

Quantitative Modelling of Adverse Outcome Pathway for Risk Assessment

WANG GAO (INERIS UTC)

SOUS DIRECTION DE

GHISLAINE GAYRAUD (LMAC UTC)

FRÉDÉRIC Y. BOIS (INERIS)

Overview

I - Introduction

- I.1 Toxicology in general
- I.2 Project Eu-ToxRisk21
- I.3 AOP and qAOP
- I.4 Objectives of my thesis

II - Probabilistic Model

- II.1 BN : Bayesian Networks
- II.2 DBN : Dynamic Bayesian Networks
- II.3 Model family for qAOP

III - Learning problem

IV - Conclusion

I. Introduction

I.1. Toxicology in general

- Toxicology

- Application domains: Cosmetics, Drug, etc.

- Objective: Risk assessment

- Approaches (Different types of routines)

- *In vivo* test: Animal experiments
- Alternative methods
 - *In vitro* test (Classic) - fast
 - *In vitro* test (advanced): High throughput screening - very fast
 - *In silico* test: computational modelling and simulation - we are working on this



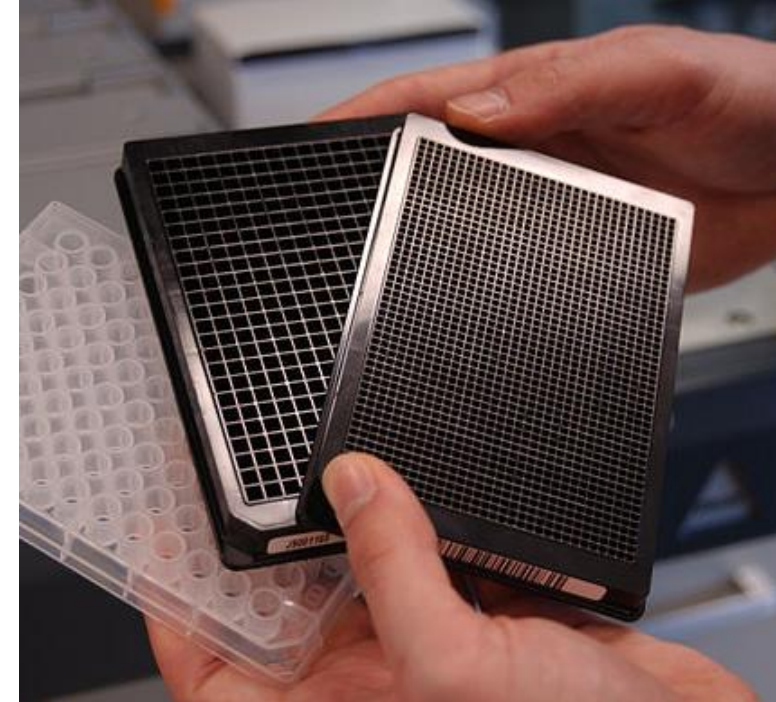
High-throughput screening

Robotic tools

384, 1536, 3456 wells

- Fast

- Low-cost



1.2. Project Eu-ToxRisk21

“An Integrated European ‘Flagship’ Programme Driving **Mechanism-based Toxicity Testing** and **Risk Assessment** for the 21st century.”

Organisation :

- ❖ Long range 2016-2020 research program
- ❖ 41 international research teams (40 EU + 1 USA) from 13 countries
- ❖ 14 Work packages with sub-objectives
 - ❖ WP 10 : **Computational Modelling for Risk assessment**

Motivation :

- ❖ Partially replace animal experimentations
- ❖ Improve the predictive methods based on high-throughput toxicity tests

15 Case studies

Liver

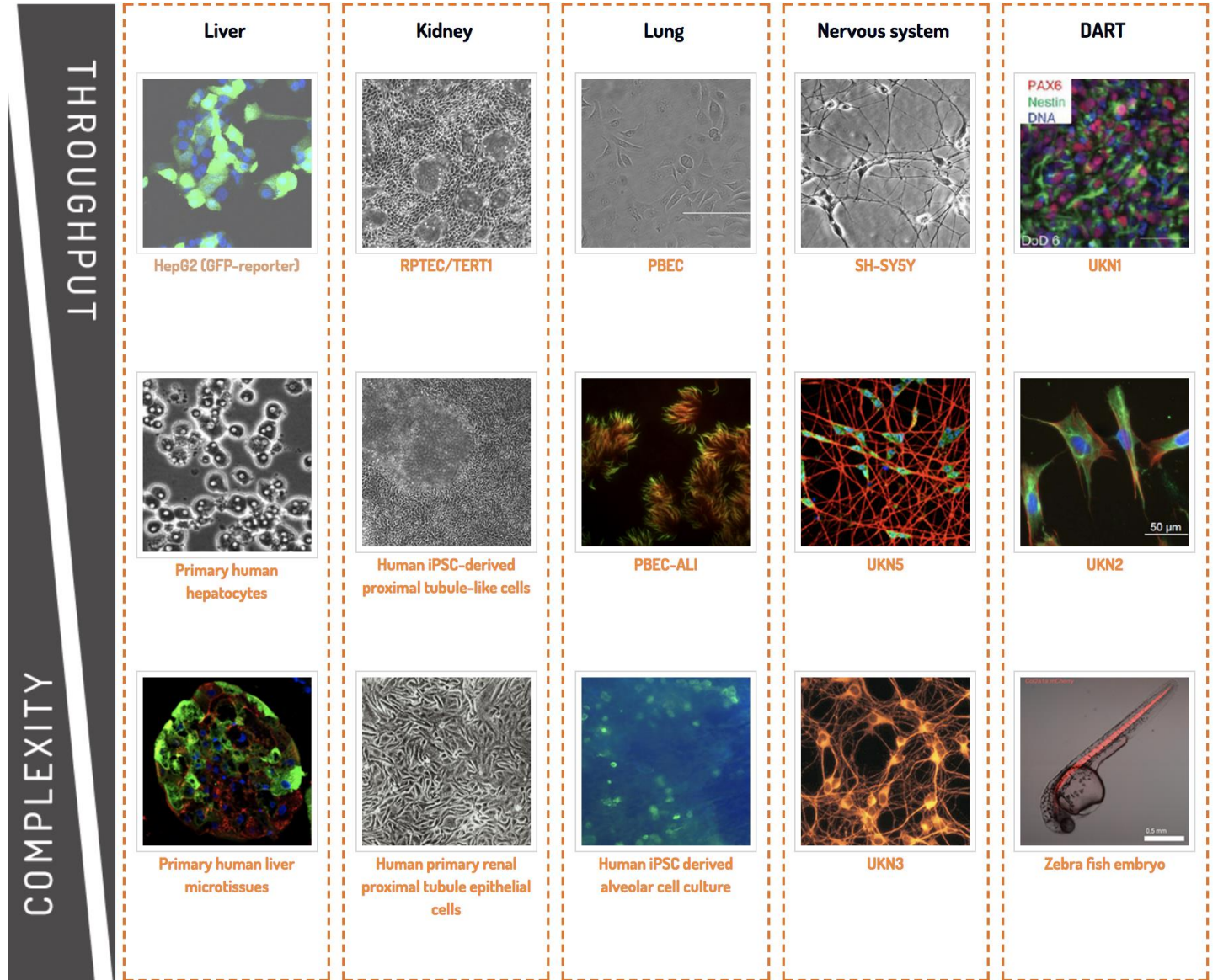
Kidney

Lung

Nervous system

Etc.

15 Cases are studied in the program
Eu-ToxRisk21



I.3. AOP and qAOP

❖ EU-ToxRisk21

- An Integrated European 'Flagship' Programme Driving **Mechanism-based** Toxicity Testing and **Risk Assessment** for the 21st century

❖ **AOP** : Adverse outcome pathway – qualitative tool for **mechanism description**

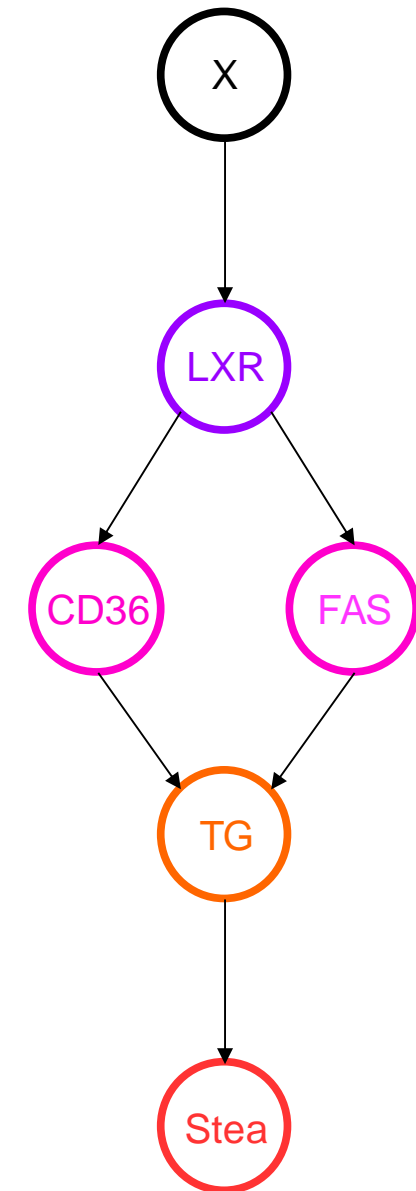
❖ **qAOP** : quantitative AOP for **risk assessment**

Remark : Each case study corresponds to a set of AOPs.

AOP and qAOP

AOP : Adverse outcome pathway

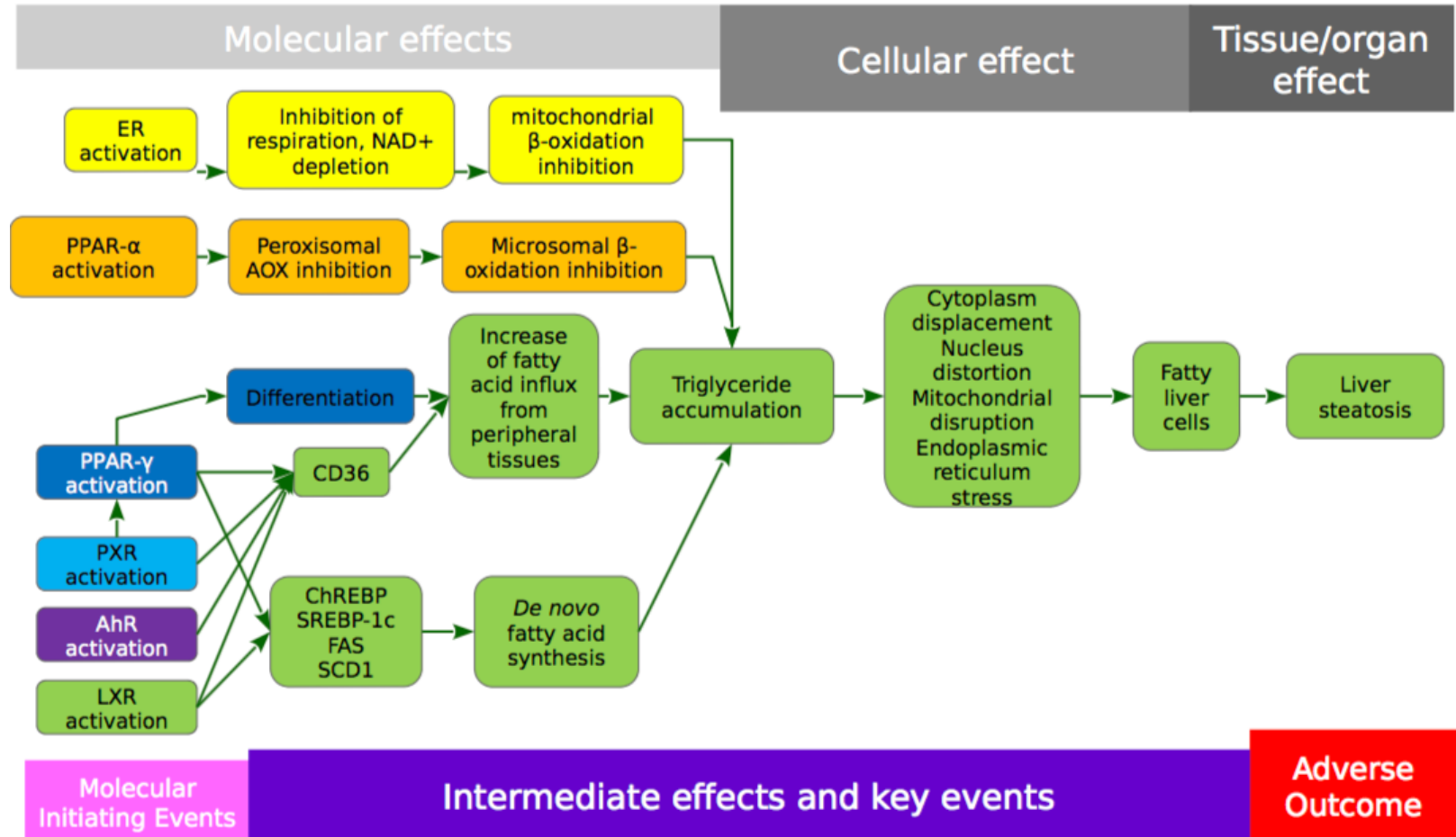
- ❖ Knowledge Exchange
- ❖ A qualitative tool for the structural representation of causal relationships (Dose-response)
 - ❖ Starting from a molecular perturbation (MIE)
 - ❖ Through key events (KE)
 - ❖ Arriving at an adverse outcome (AO)
- ❖ In this case : AO = Steatosis (fat liver)
- ❖ qAOP = quantitative AOP:
 - ❖ **Probabilistic model**
 - ❖ Predictive capacity : Decision Support Tool



A richer version of AOP : AOP Network

- Each node may have more than one parent.
- One node can be involved in different AOPs

9 AOPs are hidden here



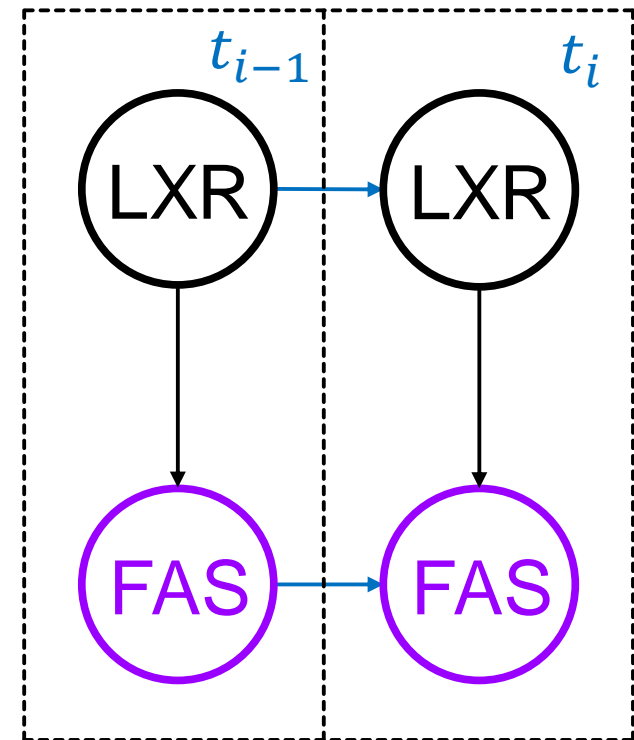
1.4. Objectives of my thesis

Objectives :

- ❖ Probabilistic modelling of AOP =: qAOP
- ❖ => Prediction of AO

Action plan :

- ❖ Fixed structure for mini AOP
 - ❖ Modelling of the strength cause-effect
 - ❖ Prediction
- ❖ Structural learning : more ambitious



II. Probabilistic Model

II.1. BN : Bayesian Networks

Def : BN : $(\mathcal{G}, P_{\mathcal{G}})$ with

- \mathcal{G} : BN Structure : DAG directed acyclic graph $\mathcal{G}(V, E)$
 - V : Set of vertices, nodes : variables $V = \{X_i | i \in 1:N\}$
 - E : Set of directed edges : causality relationships
- $P_{\mathcal{G}}$: multivariate distribution over V
 - specified as set of local conditional probability distribution (CPDs) associated with \mathcal{G} 's nodes.

$$P_{\mathcal{G}}(X) = \prod_{i=1:N} P_{\mathcal{G}}(X_i | \text{Par}(X_i))$$

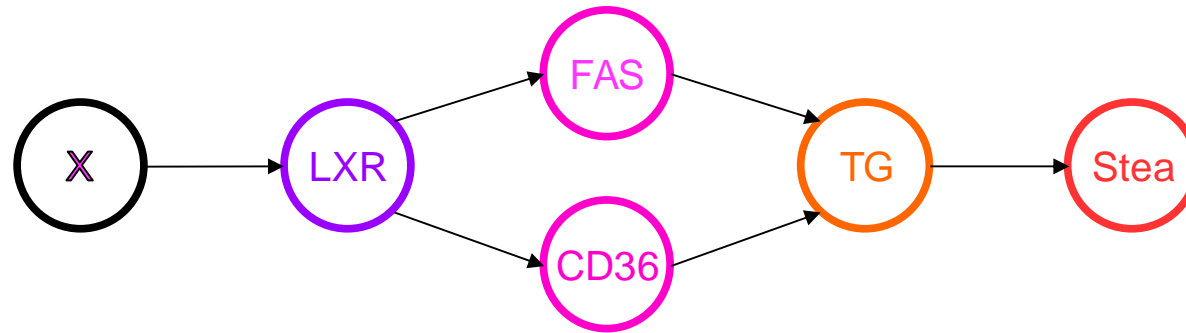
with $X = (X_1, \dots, X_i, \dots, X_N)$ system stat vector

BN : Bayesian Networks

$$P_G(X) = \prod_{i=1:N} P_G(X_i | \text{Par}(X_i))$$

Local conditional distributions (CPD) and the DAG completely determine the joint distribution

Example : steatosis case



$$P(X, L, C, F, T, S) = P(X)P(L|X)P(C|L)P(F|L)P(T|C, F)P(S|T)$$

Problem in the context of qAOP : Child node = function (Parent node(s) , **time**)

II. 2. DBN : Dynamic Bayesian Networks

Extension of BNs to handle temporal models

Assumptions:

- the timeline discretised into a set of *time slices* :

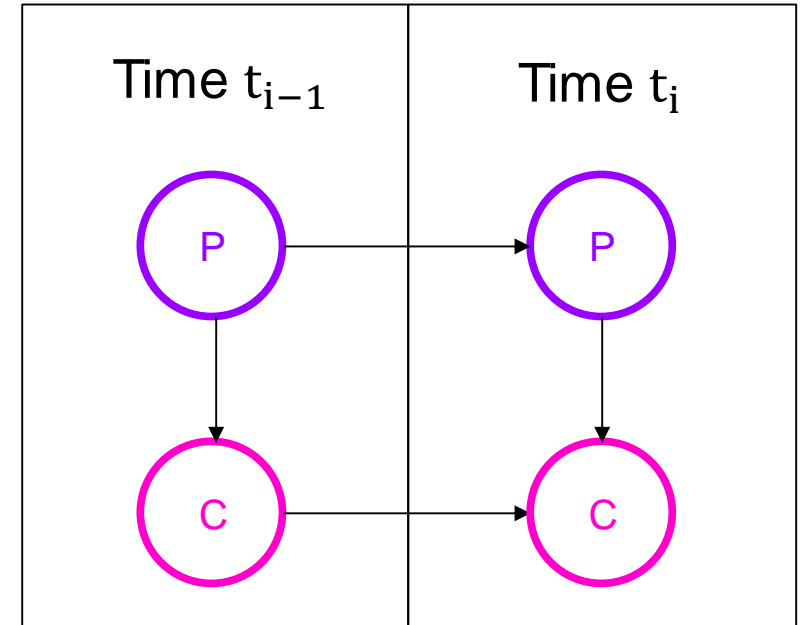
$X^{(t_0)}, X^{(t_1)}, \dots, X^{(t_m)}$, with m the number of observations

- Markov assumption for a dynamic system over the template variables $X : \forall i \in \mathbb{N}^+$

$$X^{(t_{i+1})} \perp X^{(t_0:(t_{i-1}))} | X^{(t_i)}$$

Two types of dependency :

- Inter time-slice dependency (between time-slices)
- Intra time-slice dependency (in the same time-slice)



C child node : C_t child node at time t
 P parent node : P_t parent node at t

DBN : Dynamic Bayesian Networks

Markov assumption :

$$X^{(t_{i+1})} \perp X^{(t_0:(t_{i-1}))} | X^{(t_i)}$$

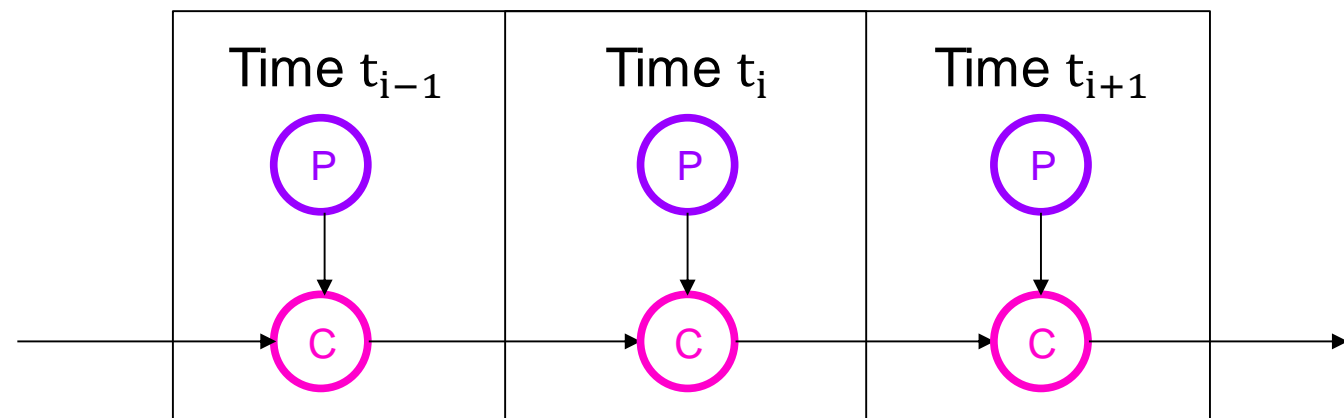
Compact definition of the joint probability distribution in DBN

$$P(X^{(t_0:t_m)}) = P(X^{(t_0)}) \prod_{i=1}^m P(X^{(t_i)} | X^{(t_{i-1})})$$

Example : Linear Dynamic System

Classic linear DBN

$$\mathbb{E}[C_{t_i}] = \alpha + \beta_{prev} C_{t_{i-1}} + \beta_{curr} P_{t_i}$$



Insights about qAOP

Data visualization DEMO : Real data for kidney disease case study

Dynamic 3D plot GSH-DCF-Time ([online](#)) ([local](#))

- ❖ S_C : Stationary state (Saturation level) of child node, denote :
- ❖ $S_C = f(P_t)$: Dependence of S_C on P_t
- ❖ C_t : Child node activity at time t
- ❖ C_t converges to S_C over time



II.3. Model family for qAOP : Embryonic form

$$S_C [P_t] - \mathbb{E}[C_t] = (S_C [P_t] - C_{t-h})e^{-\nu h}$$

- C_t : Child node activity at time t , (**observed**)
- P_t : Parent node(s) activity at time t , I could be a vector (**observed**)
- $S_C [P_t]$: stationary state of child node given its parent(s) (**unobserved**)

Questions :

$$S_C [P_t] = ?$$

$h \in \mathbb{R}^+$: non regular observation ?

$\nu > 0$: to ensure the convergence of C_t towards $S_C (P_t)$

Model family for qAOP : Linear model

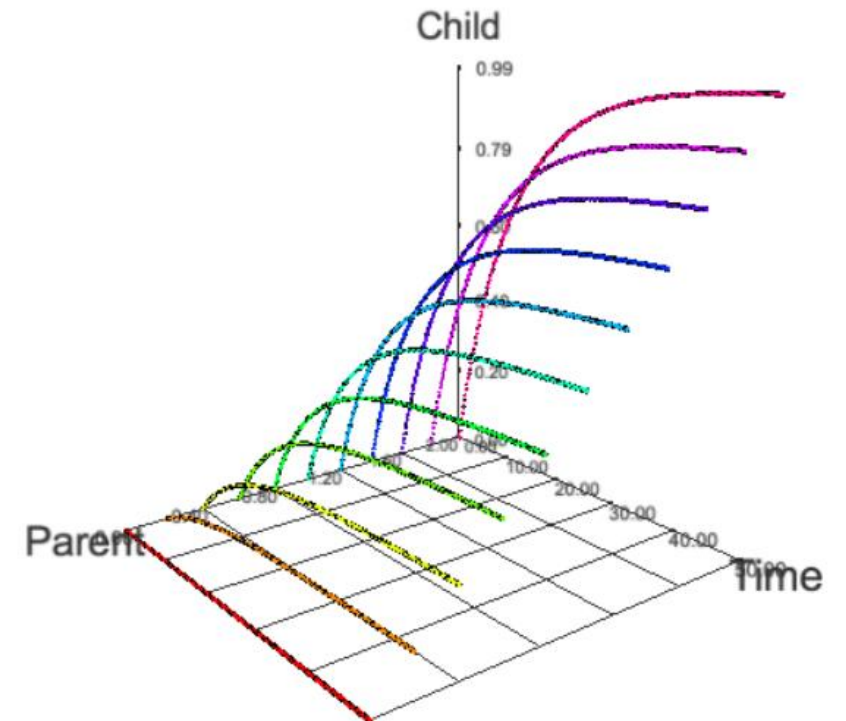
$$\begin{cases} S_C[P_t] - \mathbb{E}[C_t] = (S_C[P_t] - C_{t-h})e^{-\nu h} \\ S_C[P_t] = \beta P_t + \beta_0 \\ h \in \mathbb{R}^+ \end{cases} \quad (\mathcal{M}_L)$$

Assumption : Stationary stat of Child node is a **linear function** of parent node(s)

Remark : The classic linear DBN model is a special case of (\mathcal{M}_L)

- $S_C[P_t] = \frac{\beta_{curr}}{1-\beta_{prev}} P_t + \frac{\alpha}{1-\beta_{prev}}$
- $h = 1$
- $e^{-\nu h} = e^{-\nu} = \beta_{prev}$

$$\mathbb{E}[C_{t_i}] = \alpha + \beta_{prev} C_{t_{i-1}} + \beta_{curr} P_{t_i}$$



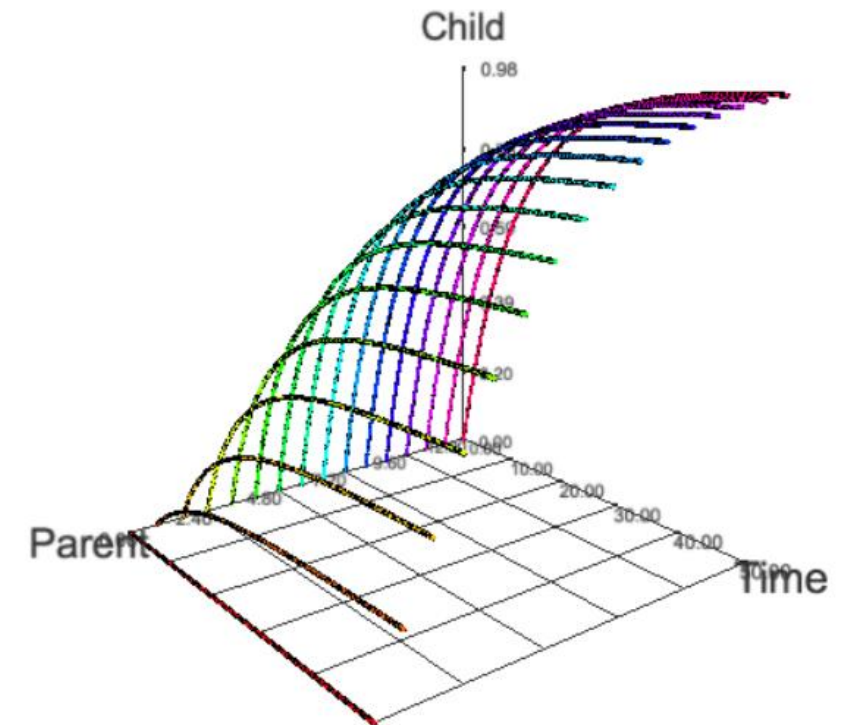
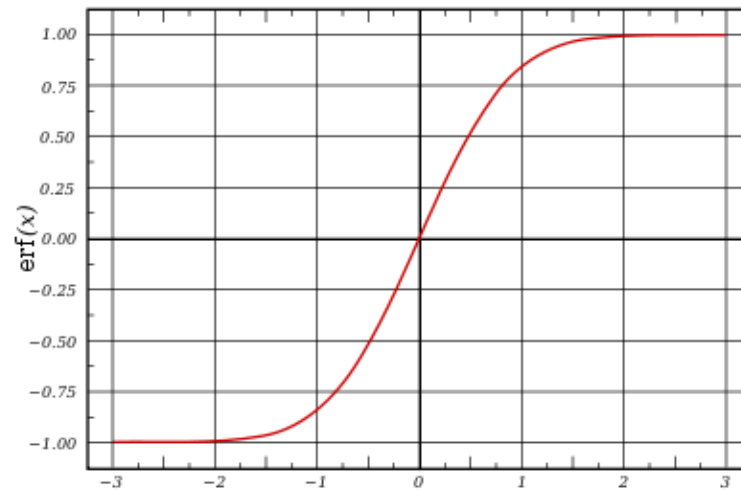
[Demo \(online\)](#)

Model family for qAOP : Sigmoid model

$$\begin{cases} S_C[P_t] - \mathbb{E}[C_t] = (S_C[P_t] - C_{t-h})e^{-\nu h} \\ S_C[P_t] = \alpha_{\max} \text{Sigmoid}[\beta P_t + \beta_0] \end{cases} \quad (\mathcal{M}_S) \quad h \in \mathbb{R}^+$$

With

- $\text{Sigmoid}[x] = \frac{1}{1+e^{-x}} - \frac{1}{2}$
- Odd
- Bounded $[-1,1]$



Assumption : The stationary stat of Child node^x is a **sigmoid function** of parent node(s)

[Demo \(online\)](#)

III. Inference

III.1. Input – Steatosis (real)

Real public domain databases (available before my thesis)

- ToxCast
- TG-Gate

DEMO : Steatosis real data (use [online DEMO link](#))

Problems :

The experiments are not designed for qAOP modelling.

- Very few number of data : 3 endpoints
- Discretization assumption may fail because the measurements of the system state taken at intervals that are regularly spaced with a predetermined time granularity Δt
- observations on time $t = 2h$ $8h$ and $24h$

Input – Steatosis (simulated)

Data simulated from pharmacokinetics models based on ODE

Three virtual experimentation conditions:

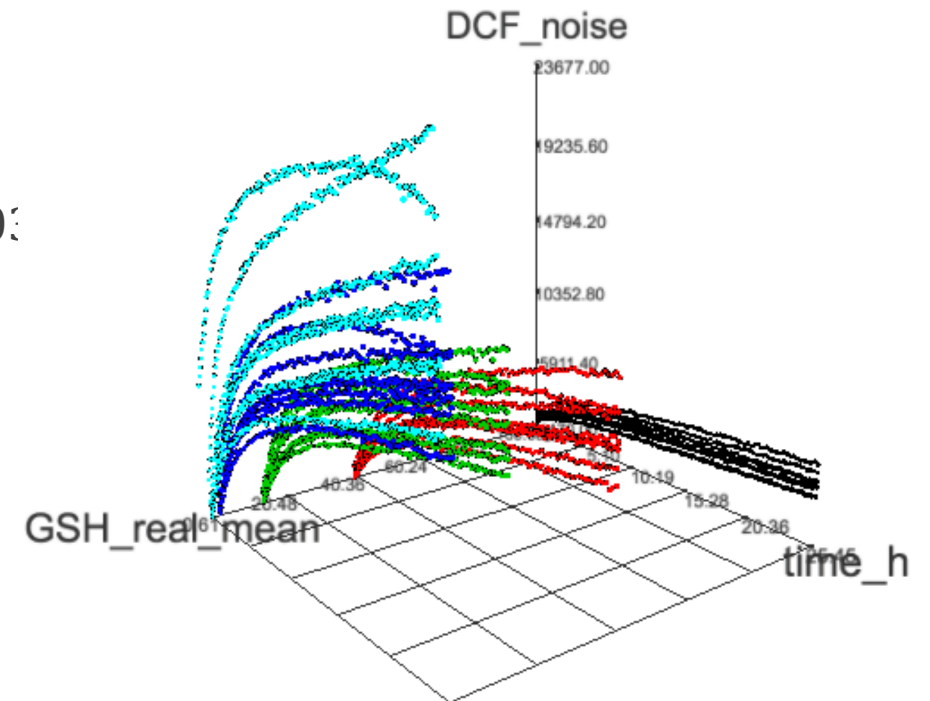
- One injection of chemical X , without decreasing feature $\forall i \in I, C_X^{(t_i)} = C_X^{(0)}$
- One injection of chemical X , with decreasing feature. $\forall i \in I, C_X^{(t_i)} < C_X^{(t_{i-1})}$
- Four injection of chemical X , with decreasing feature : $\frac{C_X^{(0)}}{4}$ each time

[online DEMO : Steatosis Generated data](#)

Input – Kidney disease (real)

Data of kidney disease from Eu-ToxRisk21

- 5 doses
- 8 replicates
- 103 observations endpoints
 - measured on time $t_i = 1 + 15 \times (i - 1)$ minutes $\forall i \in 1:103$



III.2. Parameters learning : linear model

$$\begin{cases} S_C[P_t] - \mathbb{E}[C_t] = (S_C[P_t] - C_{t-h})e^{-\nu h} \\ S_C[P_t] = \beta P_t + \beta_0 \\ h \in \mathbb{R}^+ \end{cases} \quad (\mathcal{M}_L)$$

Parameter estimation based on observations

Frequentist approach

$$\hat{\theta}_{ML} = \underset{\theta \in \Theta}{\operatorname{argmax}} L(\theta | \mathcal{D}) = \underset{\theta \in \Theta}{\operatorname{argmax}} P(\mathcal{D} | \theta)$$

Bayesian approach

$$\hat{\theta}_{MAP} = \underset{\theta \in \Theta}{\operatorname{argmax}} P(\mathcal{D} | \theta) \pi(\theta)$$

- Algorithm MCMC under the probabilistic programming language “**stan**”

III.3. Implementation: Simulated data

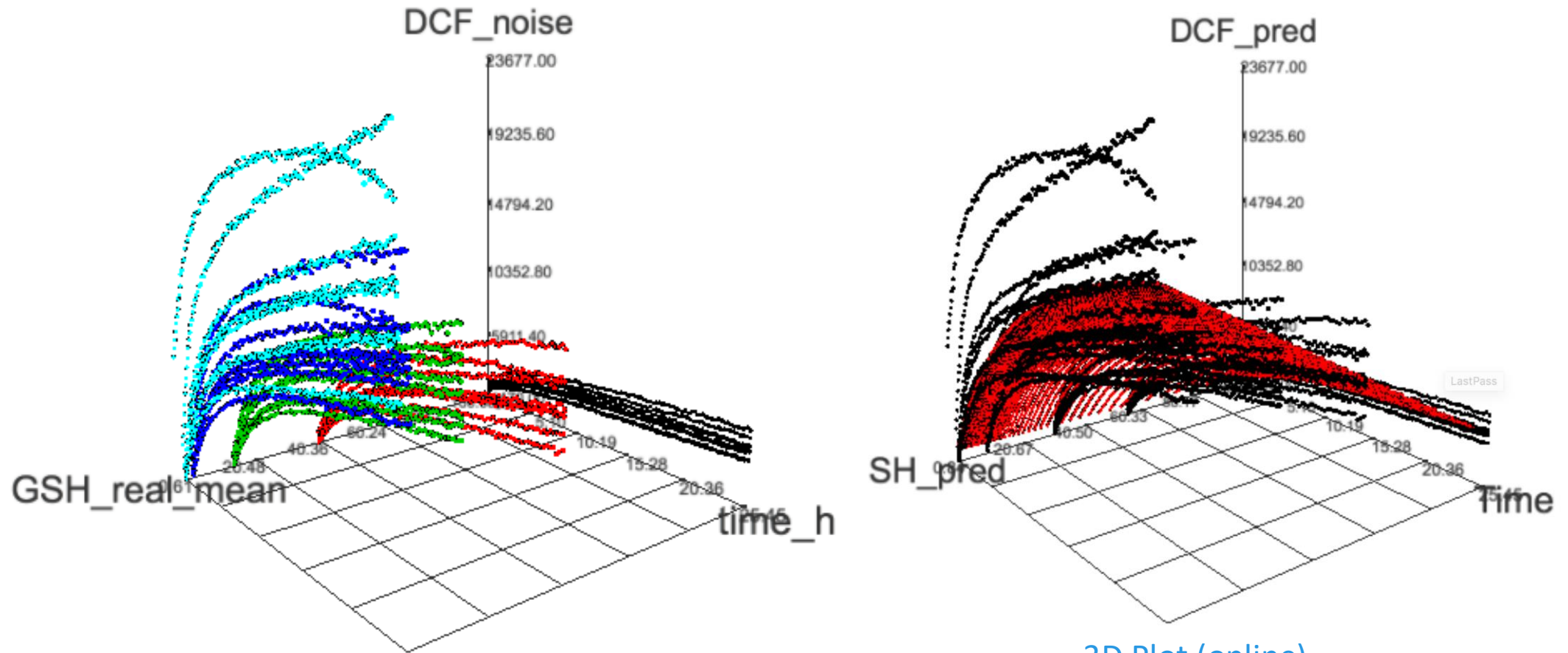
Parameter learning for a local conditional model :

$$P(FAS | LXR, t)$$

Result :

- ❖ Sigmoid model works significantly better with small set of data
- ❖ The linear model works as well as Sigmoid with large set of data

Implementation : Kidney disease (real)



[3D Plot \(online\)](#)

IV. Conclusion

Conclusion

- ❖ Proposition of qAOP model family: Linear model, Sigmoid model
- ❖ Application on Steatosis (real), steatosis (simulated), kidney disease (real).
 - ❖ Steatosis (real): Not fit well, not enough data.
 - ❖ Steatosis (simulated):
 - ❖ The sigmoid model fits better when only few data are available
 - ❖ The Linear model fits as well as the sigmoid model when more data are available
 - ❖ Kidney disease (real):
 - ❖ Linear models can well fit the database.

Future steps:

- ❖ Test sigmoid model on Kidney disease data
- ❖ Test model performance (behaviour) on simulated data with more nodes in AOP
- ❖ qAOP network problems: hierarchical DBN.

Reference

Boyen, X. and Koller R, D., Tractable Inference for Complex Stochastic Processes, In UAI-98, 1998.

Murphy, K. P., Dynamic Bayesian Networks: Representation, Inference and Learning, PhD thesis, UC Berkeley, Computer Science Division, July 2002.

Koller R, D., Friedman, K., Probabilistic Graphical Models: Principles and Techniques - Adaptive Computation and Machine Learning, The MIT Press, 2009

Friedman, N., Murphy, K., and Russel, S., Learning the structure of dynamic probabilistic networks, In 12th UAI, 1998.

Kalman, R. E., A New Approach to Linear Filtering and Prediction Problems, Transactions of the ASME--Journal of Basic Engineering, 1960.

Thank you for your attention
