

# Tutorial for CMI2NI

## 1. Introduction

CMI2NI is a MATLAB program for inferring Gene Regulatory Networks (GRN) from gene expression dataset based on Conditional Mutual Inclusive Information (DMI) which can accurately measure direct associations or causal strength between genes.

## 2. Matlab Code description

### 2.1 CMI2NI

```
[G,Gval,order]=CMI2NI(data,lamda,order0)
```

#### 2.1.1 Input

| Code          | Description  |
|---------------|--|
| <b>Data</b>   | Expression of target gene.   |
| <b>lamda</b>  | Parameter for threshold of dependence.   |
| <b>order0</b> | The parameter to end the program when order meet order_fix given; Default value is null. |

#### 2.1.2 Output

| Code         | Description                             |
|--------------|---|
| <b>G</b>     | 0-1 network or graph inferred by CMI2NI |
| <b>Gval</b>  | Network with strengthens                |
| <b>order</b> | Order number of algorithm terminated.   |

## 2.2 Example\_CMI2NI.m

As an example, 'Example\_CMI2NI.m' is to infer gene regulatory network from gene expression dataset.

```
% An example of using CMI2NI for inferring gene regulatory
networks from gene
% expression data.

clear; clc;

% Data importation
% data: Rows are genes and columns are samples.
load example_data

% Parameter setting
lamda= 0.03;

[net,net_value,order]=CMI2NI(data,lamda);
net,net_value

[net2,net_value2,order]=CMI2NI(data,lamda,1);
net2,net_value2
```

## 3. Contact information

If you encounter any problem, please do not hesitate to contact us at

zhang-xiujun@163.com