

# Bayesian Network Inference Program

## Overview

This program performs inference on a Bayesian network using four algorithms:

1. Exact Inference
2. Prior Sampling
3. Rejection Sampling
4. Likelihood Weighting

It calculates probabilities for query nodes based on given evidence.

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## Usage

### Command Format

```
python <script.py> "[<evidence1,value1><evidence2,value2>][query_node1,query_node2]" -  
-algorithm <algorithm> --sample_count <count> --runs <num_runs>
```

### Input Details

#### 1. Conditions:

- Specified in the format `<node,value>`.
- Multiple pieces of evidence are enclosed in `[]` and separated by `><`.

#### 2. Query Nodes:

- Specified as a comma-separated list of variables within `[]`.

#### 3. Algorithm:

- Specifies the inference algorithm to use from:
  - `exact` : Performs exact inference.
  - `prior` : Uses prior sampling.
  - `rejection` : Uses rejection sampling.
  - `likelihood` : Uses likelihood weighting.

#### 4. Sample Count ( `--sample_count` ):

- Number of samples to generate for sampling algorithms (Default is set to `10,000`).

#### 5. Runs ( `--runs` ):

- Number of iterations to run for the algorithms. Default is set to `1`.
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## Testing Examples

*The testing commands provided are examples and can be modified as needed to specify different evidence, query nodes, algorithms, sample sizes, and runs to get the desired results.*

## 1. Exact Inference

To run Exact Inference for Case 1 :

```
python script.py "<A,f>[B,J]" --algorithm exact
```

## 2. Prior Sampling

To run Prior Sampling for Case 2 :

```
python script.py "<J,t><E,f>[B,M]" --algorithm prior --sample_count 10000 --runs 1
```

## 3. Rejection Sampling

To run Rejection Sampling for Case 3 :

```
python script.py "<M,t><J,f>[B,E]" --algorithm rejection --sample_count 10 --runs 5
```

## 4. Likelihood Weighting

To run Likelihood Weighting for Case 1 :

```
python script.py "<A,f>[B,J]" --algorithm likelihood --sample_count 1000 --runs 10
```

## Output Format

The output is formatted as:

```
<node1,probability1><node2,probability2>...
```

Example:

```
<B,0.0010><J,0.0500>
```

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## Author

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