

# Package: linconGaussR (via r-universe)

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**Type** Package

**Title** Sampling Multivariate Normal Distribution under Linear Constraints

**Version** 0.1

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**Description** Sample truncated multivariate Normal distribution following Gessner, A., Kanjilal, O., & Hennig, P. (2019). Integrals over Gaussians under Linear Domain Constraints. 108. <[arxiv:1910.09328](https://arxiv.org/abs/1910.09328)>.

**License** GPL-3

**Imports** Rcpp (>= 1.0.7), MASS

**LinkingTo** Rcpp, RcppArmadillo

**URL** <https://github.com/YunyiShen/linconGaussR>

**BugReports** <https://github.com/YunyiShen/linconGaussR/issues>

**RoxygenNote** 7.1.1

**NeedsCompilation** yes

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**Repository** CRAN

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linconGauss	<i>Sample Gaussian distribution with linear constraints Taking truncated sample of Gaussian distribution over a linear constraint domain.</i>
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### Description

Sample Gaussian distribution with linear constraints Taking truncated sample of Gaussian distribution over a linear constraint domain.

### Usage

```
linconGauss(
  n,
  A,
  b,
  Sigma,
  mu,
  x_init = NULL,
  intersection = TRUE,
  n_retry_init = 1000,
  nskip = 5
)
```

### Arguments

n	number of samples to take
A	a matrix with M by D dimensions, the linear constraints, such that $Ax+b \geq 0$
b	the offset of the linear constraints with dimension M such that $Ax+b \geq 0$
Sigma	covariance matrix of the Gaussian
mu	mean vector of the Gaussian
x_init	the sample to start with, if NULL, a sample will be drawn using rejection method
intersection	bool whether sample from the intersection or the union of the linear constraints, default true, sample from the intersection
n_retry_init	how many times to try finding a initial value
nskp	how many sample to skip during the sampling routine

### Value

a matrix with truncated sample, row as samples

**Examples**

```
my_sample <- linconGauss(100, diag(2),c(0,0),diag(2),c(0,0))
MASS_sample <- MASS::mvrnorm(1000,c(0,0),diag(2))
plot(MASS_sample)
points(my_sample,col = "red")
abline(h=0)
abline(v=0)
```

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