

# Package ‘scDECO’

April 3, 2025

**Type** Package

**Title** Estimating Dynamic Correlation

**Version** 0.1.1

**Description** Implementations for two different Bayesian models of differential co-expression. `scdeco.cop()` fits the bivariate Gaussian copula model from Zichen Ma, Shannon W. Davis, Yen-Yi Ho (2023) <[doi:10.1111/biom.13701](https://doi.org/10.1111/biom.13701)>, while `scdeco.pg()` fits the bivariate Poisson-Gamma model from Zhen Yang, Yen-Yi Ho (2022) <[doi:10.1111/biom.13457](https://doi.org/10.1111/biom.13457)>.

**Imports** MASS, rjags, msm

**License** GPL (>= 3)

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**URL** <https://github.com/YenYiHo-Lab/scDECO>

**BugReports** <https://github.com/YenYiHo-Lab/scDECO/issues>

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**NeedsCompilation** no

**Author** Anderson Bussing [aut, cre],  
Yen-Yi Ho [aut, ths],  
Zichen Ma [aut],  
Zhen Yang [aut]

**Maintainer** Anderson Bussing <[abussing@email.sc.edu](mailto:abussing@email.sc.edu)>

**Repository** CRAN

**Date/Publication** 2025-04-02 22:20:02 UTC

## Contents

|                          |          |
|--------------------------|----------|
| scdeco.cop . . . . .     | 2        |
| scdeco.pg . . . . .      | 3        |
| scdeco.sim.cop . . . . . | 5        |
| scdeco.sim.pg . . . . .  | 6        |
| <b>Index</b>             | <b>8</b> |

---

`scdeco.cop`*Copula dynamic correlation fitting function*

---

**Description**

Copula dynamic correlation fitting function

**Usage**

```
scdeco.cop(  
  y,  
  x,  
  marginals,  
  w = NULL,  
  n.mcmc = 10000,  
  burn = 1000,  
  thin = 1,  
  offset1 = NULL,  
  offset2 = NULL  
)
```

**Arguments**

|                        |   |
|------------------------|---|
| <code>y</code>         | 2-column matrix of observations                   |
| <code>x</code>         | covariates  |
| <code>marginals</code> | length-2 vector with strings of the two marginals |
| <code>w</code>         | (optional)  |
| <code>n.mcmc</code>    | number of mcmc iterations to run                  |
| <code>burn</code>      | how many of the mcmc iterations to burn           |
| <code>thin</code>      | how much to thin the mcmc iterations              |
| <code>offset1</code>   | (optional) offset for link(mu1)                   |
| <code>offset2</code>   | (optional) offset for link(mu2)                   |

**Value**

matrix with mcmc samples as rows and columns corresponding to the different parameters

**Examples**

```
n <- 1000  
x.use = rnorm(n)  
w.use = runif(n,-1,1)  
eta1.use = c(-2.2, 0.7)  
eta2.use = c(-2, 0.8)  
beta1.use = c(1,0.5)
```

```

beta2.use = c(1,1)
alpha1.use = 7
alpha2.use = 3
tau.use = c(-0.2, .3)

marginals.use <- c("ZINB", "ZIGA")

y.use <- scdeco.sim.cop(marginals=marginals.use, x=x.use,
                      eta1.true=eta1.use, eta2.true=eta2.use,
                      beta1.true=beta1.use, beta2.true=beta2.use,
                      alpha1.true=alpha1.use, alpha2.true=alpha2.use,
                      tau.true=tau.use, w=w.use)
mcmc.out <- scdeco.cop(y=y.use, x=x.use, marginals=marginals.use, w=w.use,
                     n.mcmc=10, burn=0, thin=1) # n.mcmc=1000, burn=100, thin=5)

lowerupper <- t(apply(mcmc.out, 2, quantile, c(0.025, 0.5, 0.975)))
estmat <- cbind(lowerupper[,1],
                c(eta1.use, eta2.use, beta1.use, beta2.use, alpha1.use, alpha2.use, tau.use),
                lowerupper[,c(2,3)])
colnames(estmat) <- c("lower", "trueval", "estval", "upper")
estmat

```

---

scdeco.pg

*ZENCO Poisson Gamma dynamic correlation fitting function*


---

## Description

ZENCO Poisson Gamma dynamic correlation fitting function

## Usage

```

scdeco.pg(
  dat,
  b0,
  b1,
  adapt_iter = 100,
  update_iter = 100,
  coda_iter = 1000,
  coda_thin = 5,
  coda_burnin = 100
)

```

## Arguments

|     |  |
|-----|--|
| dat | matrix containing expression values as first two columns and covariate as third column |
| b0  | intercept of zinf parameter  |

|             |   |
|-------------|---|
| b1          | slope of zinf parameter   |
| adapt_iter  | number of adaptation iterations in the jags.model function          |
| update_iter | update iterations in the update function                            |
| coda_iter   | number of iterations for the coda.sample function                   |
| coda_thin   | how much to thin the resulting MCMC output                          |
| coda_burnin | how many iterations to burn before beginning coda sample collection |

### Value

MCMC samples that have been adapted, burned, and thinned

### Examples

```

phi1_use <- 4
phi2_use <- 4
phi3_use <- 1/7
mu1_use <- 15
mu2_use <- 15
mu3_use <- 7
b0_use <- -3
b1_use <- 0.1
tau0_use <- -2
tau1_use <- 0.4

simdat <- scdeco.sim.pg(N=1000, b0=b0_use, b1=b1_use,
                      phi1=phi1_use, phi2=phi2_use, phi3=phi3_use,
                      mu1=mu1_use, mu2=mu2_use, mu3=mu3_use,
                      tau0=tau0_use, tau1=tau1_use)

zenco_out <- scdeco.pg(dat=simdat,
                    b0=b0_use, b1=b1_use,
                    adapt_iter=1, # 500,
                    update_iter=1, # 500,
                    coda_iter=5, # 5000,
                    coda_thin=1, # 10,
                    coda_burnin=0) # 1000

boundsmat <- cbind(zenco_out$quantiles[,1],
                  c(1/phi1_use, 1/phi2_use, 1/phi3_use,
                    mu1_use, mu2_use, mu3_use,
                    tau0_use, tau1_use),
                  zenco_out$quantiles[,c(3,5)])

colnames(boundsmat) <- c("lower", "true", "est", "upper")

boundsmat

```

---

scdeco.sim.cop      *Simulating from copula model*

---

**Description**

Simulating from copula model

**Usage**

```
scdeco.sim.cop(  
  marginals,  
  x,  
  eta1.true,  
  eta2.true,  
  beta1.true,  
  beta2.true,  
  alpha1.true,  
  alpha2.true,  
  tau.true,  
  w = NULL  
)
```

**Arguments**

|             |  |
|-------------|--|
| marginals   | provide vector of length 2 of which marginals to use   |
| x           | covariate matrix                                       |
| eta1.true   | zero-inflation parameters for marginal 1               |
| eta2.true   | zero-inflation parameters for marginal 2               |
| beta1.true  | mean coefficients for marginal 1                       |
| beta2.true  | mean coefficients for marginal 2                       |
| alpha1.true | second parameter coefficients for marginal 1           |
| alpha2.true | second parameter coefficients for marginal 2           |
| tau.true    | coefficients for correlation                           |
| w           | (optional) covariate matrix for zero-inflation portion |

**Value**

matrix with values simulated from copula model

**Examples**

```

n <- 2500
x.use = rnorm(n)
w.use = runif(n,-1,1)
eta1.use = c(-2.2, 0.7)
eta2.use = c(-2, 0.8)
beta1.use = c(1,0.5)
beta2.use = c(1,1)
alpha1.use = 7
alpha2.use = 3
tau.use = c(-0.2, .3)

marginals.use <- c("ZINB", "ZIGA")

y.use <- scdeco.sim.cop(marginals=marginals.use, x=x.use,
                      eta1.true=eta1.use, eta2.true=eta2.use,
                      beta1.true=beta1.use, beta2.true=beta2.use,
                      alpha1.true=alpha1.use, alpha2.true=alpha2.use,
                      tau.true=tau.use, w=w.use)

y.use[1:10,]

```

---

scdeco.sim.pg

*Simulating from ZENCO Model*


---

**Description**

Simulating from ZENCO Model

**Usage**

```

scdeco.sim.pg(
  N,
  b0,
  b1,
  phi1,
  phi2,
  mu1,
  mu2,
  tau0,
  tau1,
  mu3,
  phi3,
  tau2 = NULL,
  tau3 = NULL,
  xc = NULL
)

```

**Arguments**

|      |   |
|------|---|
| N    | size of sample to be generated                                      |
| b0   | intercept of zinf parameter   |
| b1   | slope of zinf parameter   |
| phi1 | over-dispersion parameter of first marginal                         |
| phi2 | over-dispersion parameter of second marginal                        |
| mu1  | mean parameter of first marginal                                    |
| mu2  | mean parameter of second marginal                                   |
| tau0 | intercept of correlation  |
| tau1 | slope of of correlation   |
| mu3  | mean parameter of covariate vector                                  |
| phi3 | over-dispersion parameter of covariate vector                       |
| tau2 | (optional) correlation coefficient on optional xc covariate vector  |
| tau3 | (optional) correlation coefficient on interaction between x3 and xc |
| xc   | (optional) secondary covariate to be regressed                      |

**Value**

a matrix with expressions as first two columns and covariates as remaining columns

**Examples**

```
phi1_use <- 4
phi2_use <- 4
phi3_use <- 1/6
mu1_use <- 15
mu2_use <- 15
mu3_use <- 7
b0_use <- 0.6882
b1_use <- -0.2995
tau0_use <- 0.07
tau1_use <- 0.05

simdat <- scdeco.sim.pg(N=1000, b0=b0_use, b1=b1_use,
                       phi1=phi1_use, phi2=phi2_use, phi3=phi3_use,
                       mu1=mu1_use, mu2=mu2_use, mu3=mu3_use,
                       tau0=tau0_use, tau1=tau1_use)

simdat[1:10,]
```

# Index

scdeco.cop, [2](#)  
scdeco.pg, [3](#)  
scdeco.sim.cop, [5](#)  
scdeco.sim.pg, [6](#)