

# Package: zoid (via r-universe)

November 26, 2024

**Title** Bayesian Zero-and-One Inflated Dirichlet Regression Modelling

**Version** 1.3.1

**Description** Fits Dirichlet regression and zero-and-one inflated Dirichlet regression with Bayesian methods implemented in Stan. These models are sometimes referred to as trinomial mixture models; covariates and overdispersion can optionally be included.

**License** GPL (>=3)

**Encoding** UTF-8

**LazyData** true

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.2.3

**Biarch** true

**URL** <https://noaa-nwfsc.github.io/zoid/>

**BugReports** <https://github.com/noaa-nwfsc/zoid/issues>

**Depends** R (>= 3.4.0)

**Imports** gtools, methods, Rcpp (>= 0.12.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.0), rstantools (>= 2.1.1)

**Suggests** testthat, knitr, rmarkdown

**LinkingTo** BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.0), StanHeaders (>= 2.26.0)

**SystemRequirements** GNU make

**VignetteBuilder** knitr

**Config/pak/sysreqs** make

**Repository** <https://nmfs-opensci.r-universe.dev>

**RemoteUrl** <https://github.com/noaa-nwfsc/zoid>

**RemoteRef** HEAD

**RemoteSha** 010fba15fa388f9de711d59c24c2e0cccf01c92b

## Contents

zoid-package . . . . .	2
broken_stick . . . . .	2
chinook . . . . .	3
coddiet . . . . .	4
fit_dirichlet . . . . .	4
fit_prior . . . . .	5
fit_zoid . . . . .	5
get_fitted . . . . .	7
get_pars . . . . .	8
parse_re_formula . . . . .	8
rmspe_calc . . . . .	9

<b>Index</b>	<b>10</b>
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zoid-package	<i>The 'zoid' package.</i>
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### Description

A DESCRIPTION OF THE PACKAGE

### References

Stan Development Team (2020). RStan: the R interface to Stan. R package version 2.21.2.  
<https://mc-stan.org>

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broken_stick	<i>Random generation of datasets using the dirichlet broken stick method</i>
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### Description

Random generation of datasets using the dirichlet broken stick method

### Usage

```
broken_stick(
  n_obs = 1000,
  n_groups = 10,
  ess_fraction = 1,
  tot_n = 100,
  p = NULL
)
```

**Arguments**

n_obs	Number of observations (rows of data matrix to simulate). Defaults to 10
n_groups	Number of categories for each observation (columns of data matrix). Defaults to 10
ess_fraction	The effective sample size fraction, defaults to 1
tot_n	The total sample size to simulate for each observation. This is approximate and the actual simulated sample size will be slightly smaller. Defaults to 100
p	The stock proportions to simulate from, as a vector. Optional, and when not included, random draws from the dirichlet are used

**Value**

A 2-element list, whose 1st element  $\chi_{\text{obs}}$  is the simulated dataset, and whose 2nd element is the underlying vector of proportions  $p$  used to generate the data

**Examples**

```
y <- broken_stick(n_obs = 3, n_groups = 5, tot_n = 100)

# add custom proportions
y <- broken_stick(
  n_obs = 3, n_groups = 5, tot_n = 100,
  p = c(0.1, 0.2, 0.3, 0.2, 0.2)
)
```

---

chinook

*Data from Satterthwaite, W.H., Ciancio, J., Crandall, E., Palmer-Zwahlen, M.L., Grover, A.M., O'Farrell, M.R., Anson, E.C., Mohr, M.S. & Garza, J.C. (2015). Stock composition and ocean spatial distribution from California recreational chinook salmon fisheries using genetic stock identification. Fisheries Research, 170, 166–178. The data genetic data collected from port-based sampling of recreationally-landed Chinook salmon in California from 1998-2002.*

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**Description**

Data from Satterthwaite, W.H., Ciancio, J., Crandall, E., Palmer-Zwahlen, M.L., Grover, A.M., O'Farrell, M.R., Anson, E.C., Mohr, M.S. & Garza, J.C. (2015). Stock composition and ocean spatial distribution from California recreational chinook salmon fisheries using genetic stock identification. Fisheries Research, 170, 166–178. The data genetic data collected from port-based sampling of recreationally-landed Chinook salmon in California from 1998-2002.

**Usage**

```
chinook
```

**Format**

A data frame.

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coddiet	<i>Data from Magnussen, E. 2011. Food and feeding habits of cod (Gadus morhua) on the Faroe Bank. – ICES Journal of Marine Science, 68: 1909–1917. The data here are Table 3 from the paper, with sample proportions (columns w) multiplied by total weight to yield total grams (g) for each sample-diet item combination. Dashes have been replaced with 0s.</i>
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**Description**

Data from Magnussen, E. 2011. Food and feeding habits of cod (*Gadus morhua*) on the Faroe Bank. – *ICES Journal of Marine Science*, 68: 1909–1917. The data here are Table 3 from the paper, with sample proportions (columns w) multiplied by total weight to yield total grams (g) for each sample-diet item combination. Dashes have been replaced with 0s.

**Usage**

coddiet

**Format**

A data frame.

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fit_dirichlet	<i>Extract point estimates of compositions from fitted model.</i>
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**Description**

Extract point estimates of compositions from fitted model.

**Usage**

```
fit_dirichlet(data)
```

**Arguments**

data	The data to fit the dirichlet distribution to
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fit_prior	<i>Find appropriate standard deviations for prior</i>
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**Description**

Find appropriate standard deviations for prior

**Usage**

```
fit_prior(n_bins, n_draws = 10000, target = 1/n_bins, iterations = 5)
```

**Arguments**

n_bins	Bins for the Dirichlet distribution
n_draws	Numbers of samples to use for doing calculation
target	The goal of the specified prior, e.g. 1 or 1/n_bins
iterations	to try, to ensure robust solution. Defaults to 5

**Value**

A 3-element list consisting of sd (the approximate standard deviation in transformed space that gives a similar prior to that specified), value (the value of the root mean squared percent error function being minimized), and convergence (0 if convergence occurred, error code from `optim()` otherwise)

**Examples**

```
# fit model with 3 components / alpha = 1
set.seed(123)
f <- fit_prior(n_bins = 3, n_draws = 1000, target = 1)
# fit model with 20 components / alpha = 1/20
f <- fit_prior(n_bins = 20, n_draws = 1000, target = 1 / 20)
```

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fit_zoid	<i>Fit a trinomial mixture model with Stan</i>
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**Description**

Fit a trinomial mixture model that optionally includes covariates to estimate effects of factor or continuous variables on proportions.

**Usage**

```
fit_zoid(
  formula = NULL,
  design_matrix,
  data_matrix,
  chains = 3,
  iter = 2000,
  warmup = floor(iter/2),
  overdispersion = FALSE,
  overdispersion_sd = 5,
  posterior_predict = FALSE,
  moment_match = FALSE,
  prior_sd = NA,
  ...
)
```

**Arguments**

formula	The model formula for the design matrix. Does not need to have a response specified. If =NULL, then the design matrix is ignored and all rows are treated as replicates
design_matrix	A data frame, dimensioned as number of observations, and covariates in columns
data_matrix	A matrix, with observations on rows and number of groups across columns
chains	Number of mcmc chains, defaults to 3
iter	Number of mcmc iterations, defaults to 2000
warmup	Number iterations for mcmc warmup, defaults to 1/2 of the iterations
overdispersion	Whether or not to include overdispersion parameter, defaults to FALSE
overdispersion_sd	Prior standard deviation on 1/overdispersion parameter, Defaults to inv-Cauchy(0,5)
posterior_predict	Whether or not to return draws from posterior predictive distribution (requires more memory)
moment_match	Whether to do moment matching via <code>loo::loo_moment_match()</code> . This increases memory by adding all temporary parameters to be saved and returned
prior_sd	Parameter to be passed in to use as standard deviation of the normal distribution in transformed space. If covariates are included this defaults to 1, but for models with single replicate, defaults to 1/n_bins.
...	Any other arguments to pass to <code>rstan::sampling()</code> .

**Examples**

```
y <- matrix(c(3.77, 6.63, 2.60, 0.9, 1.44, 0.66, 2.10, 3.57, 1.33),
  nrow = 3, byrow = TRUE
)
# fit a model with no covariates
fit <- fit_zoid(data_matrix = y, chains = 1, iter = 100)
```

```

# fit a model with 1 factor
design <- data.frame("fac" = c("spring", "spring", "fall"))
fit <- fit_zoid(formula = ~fac, design_matrix = design, data_matrix = y, chains = 1, iter = 100)

# try a model with random effects
set.seed(123)
y <- matrix(runif(99,1,4), ncol=3)
design <- data.frame("fac" = sample(letters[1:5], size=nrow(y), replace=TRUE))
design$fac <- as.factor(design$fac)
fit <- fit_zoid(formula = ~(1|fac), design_matrix = design, data_matrix = y, chains = 1, iter = 100)

```

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get\_fitted

*Extract estimates of predicted latent proportions.*


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### Description

Extract point estimates of compositions from fitted model.

### Usage

```
get_fitted(fitted_model, conf_int = 0.05)
```

### Arguments

fitted_model	The fitted model returned as an rstan object from the call to zoid
conf_int	Parameter controlling confidence intervals calculated, defaults to 0.05 for 95% intervals

### Value

A list containing the posterior summaries of estimated parameters, with element mu (the predicted values in normal space). For predictions in transformed space, or overdispersion, see [get\\_pars\(\)](#)

### Examples

```

y <- matrix(c(3.77, 6.63, 2.60, 0.9, 1.44, 0.66, 2.10, 3.57, 1.33),
  nrow = 3, byrow = TRUE
)
# fit a model with no covariates
fit <- fit_zoid(data_matrix = y)
p_hat <- get_fitted(fit)

```

---

get_pars	<i>Extract parameters from fitted model.</i>
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---

### Description

Extract estimated parameters from fitted model.

### Usage

```
get_pars(fitted_model, conf_int = 0.05)
```

### Arguments

fitted_model	The fitted model returned as an rstan object from the call to zoid
conf_int	Parameter controlling confidence intervals calculated, defaults to 0.05 for 95% intervals

### Value

A list containing the posterior summaries of estimated parameters. At minimum, this will include `p` (the estimated proportions) and `betas` (the predicted values in transformed space). For models with overdispersion, an extra element `phi` will also be returned, summarizing overdispersion. For models with random intercepts, estimates of the group level effects will also be returned as `zetas` (again, in transformed space). For predictions in normal space, see [get\\_fitted\(\)](#)

### Examples

```
y <- matrix(c(3.77, 6.63, 2.60, 0.9, 1.44, 0.66, 2.10, 3.57, 1.33),
  nrow = 3, byrow = TRUE
)
# fit a model with no covariates
fit <- fit_zoid(data_matrix = y)
p_hat <- get_pars(fit)
```

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parse_re_formula	<i>Fit a trinomial mixture model that optionally includes covariates to estimate effects of factor or continuous variables on proportions.</i>
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### Description

Fit a trinomial mixture model that optionally includes covariates to estimate effects of factor or continuous variables on proportions.



**Usage**

```
parse_re_formula(formula, data)
```

**Arguments**

formula	The model formula for the design matrix.
data	The data matrix used to construct RE design matrix

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rmspe_calc	<i>Find appropriate prior for a given target distribution.</i>
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---

**Description**

Extract point estimates of compositions from fitted model.

**Usage**

```
rmspe_calc(par, n_bins, n_draws, target)
```

**Arguments**

par	The parameter (standard deviation) to be searched over to find a Dirichlet equivalent
n_bins	Bins for the Dirichlet distribution
n_draws	Numbers of samples to use for doing calculation
target	The goal of the specified prior, e.g. 1 or 1/n_bins

# Index

## \* datasets

chinook, 3

coddiet, 4

broken\_stick, 2

chinook, 3

coddiet, 4

fit\_dirichlet, 4

fit\_prior, 5

fit\_zoid, 5

get\_fitted, 7

get\_fitted(), 8

get\_pars, 8

get\_pars(), 7

loo::loo\_moment\_match(), 6

optim(), 5

parse\_re\_formula, 8

rmspe\_calc, 9

rstan::sampling(), 6

trinomix (zoid-package), 2

zoid-package, 2