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A Vignette

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CRPclustering: An R Package for Bayesian Nonparametric Chinese Restaurant Process Clustering with Entropy

Abstract

Clustering is a scientific method which finds the clusters of data and many related methods are traditionally researched for a long time. Bayesian nonparametrics is statistics which can treat models having infinite parameters. Chinese restaurant process is used in order to compose Dirichlet process. The clustering which uses Chinese restaurant process does not need to decide the number of clusters in advance. This algorithm automatically adjusts it. Then, this package can calculate clusters in addition to entropy as the ambiguity of clusters.

Introduction

Clustering is a traditional method in order to find the clusters of data and many related methods are studied for several decades. The most popular method is called as K-means[2]. K-means is an algorithmic way in order to search the clusters of data. However its method needs to decide the number of clusters in advance. Therefore if the data is both high dimensions and a complex, deciding the accurate number of clusters is difficult and normal Bayesian methods are too. For that reason, Bayesian nonparametric methods are gradually important as computers are faster than ever. In this package, we implemented Chinese restaurant process clustering (CRP)[4]. CRP can compose infinite dimensional parameters as Dirichlet process[1]. It acts like customers who sit at tables in a restaurant and has a probability to sit at a new table. As a result, Its model always automates clustering. Moreover, we added the method which calculates the entropy[5] of clusters into this package. It can check the ambiguity of the result. Then, we explain the clustering model and how to use it in detail. Finally, an example is plotted on a graph.

Background

Chinese Restaurant Process

Chinese restaurant process is a metaphor looks like customers sit at a table in Chinese restaurant. All customers except for x_i have already sat at finite tables. A new customer x_i will sit at either a table which other customers have already sat at or a new table. A new customer tends to sit at a table which has the number of customers more than other tables. A probability equation is given by

$$p(z_i = k | x_{1:n}, z_{1:n}^{\setminus i}, \alpha, \mu_0, \rho_0, a_0, b_0) = \begin{cases} p(x_i | \mu_k, \tau) \times \frac{n_k^{\setminus i}}{n-1+\alpha} & \text{if } k \in K^+(Z_{1:n}^{\setminus i}), \\ p(x_i | \mu_k, \tau) \times \frac{\alpha}{n-1+\alpha} & \text{if } k = |K^+(Z_{1:n}^{\setminus i})| + 1. \end{cases} \quad (1)$$

where $n_k^{\setminus i}$ denotes the number of the customers at a table k except for i and α is a concentration parameter.

Markov Chain Monte Carlo Methods for Clustering

Markov chain Monte Carlo (MCMC) methods[3] are algorithmic methods to sample from posterior distributions. If conditional posterior distributions are given by models, it is the best way in order to acquire parameters as posterior distributions. The algorithm for this package is given by

i) Sampling z_i for each i ($i = 1, 2, \dots, n$)

$$p(z_i = k | x_{1:n}, z_{1:n}^{\setminus i}, \alpha, \mu_k, \mu_0, \Sigma_k, \rho_0) = \begin{cases} p(x_i | \mu_k, \Sigma_k) \times \frac{n_k^{\setminus i}}{n-1+\alpha}, \\ p(x_i | \mu_k, \Sigma_k) \times \frac{\alpha}{n-1+\alpha} \end{cases} \quad \mu_k \sim N(\mu_0, \Sigma_{table}). \quad (2)$$

$$z_i \sim \text{Multi}(p(z_i = 1), p(z_i = 2), \dots, p(z_i = \infty)), \quad (3)$$

where k is a k th cluster and i is a i th data. *mu* and *sigmatable* arguments in the "crp_gibbs" function are generating μ_k parameter for new table.

ii) Sampling μ_k for each k ($k = 1, 2, \dots, \infty$)

$$\mu_k \sim p(\mu_k | x_{1:n}, z_{1:n}, \Sigma_k, \mu_0, \rho_0) = N(\mu_k | \frac{n_k}{n_k + \rho_0} \bar{x}_k + \frac{\rho_0}{n_k + \rho_0} \mu_0, \Sigma_k), \quad (4)$$

$$\Sigma_{k_{ij}} = \frac{n_k}{n_k + \rho_0} \text{Cov}(x_{ki}, x_{kj}) + \frac{\rho_0}{n_k + \rho_0} (\bar{x}_{ki} - \mu_{0i})(\bar{x}_{kj} - \mu_{0j}), \quad (5)$$

$$\bar{x}_k = \frac{1}{n_k} \sum_{i=1}^n \delta(z_i = k) x_i. \quad (6)$$

Iterations i) ii) continue on many times, and Σ_k is a variance-covariance matrix of k th cluster. i and j are rows and columns' number of Σ_k . ρ_0 is a argument *rho0* in "crp_gibbs" function. First several durations of iterations which are called as "burn in" are error ranges. For that reason, "burn in" durations are abandoned.

Clusters Entropy

Entropy denotes the ambiguity of clustering. As a result of a simulation, data x_i joins in a particular table. From the total numbers n_k of the particular table k at the last iteration, a probability p_k at each cluster k is calculated. The entropy equation is given by

$$\text{Entropy} = - \sum_{k=1}^{\infty} \frac{n_k}{n} \log_2 \frac{n_k}{n}. \quad (7)$$

Installation

CRPclustering is available through GitHub (<https://github.com/jirotubuyaki/CRPclustering>) or CRAN (<https://CRAN.R-project.org/package=CRPclustering>). If download from GitHub, you can use devtools by the commands:

```
> library(devtools)
> install_github("jirotubuyaki/CRPclustering")
```

Once the packages are installed, it needs to be made accessible to the current R session by the commands:

```
> library(CRPclustering)
```

For online help facilities or the details of a particular command (such as the function `crp_gibbs`) you can type:

```
> help(package="CRPclustering")
```

Methods

Implemented by Scala

This package is implemented by Scala. Please install Scala compiler version 2.12.6. Programs are compiled java archive. If you are interested in source codes by Scala. Please check GitHub (<https://github.com/jirotubuyaki/CRPCLustering>).

Method for Chinese Restaurant Process Clustering

```
> result <- crp_train(as.matrix(data),
                      mu=c(0,0),
                      sigma_table=1,
                      alpha=1,
                      ro_0=1,
                      burn_in=100,
                      iteration=1000
                      )
```

This method calculates CRP clustering. Let arguments be:

- data: a matrix of data for clustering. row is each data i and column is dimensions of each data i .
- mu: a vector of center points of data. If data is 3 dimensions, a vector of 3 elements like "c(2,4,7)".
- sigma_table: a numeric of table position variance.
- alpha: a numeric of a CRP concentration rate.
- ro_0: a numeric of a CRP mu change rate.
- burn_in: an iteration integer of burn in.
- iteration: an iteration integer.

Let return be:

- result: an array denotes result clusters. First column is cluster number. Second is joined data number for each cluster. From next, result clusters' mean and variance matrix.

Predict Cluster Data Joined

```
> predict <- crp_predict(as.matrix(data), result)
```

Let arguments be:

- data: a matrix of data for clustering. row is each data i and column is dimensions of each data i .
- result: return result from method "crp_train".

Let return be:

- predict: an array denotes first column is joined cluster and nexts are joined probability for each result cluster.

Visualization Method

```
> crp_plot(as.matrix(data), predict)
```

This method exhibits multi dimensional plot matrix. Let arguments be:

- data: a matrix of data for clustering. Row is each data i and column is dimensions of each data i .
- predict: return predict from method "crp_predict".

Example

Data is generated from ten normal distributions and parameters are set as $\mu = c(0, 0, 0, 0, 0, 0, 0, 0, 0, 0)$, $\alpha = 1$, $\text{sigmatable} = 1$, $\rho_0 = 1$, $\text{burnin} = 100$, $\text{iteration} = 1000$. The result is plotted on a graph and each data joins in any cluster. The graph is given by below:

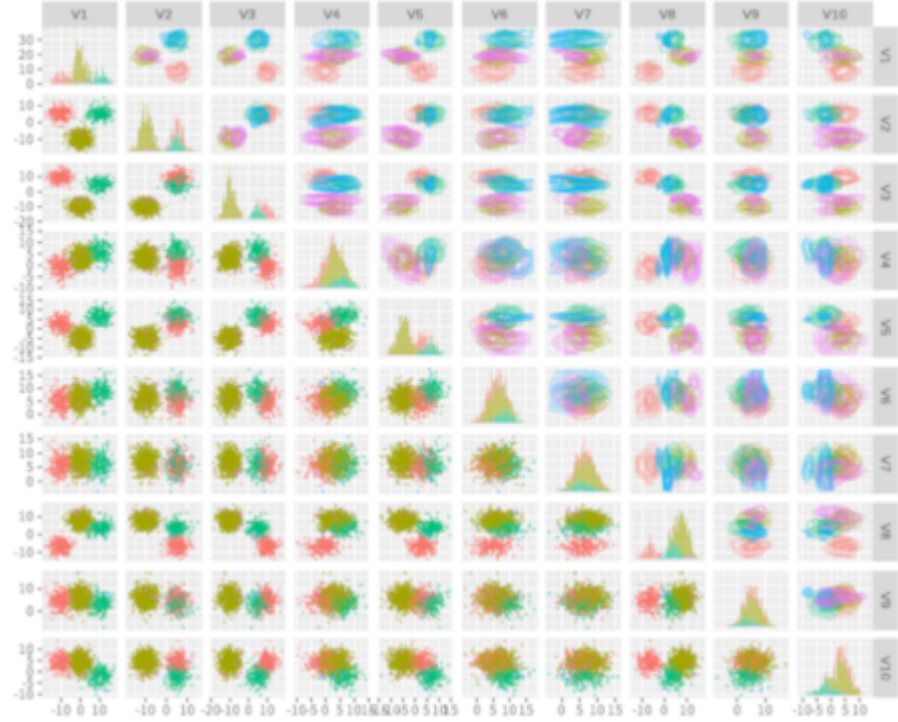


Figure 1: Result of clustering

Conclusions

Chinese restaurant process clustering was implemented and explained how to use it. Computer resources are limited. Computer processing power is the most important problem. After this, several improvements are planned. Please send suggestions and report bugs to okadaalgorithm@gmail.com.

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References

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