



geneticLHS {lhs}

R Documentation

## Latin Hypercube Sampling with a Genetic Algorithm

### Description

Draws a Latin Hypercube Sample from a set of uniform distributions for use in creating a Latin Hypercube Design. This function attempts to optimize the sample with respect to the S optimality criterion through a genetic type algorithm.

### Usage

```
geneticLHS(n=10, k=2, pop=100, gen=4, pMut=.1, verbose=FALSE)
```

### Arguments

<code>n</code>	The number of partitions (simulations or design points)
<code>k</code>	The number of replications (variables)
<code>pop</code>	The number of designs in the initial population
<code>gen</code>	The number of generations over which the algorithm is applied
<code>pMut</code>	The probability with which a mutation occurs in a column of the progeny

verbose Print informational messages

## Details

Latin hypercube sampling (LHS) was developed to generate a distribution of collections of parameter values from a multidimensional distribution. A square grid containing possible sample points is a Latin square iff there is only one sample in each row and each column. A Latin hypercube is the generalisation of this concept to an arbitrary number of dimensions. When sampling a function of  $k$  variables, the range of each variable is divided into  $n$  equally probable intervals.  $n$  sample points are then drawn such that a Latin Hypercube is created. Latin Hypercube sampling generates more efficient estimates of desired parameters than simple Monte Carlo sampling.

This program generates a Latin Hypercube Sample by creating random permutations of the first  $n$  integers in each of  $k$  columns and then transforming those integers into  $n$  sections of a standard uniform distribution. Random values are then sampled from within each of the  $n$  sections. Once the sample is generated, the uniform sample from a column can be transformed to any distribution by using the quantile functions, e.g. `qnorm()`. Different columns can have different distributions.

S-optimality seeks to maximize the mean distance from each design point to all the other points in the design, so the points are as spread out as possible.

Genetic Algorithm:

1. Generate `pop` random latin hypercube designs of size  $n$  by  $k$
2. Calculate the S optimality measure of each design
3. Keep the best design in the first position and throw away half of the rest of the population
4. Take a random column out of the best matrix and place it in a random column of each of the other matrices, and take a random column out of each of the other matrices and put it in copies of the best matrix thereby causing the progeny

5. For each of the progeny, cause a genetic mutation  $p_{Mut}$  percent of the time. The mutation is accomplished by switching two elements in a column

## Value

An  $n$  by  $k$  Latin Hypercube Sample matrix with values uniformly distributed on  $[0,1]$

## Author (s)

Rob Carnell

## References

Stocki, R. (2005) A method to improve design reliability using optimal Latin hypercube sampling *Computer Assisted Mechanics and Engineering Sciences* **12**, 87–105.

Stein, M. (1987) Large Sample Properties of Simulations Using Latin Hypercube Sampling. *Technometrics*. **29**, 143–151.

## See Also

[randomLHS](#), [improvedLHS](#), [maximinLHS](#), and [optimumLHS](#) to generate Latin Hypercube Samples. [optAugmentLHS](#), [optSeededLHS](#), and [augmentLHS](#) to modify and augment existing designs.

## Examples

```
geneticLHS(4, 3, 50, 5, .25)
```