Package 'distspec'

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`+.dist_spec` *Creates a delay distribution as the sum of two other delay distributions.*

Description

Creates a delay distribution as the sum of two other delay distributions.

Usage

```
## S3 method for class 'dist_spec'
e1 + e2
```

Arguments

e1	The first delay distribution (of type <dist_spec>) to combine.
e2	The second delay distribution (of type <dist_spec>) to combine.

Value

A delay distribution representing the sum of the two delays

Examples

```
# A fixed lognormal distribution with mean 5 and sd 1.
dist1 <- LogNormal(
  meanlog = 1.6, sdlog = 1, max = 20
)
dist1 + dist1

# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- Gamma(
  shape = Normal(3, 0.5),
  rate = Normal(2, 0.5),
  max = 20
)
dist1 + dist2
```

==.dist_spec *Compares two delay distributions*

Description

Compares two delay distributions

Usage

```
## S3 method for class 'dist_spec'
e1 == e2

## S3 method for class 'dist_spec'
e1 != e2
```

Arguments

e1	The first delay distribution (of type <dist_spec>) to combine.
e2	The second delay distribution (of type <dist_spec>) to combine.

Value

TRUE or FALSE

Examples

```
Fixed(1) == Normal(1, 0.5)
```

`Beta` *Beta distribution*

Description

A beta distribution as a `<dist_spec>`, given either by its shape parameters `shape1`/`shape2` or by its `mean`/`sd`. It is not discretised.

Usage

```
Beta(shape1, shape2, mean, sd, ...)
```

Arguments

`shape1`, `shape2`	Shape parameters of the beta distribution.
`mean`, `sd`	Mean and standard deviation of the distribution, as an alternative to `shape1`/`shape2`.
`...`	Limits of the distribution, passed to `bound_dist()`.

Value

A `<dist_spec>`.

See Also

Distributions for an overview and the other distributions.

Examples

```
Beta(shape1 = 2, shape2 = 5)
Beta(mean = 0.3, sd = 0.15)
```

bound_dist *Define bounds of a* <dist_spec>

Description

Set the bounds that constrain a distribution when it is discretised: `max` truncates the support at that value, while `cdf_cutoff` trims the tail by keeping the distribution only up to its `cdf_cutoff` quantile. Either bound drops the mass beyond it and renormalises the remaining PMF to sum to one.

Usage

```
bound_dist(x, max = Inf, cdf_cutoff = 1)
```

Arguments

`x`	A <dist_spec>.
`max`	Numeric, maximum value of the distribution. The distribution will be truncated at this value. Default: `Inf`, i.e. no maximum.
`cdf_cutoff`	Numeric in `(0, 1]`; the cumulative probability up to which the distribution is kept, i.e. it is truncated at the `cdf_cutoff` quantile. For example `cdf_cutoff = 0.999` keeps the distribution up to its 99.9th percentile. Default: `1`, i.e. keep the full distribution. A value below `0.5` is rejected, as it is almost certainly the tail probability to *drop* rather than the CDF level to keep (use `1 - x` instead).

Value

a <dist_spec> with relevant attributes set that define its bounds

See Also

`discretise()`, which applies these bounds when producing a PMF.

Examples

```
# Truncate a gamma distribution at 20
bound_dist(Gamma(mean = 5, sd = 1), max = 20)
# Keep it up to its 99.9th percentile
bound_dist(Gamma(mean = 5, sd = 1), cdf_cutoff = 0.999)
```

`c.dist_spec` *Combines multiple delay distributions for further processing*

Description

This combines the given distributions into a single composite <dist_spec> holding multiple delay distributions.

Note that distributions that already are combinations of other distributions cannot be combined with other combinations of distributions.

Usage

```
## S3 method for class 'dist_spec'
c(...)
```

Arguments

`...`	The delay distributions to combine

Value

Combined delay distributions (with class <dist_spec>)

Examples

```
# A fixed lognormal distribution with mean 5 and sd 1.
dist1 <- LogNormal(
  meanlog = 1.6, sdlog = 1, max = 20
)
dist1 + dist1

# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- Gamma(
  shape = Normal(3, 0.5),
  rate = Normal(2, 0.5),
  max = 20
)
c(dist1, dist2)
```

collapse *Collapse nonparametric distributions in a <dist_spec>*

Description

This convolves any consecutive nonparametric distributions contained in the <dist_spec>.

Usage

```
## S3 method for class 'dist_spec'
collapse(x, ...)
```

Arguments

x	A <dist_spec>
...	ignored

Value

A <dist_spec> where consecutive nonparametric distributions have been convolved

See Also

discretise() to produce the nonparametric components this convolves. The vignette("distspec") shows the full get_pmf(collapse(discretise(d1 + d2))) pipeline.

Examples

```
# A fixed gamma distribution with mean 5 and sd 1.
dist1 <- Gamma(mean = 5, sd = 1, max = 20)

# An uncertain lognormal distribution with meanlog and sdlog normally
# distributed as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- LogNormal(
  meanlog = Normal(3, 0.5),
  sdlog = Normal(2, 0.5),
  max = 20
)

# The sum of two distributions
collapse(discretise(dist1 + dist2, strict = FALSE))
```

convert_to_logmean *Convert mean and sd to log mean for a log normal distribution*

Description

Convert from mean and standard deviation to the log mean of the lognormal distribution. Useful for defining a `LogNormal()` distribution from a mean and standard deviation on the natural scale.

Usage

```
convert_to_logmean(mean, sd)
```

Arguments

mean	Numeric, mean of a distribution
sd	Numeric, standard deviation of a distribution

Value

The log mean of a lognormal distribution

Examples

```
convert_to_logmean(2, 1)
```

convert_to_logsd *Convert mean and sd to log standard deviation for a log normal distribution*

Description

Convert from mean and standard deviation to the log standard deviation of the lognormal distribution. Useful for defining a `LogNormal()` distribution from a mean and standard deviation on the natural scale.

Usage

```
convert_to_logsd(mean, sd)
```

Arguments

mean	Numeric, mean of a distribution
sd	Numeric, standard deviation of a distribution

Value

The log standard deviation of a lognormal distribution

Examples

```
convert_to_logsd(2, 1)
```

Dirichlet *Dirichlet prior for a nonparametric distribution*

Description

A Dirichlet prior over the weights of a nonparametric probability mass function, used to specify an uncertain NonParametric() distribution whose PMF is left uncertain given the Dirichlet prior. Give either alpha directly, or a reference prior PMF together with a concentration.

Usage

```
Dirichlet(alpha, prior, concentration, ...)
```

Arguments

alpha	A positive numeric vector of concentration parameters.
prior	Either a numeric PMF vector (zero-indexed, i.e. the first entry represents probability mass at zero) or a dist_spec object. If a dist_spec object is provided it will be discretised and the PMF extracted. If numeric, it will be normalised to sum to one internally.
concentration	A positive scalar controlling how tightly the Dirichlet prior concentrates around the supplied PMF. The Dirichlet alpha vector is computed as alpha_i = concentration * p_i where p_i is the prior PMF. Guidance on values: • concentration = 1: weak prior, each alpha equals the PMF value (near-uniform for roughly equal PMF entries) • concentration = 5-20: moderate flexibility around the reference shape • concentration = 50+: strong anchoring to the reference PMF
...	Not used.

Value

A <dist_spec>.

See Also

NonParametric() to use the prior, and Distributions for an overview.

Examples

```
Dirichlet(c(1, 1, 1, 1))
Dirichlet(prior = c(0.1, 0.3, 0.4, 0.2), concentration = 10)
```

`discretise` *Discretise a <dist_spec>*

Description

Discretise a <dist_spec>

Usage

```
## S3 method for class 'dist_spec'
discretise(x, strict = TRUE, remove_trailing_zeros = TRUE, ...)

discretize(x, ...)
```

Arguments

`x`	A `<dist_spec>`
`strict`	Logical; If `TRUE` (default) an error will be thrown if a distribution cannot be discretised (e.g., because no finite maximum has been specified or parameters are uncertain). If `FALSE` then any distribution that cannot be discretised will be returned as is.
`remove_trailing_zeros`	Logical; If `TRUE` (default), trailing zeroes in the resulting PMF will be removed. If `FALSE`, trailing zeroes will be retained.
`...`	ignored

Value

A `<dist_spec>` where all distributions with constant parameters are nonparametric. Extract the resulting PMF vector with `get_pmf()`.

Methodological details

The probability mass function is computed using the `{primarycensored}` package, which provides double censored PMF calculations. This correctly represents the probability mass function of a double censored distribution arising from the difference of two censored events.

The probability mass function of the discretised probability distribution is a vector where the first entry corresponds to the integral over the (0,1] interval of the corresponding continuous distribution (probability of integer 0), the second entry corresponds to the (0,2] interval (probability mass of integer 1), the third entry corresponds to the (1, 3] interval (probability mass of integer 2), etc.

The maximum value truncates the distribution: mass beyond it is dropped and the remaining PMF is renormalised to sum to one. A `cdf_cutoff` below `1` additionally trims the tail, keeping the distribution only up to its `cdf_cutoff` quantile.

Fixed distributions:

A `Fixed()` (point-mass) distribution is not discretised through a CDF but by its own method: an integer value places all of the mass on that integer, while a fractional value splits the mass proportionally across the two adjacent integers. For example `Fixed(2.25)` places 0.75 on 2 and 0.25 on 3.

References

Charniga, K., et al. "Best practices for estimating and reporting epidemiological delay distributions of infectious diseases using public health surveillance and healthcare data", *arXiv e-prints*, 2024. doi:10.48550/arXiv.2405.08841 Park, S. W., et al., "Estimating epidemiological delay distributions for infectious diseases", *medRxiv*, 2024. doi:10.1101/2024.01.12.24301247 Abbott S., et al., "primarycensored: Primary Event Censored Distributions", 2025. doi:10.5281/zenodo.13632839

See Also

`collapse()` to convolve the discretised components of a composite distribution into a single PMF, and `sample_dist()` to draw random samples. The `vignette("distspec")` shows the full `get_pmf(collapse(discretise + d2)))` pipeline.

Examples

```
# A fixed gamma distribution with mean 5 and sd 1, discretised to a PMF.
dist1 <- Gamma(mean = 5, sd = 1, max = 20)
get_pmf(discretise(dist1))

# An uncertain lognormal distribution cannot be discretised, so with
# `strict = FALSE` it is returned unchanged.
dist2 <- LogNormal(
  meanlog = Normal(3, 0.5),
  sdlog = Normal(2, 0.5),
  max = 20
)
discretise(dist2, strict = FALSE)

# A fractional fixed value splits its mass across the two adjacent integers.
get_pmf(discretise(Fixed(2.25)))
```

`Distributions` *Probability distributions*

Description

distspec represents probability distributions (typically epidemiological delays, such as generation times or reporting delays) as `<dist_spec>` objects. Each supported distribution has its own constructor: `LogNormal()`, `Gamma()`, `Normal()`, `Exponential()`, `Weibull()`, `Beta()`, `Fixed()`, `NonParametric()` and `Dirichlet()`.

Details

A parameter can be given either as a fixed numeric value or as an uncertain value (another `<dist_spec>`); currently only normally distributed uncertain parameters (from `Normal()`) are supported.

Each distribution has a "natural" (canonical) parameterisation, such as `shape` and `rate` for `Gamma()` or `meanlog` and `sdlog` for `LogNormal()`. It can sometimes also be specified using other parameters, such as its mean and standard deviation, which are then converted to the natural parameters (propagating any uncertainty with a first-order delta-method approximation).

See Also

`discretise()` and `collapse()` to discretise and convolve distributions, `sample_dist()` to draw samples, and `get_parameters()` / `get_pmf()` to inspect them.

Exponential *Exponential distribution*

Description

An exponential distribution as a `<dist_spec>`, given either by its `rate` or by its `mean`.

Usage

```
Exponential(rate, mean, ...)

Exp(...)
```

Arguments

`rate`	vector of rates.
`mean`	Mean of the distribution, as an alternative to `rate`.
`...`	Limits of the distribution, passed to `bound_dist()`.

Value

A `<dist_spec>`.

See Also

Distributions for an overview and the other distributions.

Examples

```
Exponential(rate = 1)
Exponential(mean = 4)
```

`Fixed` *Fixed (point-mass) distribution*

Description

A fixed (delta) distribution as a `<dist_spec>`, placing all of its mass on a single `value`.

Usage

```
Fixed(value, ...)
```

Arguments

`value`	Value of the fixed (delta) distribution.
`...`	Limits of the distribution, passed to `bound_dist()`.

Value

A `<dist_spec>`.

See Also

Distributions for an overview and the other distributions.

Examples

```
Fixed(value = 3)
Fixed(value = 3.5)
```

`fix_parameters` *Fix the parameters of a* `<dist_spec>`

Description

If the given `<dist_spec>` has any uncertainty, it is removed and the corresponding distribution converted into a fixed one.

Call this before `sample_dist()` or `get_pmf()` on an uncertain distribution, as neither can operate on a distribution that still carries a prior.

Usage

```
## S3 method for class 'dist_spec'
fix_parameters(x, strategy = c("mean", "sample"), ...)
```

Arguments

x	A <dist_spec>
strategy	Character; either "mean" (use the mean estimates of the mean and standard deviation) or "sample" (randomly sample mean and standard deviation from uncertainty given in the <dist_spec>
...	ignored

Value

A <dist_spec> object without uncertainty

Examples

```
# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist <- Gamma(
  shape = Normal(3, 0.5),
  rate = Normal(2, 0.5),
  max = 20
)

fix_parameters(dist)
```

Gamma *Gamma distribution*

Description

A gamma distribution as a <dist_spec>, given either by its natural parameters shape/rate (or shape/scale) or by its mean/sd.

Usage

```
Gamma(shape, rate, scale, mean, sd, ...)
```

Arguments

shape, scale	shape and scale parameters. Must be positive, scale strictly.
rate	an alternative way to specify the scale.
mean, sd	Mean and standard deviation of the distribution, as an alternative to shape/rate.
...	Limits of the distribution, passed to bound_dist().

Value

A <dist_spec>.

See Also

Distributions for an overview and the other distributions.

Examples

```
Gamma(mean = 4, sd = 1)
Gamma(shape = 16, rate = 4)
Gamma(shape = Normal(16, 2), rate = Normal(4, 1))
```

`get_distribution` *Get the distribution of a* `<dist_spec>`

Description

Get the distribution of a `<dist_spec>`

Usage

```
get_distribution(x, id = NULL)
```

Arguments

`x`	A `<dist_spec>`.
`id`	Integer; the id of the distribution to use (if x is a composite distribution). If `x` is a single distribution this is ignored and can be left at its default value of `NULL`.

Value

A character string naming the distribution (or "nonparametric")

Examples

```
dist <- Gamma(shape = 3, rate = 2, max = 10)
get_distribution(dist)
```

get_parameters *Get parameters of a parametric distribution*

Description

Generic function to extract the distribution parameters (e.g. shape and rate for Gamma) from a dist_spec object.

Usage

```
get_parameters(x, ...)

## S3 method for class 'dist_spec'
get_parameters(x, id = NULL, ...)
```

Arguments

x	A dist_spec object
...	Additional arguments passed to methods
id	Integer; the id of the distribution to use (if x is a composite distribution). If x is a single distribution this is ignored and can be left at its default value of NULL.

Value

A list of parameters of the distribution.

Examples

```
dist <- Gamma(shape = 3, rate = 2)
get_parameters(dist)
```

get_pmf *Get the probability mass function of a nonparametric distribution*

Description

Get the probability mass function of a nonparametric distribution

Usage

```
get_pmf(x, id = NULL)
```

Arguments

x	A <dist_spec>.
id	Integer; the id of the distribution to use (if x is a composite distribution). If x is a single distribution this is ignored and can be left at its default value of NULL.

Details

An uncertain (Dirichlet-backed) nonparametric distribution has no concrete PMF, so calling `get_pmf()` on one is an error. Resolve it to a fixed PMF first with `fix_parameters()` (e.g. `strategy = "mean"`).

Value

The pmf of the distribution

Examples

```
dist <- discretise(Gamma(shape = 3, rate = 2, max = 10))
get_pmf(dist)
```

---

is_constrained *Check if a <dist_spec> is constrained, i.e. has a finite maximum or nonzero CDF cutoff.*

Description

Check if a <dist_spec> is constrained, i.e. has a finite maximum or nonzero CDF cutoff.

Usage

```
## S3 method for class 'dist_spec'
is_constrained(x, ...)
```

Arguments

`x`	A `<dist_spec>`
`...`	ignored

Value

Logical; TRUE if `x` is constrained

Examples

```
# A fixed gamma distribution with mean 5 and sd 1.
dist1 <- Gamma(mean = 5, sd = 1, max = 20)

# An uncertain lognormal distribution with meanlog and sdlog normally
# distributed as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- LogNormal(
  meanlog = Normal(3, 0.5),
  sdlog = Normal(2, 0.5),
  max = 20
)
```

```
# both distributions are constrained and therefore so is the sum
is_constrained(dist1 + dist2)
```

`LogNormal` *Lognormal distribution*

Description

A lognormal distribution as a `<dist_spec>`, given either by its natural parameters `meanlog`/`sdlog` or by its `mean`/`sd`.

Usage

```
LogNormal(meanlog, sdlog, mean, sd, ...)
```

Arguments

`meanlog`, `sdlog`	mean and standard deviation of the distribution on the log scale with default values of `0` and `1` respectively.
`mean`, `sd`	Mean and standard deviation of the distribution, as an alternative to `meanlog`/`sdlog`.
`...`	Limits of the distribution, passed to `bound_dist()`.

Value

A `<dist_spec>`.

See Also

Distributions for an overview and the other distributions.

Examples

```
LogNormal(mean = 4, sd = 1)
LogNormal(mean = 4, sd = 1, max = 10)
# Uncertain parameters must be given as the natural parameters
LogNormal(meanlog = Normal(1.5, 0.5), sdlog = 0.25, max = 10)
```

`max.dist_spec` *Returns the maximum of one or more delay distribution*

Description

This works out the maximum of all the (parametric / nonparametric) delay distributions combined in the passed <dist_spec> (ignoring any uncertainty in parameters)

Usage

```
## S3 method for class 'dist_spec'
max(x, ...)
```

Arguments

`x`	The <dist_spec> to use
`...`	Not used

Value

A numeric vector of maxima, one per component.

Examples

```
# A fixed gamma distribution with mean 5 and sd 1.
dist1 <- Gamma(mean = 5, sd = 1, max = 20)
max(dist1)

# An uncertain lognormal distribution with meanlog and sdlog normally
# distributed as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- LogNormal(
  meanlog = Normal(3, 0.5),
  sdlog = Normal(2, 0.5),
  max = 20
)
max(dist2)

# The max the sum of two distributions
max(dist1 + dist2)
```

`mean.dist_spec` *Returns the mean of one or more delay distribution*

Description

This works out the mean of all the (parametric / nonparametric) delay distributions combined in the passed <dist_spec>.

Usage

```
## S3 method for class 'dist_spec'
mean(x, ..., ignore_uncertainty = FALSE)
```

Arguments

`x`	The `<dist_spec>` to use
`...`	Not used
`ignore_uncertainty`	Logical; whether to ignore any uncertainty in parameters. If set to FALSE (the default) then the mean of any uncertain parameters will be returned as NA.

Value

A numeric vector of means, one per component of the `<dist_spec>`; `NA` for any component with uncertain parameters unless `ignore_uncertainty = TRUE`.

Examples

```
# A fixed lognormal distribution with mean 5 and sd 1.
dist1 <- LogNormal(mean = 5, sd = 1, max = 20)
mean(dist1)

# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- Gamma(
  shape = Normal(3, 0.5),
  rate = Normal(2, 0.5),
  max = 20
)
mean(dist2)

# The mean of the sum of two distributions
mean(dist1 + dist2)
```

`new_dist_spec` *Internal function for generating a* `dist_spec` *given parameters and a distribution.*

Description

This will convert all parameters to natural parameters before generating a `dist_spec`. If they are uncertain the uncertainty is propagated to the natural parameters with a first-order (delta-method) approximation (see `convert_to_natural()`).

Usage

```
new_dist_spec(params, distribution, max = Inf, cdf_cutoff = 1)
```

Arguments

`params`	Parameters of the distribution (including `max`)
`distribution`	Character; the distribution type (e.g. `"gamma"`, `"lognormal"`, `"nonparametric"`).
`max`	Numeric, maximum value of the distribution. The distribution will be truncated at this value. Default: `Inf`, i.e. no maximum.
`cdf_cutoff`	Numeric in `(0, 1]`; the cumulative probability up to which the distribution is kept, i.e. it is truncated at the `cdf_cutoff` quantile. For example `cdf_cutoff = 0.999` keeps the distribution up to its 99.9th percentile. Default: `1`, i.e. keep the full distribution. A value below `0.5` is rejected, as it is almost certainly the tail probability to *drop* rather than the CDF level to keep (use `1 - x` instead).

Value

A `dist_spec` of the given specification.

Examples

```
new_dist_spec(
  params = list(mean = 2, sd = 1),
  distribution = "normal"
)
```

NonParametric *Nonparametric distribution*

Description

A nonparametric distribution as a <dist_spec>, defined directly by its probability mass function. The PMF can instead be left uncertain by passing a Dirichlet() prior.

Usage

```
NonParametric(pmf, ...)
```

Arguments

pmf	Probability mass function, as a zero-indexed numeric vector (the first entry is the mass at zero) or a <dist_spec> (e.g. from Dirichlet()). A numeric vector is normalised to sum to one.
...	Limits of the distribution, passed to bound_dist().

Value

A <dist_spec>.

See Also

Distributions for an overview and the other distributions.

Examples

```
NonParametric(c(0.1, 0.3, 0.2, 0.4))

# With a Dirichlet prior (PMF left uncertain)
NonParametric(pmf = Dirichlet(c(1, 1, 1, 1)))
```

Normal *Normal distribution*

Description

A normal distribution as a <dist_spec>, given by its mean and sd. Also used to give an uncertain parameter of another distribution.

Usage

```
Normal(mean, sd, ...)
```

Arguments

mean, sd	Mean and standard deviation of the distribution.
...	Limits of the distribution, passed to bound_dist().

Value

A <dist_spec>.

See Also

Distributions for an overview and the other distributions.

Examples

```
Normal(mean = 4, sd = 1)
Normal(mean = 4, sd = 1, max = 10)
```

---

plot.dist_spec *Plot PMF and CDF for a dist_spec object*

Description

This function takes a <dist_spec> object and plots its probability mass function (PMF) and cumulative distribution function (CDF) using {ggplot2}.

Usage

```
## S3 method for class 'dist_spec'
plot(x, samples = 50L, res = 1, cumulative = TRUE, ...)
```

Arguments

x	A <dist_spec> object
samples	Integer; Number of samples to generate for distributions with uncertain parameters (default: 50).
res	Numeric; Resolution of the PMF and CDF (default: 1, i.e. integer discretisation). This applies only to components discretised from a continuous distribution; a nonparametric component is already discretised on its integer support and is unaffected by res.
cumulative	Logical; whether to plot the cumulative distribution in addition to the probability mass function
...	ignored

Details

A component must have a finite range to be plotted. One with no finite max and no cdf_cutoff of its own raises an error; bound it first (e.g. with bound_dist()).

Value

A {ggplot2} object showing the PMF (and, if cumulative = TRUE, the CDF) of each component, faceted by distribution.

Examples

```
# A fixed lognormal distribution with mean 5 and sd 1.
dist1 <- LogNormal(mean = 1.6, sd = 0.5, max = 20)
# Plot discretised distribution with 1 day discretisation window
plot(dist1)
# Plot discretised distribution with 0.01 day discretisation window
plot(dist1, res = 0.01, cumulative = FALSE)

# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- Gamma(
  shape = Normal(3, 0.5),
  rate = Normal(2, 0.5),
  max = 20
)
plot(dist2)

# Multiple distributions with 0.1 discretisation window and do not plot the
# cumulative distribution
plot(dist1 + dist2, res = 0.1, cumulative = FALSE)
```

---

print.dist_spec *Prints the parameters of one or more delay distributions*

Description

This displays the parameters of the uncertain and probability mass functions of fixed delay distributions combined in the passed <dist_spec>.

Usage

```
## S3 method for class 'dist_spec'
print(x, ...)
```

Arguments

x	The <dist_spec> to use
...	Not used

Value

invisible

Examples

```
#' # A fixed lognormal distribution with mean 5 and sd 1.
dist1 <- LogNormal(mean = 1.5, sd = 0.5, max = 20)
print(dist1)

# An uncertain gamma distribution with shape and rate normally distributed
# as Normal(3, 0.5) and Normal(2, 0.5) respectively
dist2 <- Gamma(
  shape = Normal(3, 0.5), rate = Normal(2, 0.5), max = 20
)
print(dist2)
```

`sample_dist` *Sample from a distribution*

Description

Draws random samples from a <dist_spec> whose parameters are fixed numbers, using the base-R random-generation function for its family (e.g. `rgamma()` for a gamma distribution). A discretised distribution is sampled on its integer support.

Only distributions with fixed parameters can be sampled. If any parameter is itself a distribution (a prior), there is no single distribution to sample from and an error is raised.

A composite (multi-component) distribution is sampled per component, in keeping with `mean()`/`sd()`, which also return one value per component. Use `rowSums()` on the result to obtain samples of the combined (convolved) distribution.

Usage

```
sample_dist(x, n, ...)

## S3 method for class 'dist_spec'
sample_dist(x, n, ...)

## S3 method for class 'multi_dist_spec'
sample_dist(x, n, ...)
```

Arguments

x	A <dist_spec>.
n	The number of samples to draw.
...	Not used.

Value

For a single distribution, a numeric vector of `n` samples. For a composite distribution of `k` components, an `n` by `k` matrix, one column of `n` samples per component (`rowSums()` gives `n` samples of the combined distribution).

See Also

fix_parameters() to resolve an uncertain distribution to fixed parameters before sampling, and discretise() to obtain a PMF instead.

Examples

```
# Samples from a fixed gamma distribution
sample_dist(Gamma(shape = 2, rate = 1), 10)

# Samples from a discretised distribution, drawn on its integer support
sample_dist(discretise(Gamma(shape = 2, rate = 1, max = 20)), 10)

# A fixed distribution always returns the same value
sample_dist(Fixed(3), 5)

# A composite: an n-by-k matrix, one column per component
sample_dist(Gamma(shape = 2, rate = 1) + Gamma(shape = 3, rate = 1), 10)
```

Weibull *Weibull distribution*

Description

A Weibull distribution as a <dist_spec>, given either by its shape/scale or by its mean/sd.

Usage

```
Weibull(shape, scale, mean, sd, ...)
```

Arguments

shape, scale	shape and scale parameters, the latter defaulting to 1.
mean, sd	Mean and standard deviation of the distribution, as an alternative to shape/scale.
...	Limits of the distribution, passed to bound_dist().

Value

A <dist_spec>.

See Also

Distributions for an overview and the other distributions.

Examples

```
Weibull(shape = 1, scale = 1)
Weibull(shape = 1, scale = 1, max = 10)
Weibull(mean = 4, sd = 1)
```

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