



# Mocapy++

implementation of a  
mixed discrete/continuous node



# Mocapy: Graphical Models: Dynamic Bayesian Networks

## Mocapy: Dynamic Bayesian Networks topology

Directed Acyclic Graphs

nodes are probability distributions  
arcs encode conditional independences  
all the slices are equal  
first order dependences between slides

Topological independences → computational efficiency

## First implementation: **Monte Carlo Python**

now faster implementation in C++: Mocapy++ (same architecture)

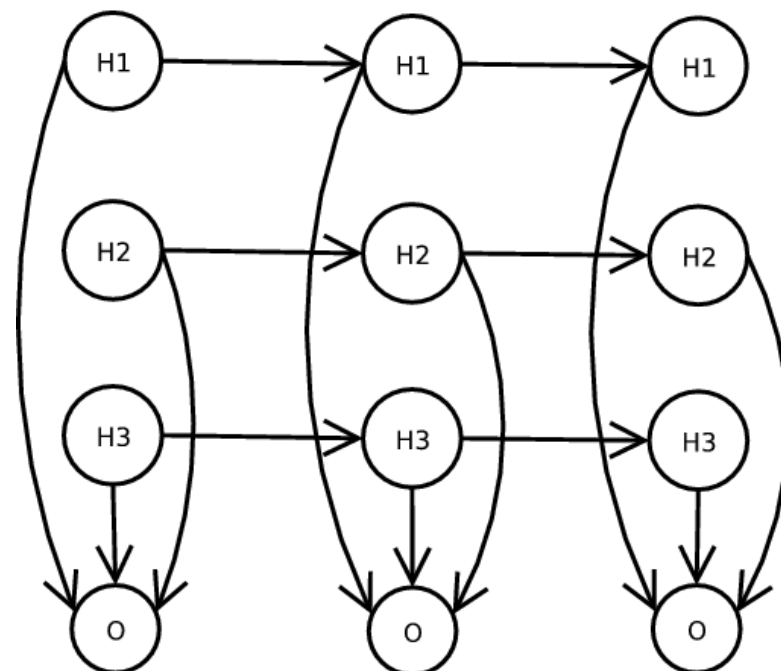
## Your task: to extend Mocapy++, defining a new node type

Maximum likelihood estimation

Density calculation

Sampling

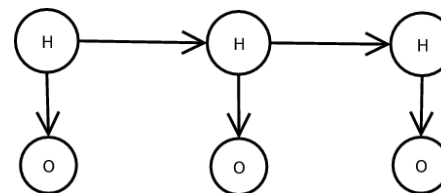
Technicality: data storage: save/load data, and save/restore models



# Mocapy++: background overview

## Mocapy: Inference

identify the distribution of hidden nodes  $\mathbf{h}$   
given the model and the observed quantities



$$P(\mathbf{h} \mid \mathbf{x}, \theta)$$

## Mocapy: Learning

estimate the parameters of the model

*approximate* Expectation-Maximization:  
iterative Maximum Likelihood while (Markov) blanket sampling

$$\theta_{\text{ML}} = \arg \max_{\theta} \sum_{\mathbf{h}} P(\mathbf{h}, \mathbf{d} \mid \theta)$$

$$\theta_{\text{new}} = \arg \max_{\theta} \sum_{\mathbf{h}} P(\mathbf{h} \mid \mathbf{d}, \theta_{\text{old}}) \log P(\mathbf{h}, \mathbf{d} \mid \theta)$$

*in practice: call the Stochastic-EM functionalities, in the Engine section of the framework*



# Mocapy++: framework usage

Your test case should mimic *examples/hmm\_simple.cpp*

Nodes: definitions, parameters

DBN: topology definition

Sampling algorithm: Gibbs MC

EM engine: sampling + DBN object (Stochastic-EM)

load (in this order):

mismask: label for the node values (**missing mask**)

0: MOCAPY\_OBSERVED

1: MOCAPY\_HIDDEN

2: MOCAPY\_MISSING

data: one value per dimension,

in the order specified by the DBN topology

perform the learning (Stochastic-EM)

print the learned nodes

```
#include <iostream>
#include <stdio.h>
#include <stdlib.h>

#include "mocapy.h"
using namespace mocapy;
using namespace std;

int main(void) {
    // The dynamic Bayesian network
    DBN dbn;

    // Nodes in slice 1
    Node* h1 = NodeFactory::new_discrete_node(5, "h1");
    Node* o1 = NodeFactory::new_discrete_node(2, "o1");

    // Nodes in slice 2
    Node* h2 = NodeFactory::new_discrete_node(5, "h2");

    // Set architecture
    dbn.set_slices(vec(h1, o1), vec(h2, o1));

    dbn.add_intra("h1", "o1");
    dbn.add_inter("h1", "h2");
    dbn.construct();

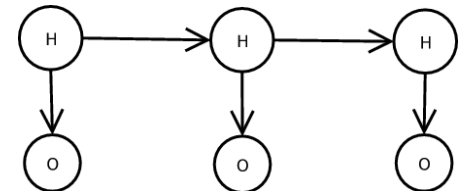
    cout << "Loading traindata" << endl;
    GibbsRandom mcmc = GibbsRandom(&dbn);

    EMEngine em = EMEngine(&dbn, &mcmc);
    em.load_mismask("data/mismask.dat");
    em.load_weights("data/weights.dat");
    em.load_sequences("data/traindata.dat");

    cout << "Starting EM loop" << endl;
    for (uint i=0; i<100; i++) {
        em.do_E_step(20, 10);
        double ll = em.get_loglik();
        cout << "LL= " << ll << endl;
        em.do_M_step();
    }

    cout << "h1: " << *h1 << endl;
    cout << "o1: " << *o1 << endl;
    cout << "h2: " << *h2 << endl;

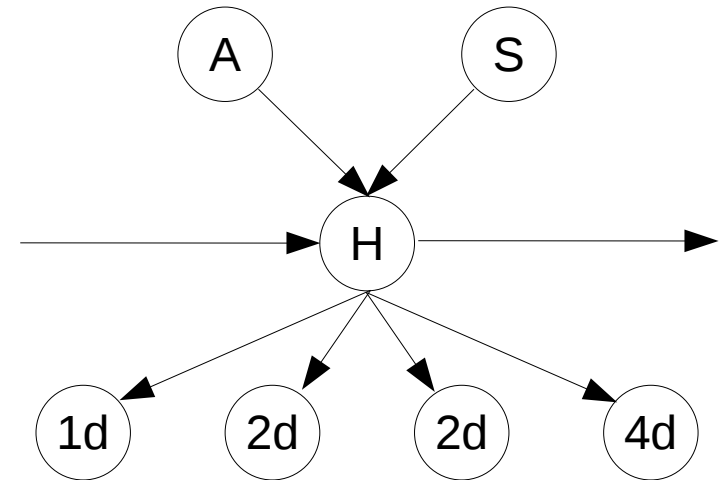
    return EXIT_SUCCESS;
}
```



# Mocapy++: datafiles

## Mocapy: format for datafiles

```
// File format:
// A line contains values in a slice
//     separated by "sep" (default space).
// A block of lines therefore represents a sequence.
// An empty line separates two sequences.
// Lines starting with # are ignored
```



input.mismask (order: has to be consistent with topology in .cpp file):

0	0	1	0	0	0	0
0	0	1	0	0	0	0
0	0	1	0	0	0	0
0	0	1	0	0	0	0

input.data (order: consistent with mismask):

8	3	0	0.4286	-4.2688	-2.6449	-2.4226	-2.3128	-0.9675	-0.7078	-0.8320	-0.6090
8	2	0	0.5385	-4.4054	-2.4140	-2.2166	-1.6194	-3.8429	-1.6061	-1.8543	-0.9892
17	2	0	0.3750	-4.4473	-2.3684	-2.6483	-1.8028	-4.5492	-2.6714	-2.5935	-2.3782
8	2	0	0.3636	-4.5259	-2.3819	-2.2348	-1.5630	-4.6230	-2.7179	-2.5563	-1.9557



# Mocapy++: download, compilation

Mocapy++: download from [sourceforge.net](https://sourceforge.net) (anonymous svn)

```
svn co https://mocapy.svn.sourceforge.net/svnroot/mocapy/Mocapy .
```

Compilation: cmake. Your test program can be compiled

as an external program:

```
<include mocapy.h>
```

link with: mocapy++, lapack, boost.serialization libraries

as an example in the examples/ directory (easier):

place your code in the examples/ directory

edit the CMakeLists.txt

```
SET(PROGS ...)
```

Manual: file /MANUAL.pdf in the root of the repository

ch. 2: installation

ch. 1,3: theoretical foundations

ch. 4: I/O, general usage

ch. 6, 7: design, extension



# Mocapy++: your task: overview

Your test case should mimic *examples/hmm\_simple.cpp*

input.mismask:

1	0
1	0
1	0

input.data (here 2 sequences):

# first column: hidden node  
 # first dimension: 0: indicator for discrete  
 # first dimension: 1: indicator for continuous

0	0	3
0	0	2
0	1	0
0	0	0
0	1	0.5
0	1	3.75

0	0	3
0	0	1

# errors: skip them, but warn the user

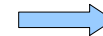
0	0	3.75
0	0	-5

```
#include <iostream>
#include <stdio.h>
#include <stdlib.h>

#include "mocapy.h"
using namespace mocapy;
using namespace std;
```

```
int main(void) {
    // The dynamic Bayesian network
    DBN dbn;

    // Nodes in slice 1
    Node* h1 = NodeFactory::new_discrete_node(5, "h1");
    Node* o1 = NodeFactory::new_discrete_node(2, "o1");
```



```
// Nodes in slice 2
Node* h2 = NodeFactory::new_discrete_node(5, "h2");
```

```
// Set architecture
dbn.set_slices(vec(h1, o1), vec(h2, o1));
```

```
dbn.add_intra("h1", "o1");
dbn.add_inter("h1", "h2");
dbn.construct();
```

```
cout << "Loading traindata" << endl;
GibbsRandom mcmc = GibbsRandom(&dbn);
```

```
EMEngine em = EMEngine(&dbn, &mcmc);
em.load_mismask("data/mismask.dat");
em.load_weights("data/weights.dat");
em.load_sequences("data/traindata.dat");
```

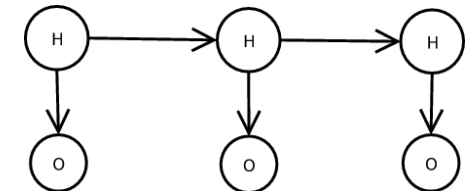


```
cout << "Starting EM loop" << endl;
for (uint i=0; i<100; i++) {
    em.do_E_step(20, 10);
    double ll = em.get_loglik();
    cout << "LL= " << ll << endl;
    em.do_M_step();
}
```

```
cout << "h1: " << *h1 << endl;
cout << "o1: " << *o1 << endl;
cout << "h2: " << *h2 << endl;
```

```
return EXIT_SUCCESS;
```

```
}
```



# Mocapy++: your task

## Mocapy: your task: high level

Input: categorical and continuous

algorithms from DiscreteNode class. Continuous estimation: extract code from GaussianNode (skip the multidimensional and shrinkage case)

Define a 2d node

first dimension: indicator (0: discrete, 1: continuous)

second dimension: value of the node

this node requires exactly one discrete parent

Learning: 2 steps (to do for each value of the single discrete parent):

learn the ratio between categorical and continuous output types

simply count the number of categorical, divided by the number of valid observations

learn the parameters for the two different sets of value types, independently

we split the input dataset in categorical dataset and Gaussian dataset

Sampling: 2 steps

sample a random number (a percentage)

lower or equal to categorical ratio threshold? Sample from the DiscreteNode algorithms

higher? Sample from GaussianNode algorithms





# Mocapy++: architecture

## Mocapy++: framework and changes

Engine: you simply need to use it

as you see, the node object is  
the glue between the engine and  
the distribution frameworks

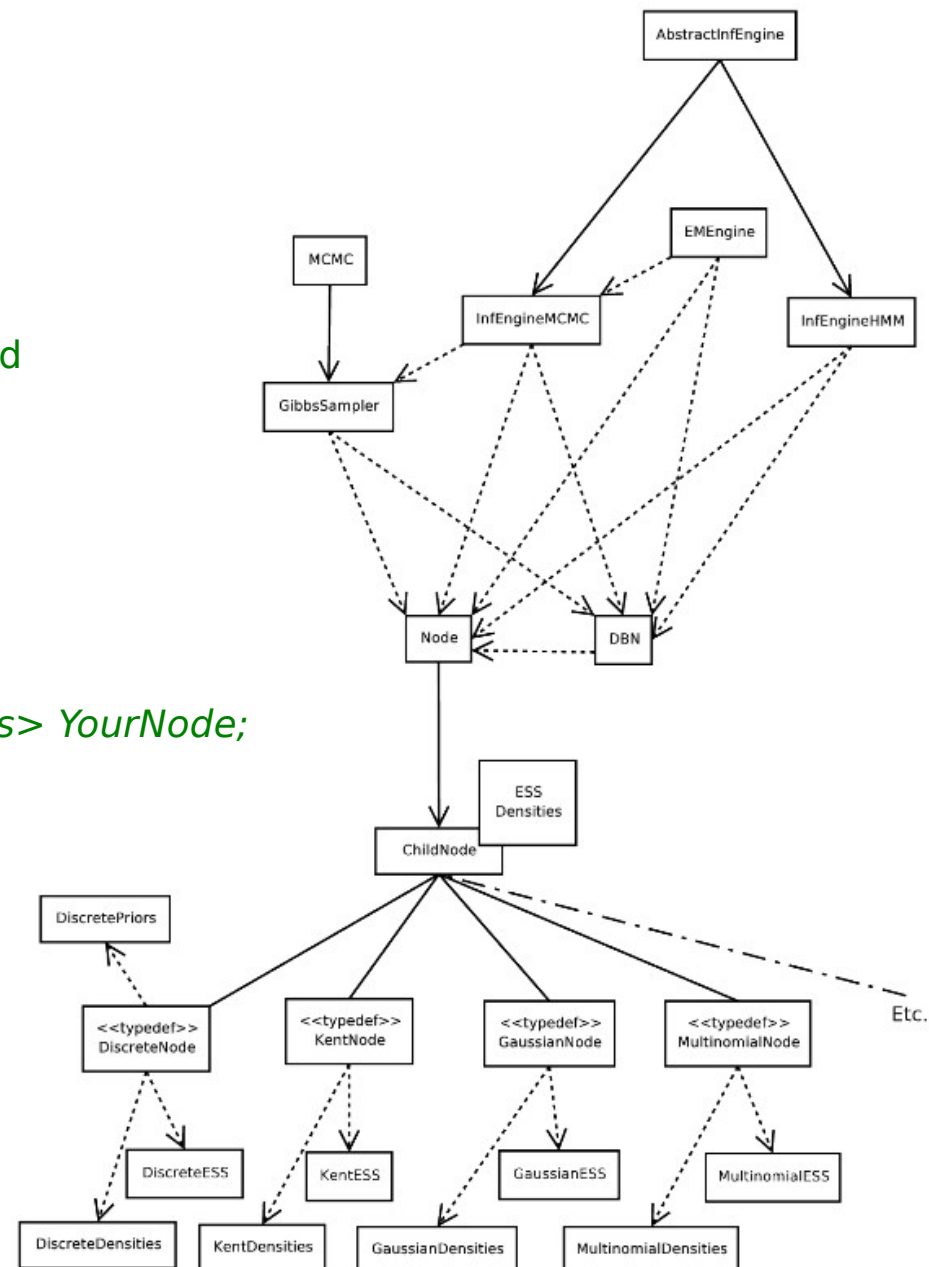
Node → ChildNode template

*template Childnode<xESS, xDensities> YourNode;*

Node Internals

ESS: Expected Sufficient Stats  
add/store sample  
for hidden node operations

densities: estimation, sampling



# Mocapy++: your task: details

Mocapy: your task: part of the framework to change and deliver

src/framework/dbn.h:

include lines

typedef ChildNode<YourESS, YourDensities> YourNode;

src/yournode/youress.{h,cpp}

YourESS class, inherits from abstract ESSBase. Override:

construct(): define the appropriate shape of the ESS

add\_ptv(): add a sample point to the ESS

src/yournode/yourdensities.{h,cpp}

YourDensities class, inherits from abstract DensitiesBase. Override:

construct(parent\_sizes): initialize parameters (mean, covariance, CPD, etc.).

estimate(ess): estimate the parameters of the node based on the ESS

sample(ptv): return a sample based on the parent values

get\_lik(ptv, log): return the likelihood  $P(\text{child}|\text{parent})$

get\_parameters(): return the parameters of the distribution

examples/yournode\_hmm.cpp

as per simple\_hmm.cpp (edit the CMakeLists.txt)

make sure the dbn.save() and dbn.load() methods work



# Mocapy++: your addenda

## Mocapy: simple example on how to wrap the two nodes

Let's suppose the categorical output has probability 0.3

### sampling

Sample the threshold. Is it lower than 0.3?

Sample from the categorical routines

Else, sample from the Gaussian routines

### Learning

Discrete set: discrete/discretedensities.\*, discrete/discreteess.\*

Continuous set: gaussian/\*

start: examples/hmm\_gauss\_1d.cpp

public functions: construct(), estimate(), get\_lik()

private functions: calc\_means(), calc\_cov\_full() (copy-and-paste is the easier option)

### Likelihood estimation for a given parent value

Note:  $\theta$  is the probability vector of the discrete distribution for that parent value

Note: G is Gaussian with mean and std. dev.  $\mu, \sigma$  for that parent value

$P(\text{categorical}, 2) = P(2 | \theta) * P(\text{categorical}) \leftarrow 0.3$

$P(\text{continuous}, 3.5) = G(3.5 | \mu, \sigma) * P(\text{continuous}) \leftarrow 0.7$

Details for the instantiation of the nodes: src/framework/nodefactory.\*, examples/\*.cpp



# Mocapy++: overview of the task

## Mocapy++: task

download  
compile

write ESS, Densities (*mocapy++ manual*)  
define node as a specialization of ChildNode

load data file, mismask

write your test program

Deliver: tar.gz archive

report  
your code (test included)  
your data files

