(lambda = 1)
events <- pp_simulate(hpp_obj, end = 50)
pp_diag(hpp_obj, events)
```

---

pp_hp	<i>Create a Hawkes process object</i>
-------	---------------------------------------

---

**Description**

Create a Hawkes Process with an exponential kernel according to the given parameters: lambda0, alpha, beta and events. If events are missing, then it means that data will be added later(simulated from this process)

**Usage**

```
pp_hp(lambda0, alpha, beta, events = NULL)
```

**Arguments**

lambda0	initial intensity at the start time
alpha	jump size in increase of intensity
beta	exponential decay of intensity
events	vector containing the event times. Note that the first event is at time zero. Alternatively, events could be specified as NULL, meaning that the data will be added later (e.g. simulated).

**Value**

hp object

**Examples**

```
pp_hp(lambda0 = 0.1, alpha = 0.45, beta = 0.5)
```

---

`pp_hpp`*Create a homogeneous Poisson process object*

---

**Description**

Create a homogeneous Poisson object according to given parameters: lambda, and events. If events are missing, then it means that data will be added later(simulated from this process).

**Usage**

```
pp_hpp(lambda, events = NULL)
```

**Arguments**

lambda	rate of the Poisson process
events	event times, optional

**Value**

hpp object

**Examples**

```
pp_hpp(lambda = 1)
```

---

`pp_kspplot`*KS plot of empirical and theoretical cdf curve of fitted point process*

---

**Description**

Plot empirical cdf plot for rescaled-inter-event-times and exponential cdf as a reference curve

**Usage**

```
pp_kspplot(r, ...)
```

**Arguments**

r	rescaled-inter-event-times
...	other arguments for plots

**Value**

no return value, KS plot for rescaled-inter-event-times and exponential cdf curve

pp\_mmhp

*Create a Markov-modulated Hawkes Process(MMHP) object***Description**

Create a Markov-modulated Hawkes Process(MMHP) model according to the given parameters: lambda0, lambda1, alpha, beta, event times and transition probability matrix. If event time events is missing, then it means that data will be added later(e.g. simulated)

**Usage**

```
pp_mmhp(lambda0, lambda1, alpha, beta, Q = NULL, delta = NULL, events = NULL)
```

**Arguments**

lambda0	intensity for homogeneous Poisson process.
lambda1	base intensity for Hawkes process.
alpha	jump size of the increase in intensity in the hawkes process
beta	exponential decrease of intensity in the hawkes process
Q	transition probability matrix.
delta	initial state probability.
events	vector containing the event times. Note that the first event is at time zero. Alternatively, events could be specified as NULL, meaning that the data will be added later (e.g. simulated).

**Value**

mmhp object

**Examples**

```
Q <- matrix(c(-0.4, 0.4, 0.2, -0.2), ncol = 2, byrow = TRUE)
pp_mmhp(Q,
  delta = c(1 / 3, 2 / 3), lambda0 = 0.9, lambda1 = 1.1,
  alpha = 0.8, beta = 1.2
)
```

---

pp\_mmpp *Create a Markov-modulated Poisson Process(MMPP) object*

---

### Description

Create a Markov-modulated Poisson Process(MMPP) model according to the given parameters: lambda0, c, q1, q2 and event times. If event time tau is missing, then it means that data will be added later(e.g. simulated)

### Usage

```
pp_mmpp(lambda0, c, Q, events = NULL, delta = NULL)
```

### Arguments

lambda0	parameters for Poisson process.
c	the proportion of intensity 1 over intensity 2
Q	transition probability matrix
events	vector containing the event times. Note that the first event is often specified as zero. Alternatively, events could be specified as NULL, meaning that the data will be added later (e.g. simulated).
delta	initial state probability.

### Value

mmpp object

### Examples

```
Q <- matrix(c(-0.4, 0.4, 0.2, -0.2), ncol = 2, byrow = TRUE)
pp_mmpp(Q = Q, lambda0 = 1, c = 1.5, delta = c(1 / 3, 2 / 3))
```

---

pp\_qqexp *Plot QQ-plot for rescaled-inter-event-times of fitted point process*

---

### Description

Generate Quantile-quantile plot for rescaled-inter-event-times, which are independently and identically distributed as exponential random variables with rate 1 under the true point process.

### Usage

```
pp_qqexp(r, ...)
```

**Arguments**

r                    rescaled-inter-event-times  
 ...                  other arguments for plots

**Value**

no return value, quantile-quantile plot for rescaled-inter-event-times

---

pp\_residual                  *Compute raw and pearson residuals for point process models*

---

**Description**

Compute raw and pearson residuals for point process models

**Usage**

```
pp_residual(object, events, start = 0, end = max(events), steps = 1000)
```

**Arguments**

object                point process model containing the parameters  
 events                vector of event times  
 start                 start of observation period (default 0)  
 end                    end of observation period (default final event)  
 steps                 number of steps for numeric integration (if needed)

**Value**

the raw and pearson residuals

**Examples**

```
Q <- matrix(c(-0.4, 0.4, 0.2, -0.2), ncol = 2, byrow = TRUE)
x <- pp_mmhp(Q,
  delta = c(1 / 3, 2 / 3), lambda0 = 0.9,
  lambda1 = 1.1, alpha = 0.8, beta = 1.2
)
y <- pp_simulate(x, n = 10)
pp_residual(x, events = y$events)
```

pp\_simulate

*Simulate events from a temporal point process***Description**

Currently available point processes are homogeneous Poisson, Hawkes with exponential kernel, MMHP and MMPP

**Usage**

```
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)

## Default S3 method:
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)

## S3 method for class 'hpp'
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)

## S3 method for class 'hp'
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)

## S3 method for class 'mmp'
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)

## S3 method for class 'mmhp'
pp_simulate(object, start = 0, end = 1, n = NULL, verbose = FALSE)
```

**Arguments**

object	point process model object of type hpp, hp, mmhp, or mmp
start	start time of events simulated. Not used for Markov modulated models
end	end time of events simulated. Not used for Markov modulated models
n	number of events simulated. Required for Markov modulated models, optional otherwise
verbose	whether to output informative messages as running

**Value**

a vector of event times for all models. For Markov modulated models, also returns details on the underlying latent process

**Examples**

```
hpp_obj <- pp_hpp(lambda = 1)
s <- pp_simulate(hpp_obj, n = 50)
```

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