

# Operator-valued Kernel-based models for Gene Regulatory Network Inference

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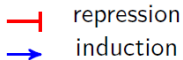
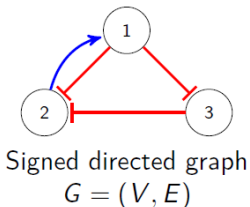
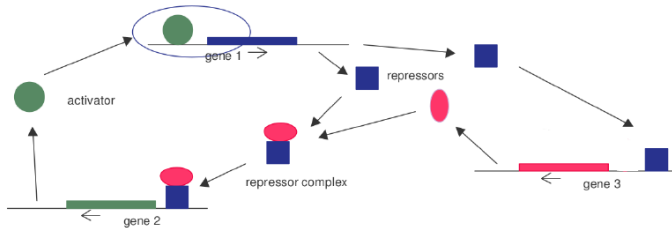
# Outline

- 1 Reverse-engineering GRN
- 2 A new framework for network inference
- 3 A novel nonlinear vector autoregressive model
- 4 Learning the OKVAR model
- 5 Numerical studies
- 6 Conclusion

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# Example: Gene Regulatory Network

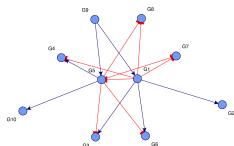
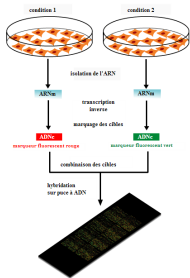


$$\begin{pmatrix}
 0 & 1 & 0 \\
 -1 & 0 & -1 \\
 -1 & 0 & 0
 \end{pmatrix}$$

Adjacency matrix

# Reverse-engineering GRN

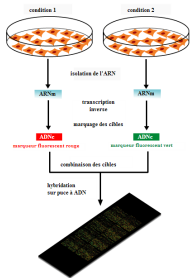
Reconstruction of gene regulatory networks = Identify **direct** interactions from gene expression data



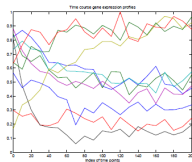
Signed directed graph  
(*E. coli* subnetwork)

# Reverse-engineering GRN

Reconstruction of gene regulatory networks = Identify **direct** interactions from gene expression data (**time-series**)

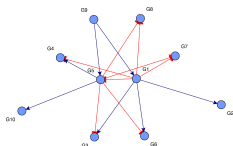


Processing



Time-course gene expression profiles

Reverse-modeling



Signed directed graph  
(*E. coli* subnetwork)

# Dynamical models and Network inference : State of the art

- Correlations, Mutual Information [Butte *et al.*, 2000; Basso *et al.*, 2005; Faith *et al.*, 2007]

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- **Linear models:**
  - ▶ Linear regression [D'Haeseleer, 1999], LASSO [van Someren *et al.*, 2006], linear autoregressive models [Fujita *et al.*, 2007; Shimamura *et al.*, 2009], several-order autoregressive models [Lozano, 2009; Bolstad *et al.*, 2011], Gaussian graphical models [Schäfer & Strimmer, 2005; Charbonnier *et al.*, 2010]
  - ▶ Granger causality [Shojaie and Michailidis, 2010 and 2011]
  - ▶ State-space models [Perrin *et al.*, 2003; Rangel *et al.*, 2004]



# Dynamical models and Network inference : State of the art

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  - ▶ Granger causality [Shojaie and Michailidis, 2010 and 2011]
  - ▶ State-space models [Perrin *et al.*, 2003; Rangel *et al.*, 2004]
- **Nonlinear models:**
  - ▶ Boolean logic [Liang *et al.*, 1998]
  - ▶ Ordinary Differential Equations (ODEs) [Chen *et al.*, 1999]
  - ▶ Bayesian networks [Murphy & Mian, 1999; Friedman *et al.*, 2000; Perrin *et al.*, 2003; Auliac *et al.*, 2008]
  - ▶ Random forests [Huynh-Thu *et al.*, 2010]
  - ▶ Non-parametric Gaussian processes [Äijö & Lähdesmäki, 2009]
  - ▶ Kernels and time-series [Ralaivola *et al.*, 2005; Principe *et al.*, 2011; Kallas *et al.*, 2011]

# Dynamical models and Network inference

## Limitations

- Specific
- Undirected graph
- Linear
- Small systems

# Dynamical models and Network inference

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- Specific
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## Requirements

- Generic
- Causality
- Nonlinear
- Scalable

## Our approach

- 1 Introduce a general framework for nonlinear multivariate modeling and network inference
- 2 Extend the framework of linear modeling to sparse nonlinear modeling

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# Modeling nonlinear dynamical systems

## Model assumptions

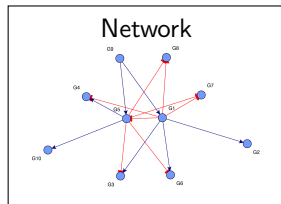
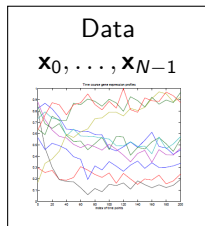
The temporal evolution of the system is ruled by a **first-order stationary nonlinear** model  $h : \mathbb{R}^d \rightarrow \mathbb{R}^d$ :

$$\mathbf{x}_{t+1} = h(\mathbf{x}_t) + \mathbf{u}_t \quad (1)$$

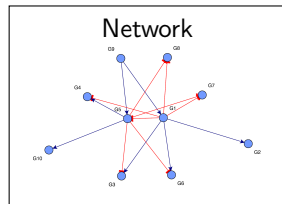
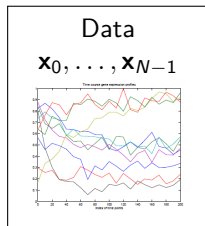
where

- $\mathbf{x}_0, \dots, \mathbf{x}_{N-1} \in \mathbb{R}^d$  : observed time series of a dynamical system comprising of  $d$  variables at time  $t = 0, \dots, N - 1$
- $\mathbf{u}_t$  is a noise term

# Network inference chart

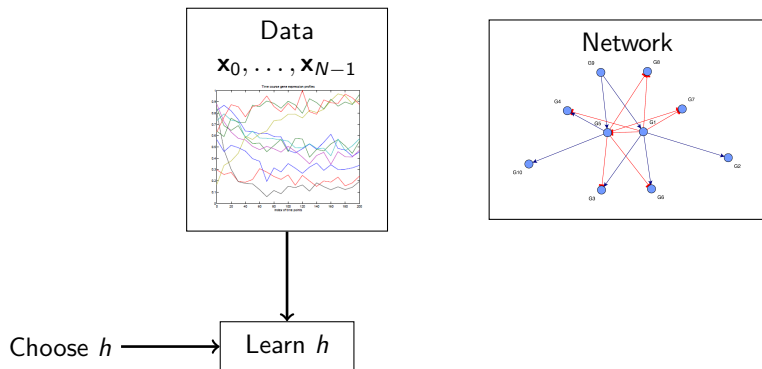


# Network inference chart



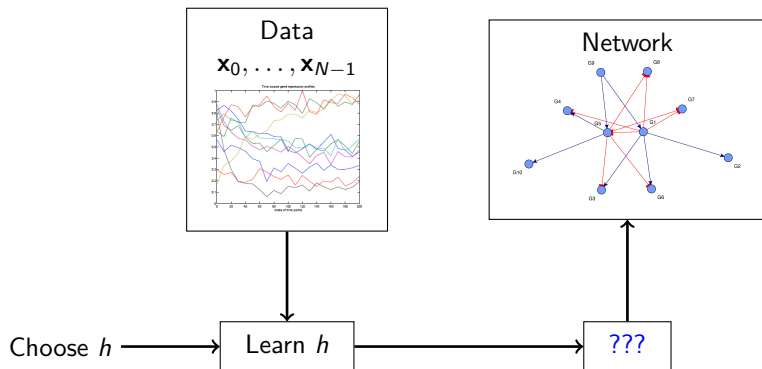
Choose  $h$

# Network inference chart





# Network inference chart



## Network inference with linear models

- 1  $\mathbf{x}_{t+1} = B\mathbf{x}_t + \mathbf{u}_t$  : learn  $B$  with a sparsity constraint
- 2 Threshold  $B$  to get an estimate of the adjacency matrix  $A$

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## Network inference with nonlinear models

- 1 Learn a nonlinear model  $h : \mathbb{R}^d \rightarrow \mathbb{R}^d$  :  $\mathbf{x}_{t+1} = h(\mathbf{x}_t) + \mathbf{u}_t$

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## Network inference with nonlinear models

- 1 Learn a nonlinear model  $h : \mathbb{R}^d \rightarrow \mathbb{R}^d : \mathbf{x}_{t+1} = h(\mathbf{x}_t) + \mathbf{u}_t$
- 2 Compute the average empirical **Jacobian** matrix of  $h$  :

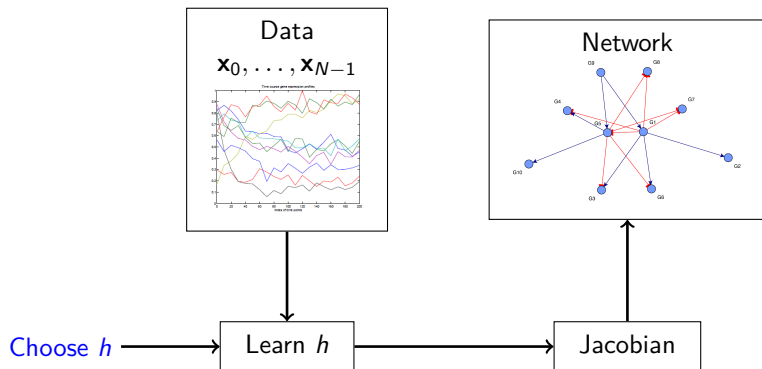
$$J(h)_{ij} = \frac{1}{N-1} \sum_{t=0}^{N-2} \frac{\partial h(\mathbf{x}_t)_i}{\partial (\mathbf{x}_t)_j}$$

- 3 Threshold  $J(h)$  to get an estimate of the adjacency matrix  $A$

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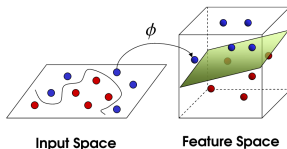


*From single-valued functions ...*

- Binary classification
- Real-valued regression

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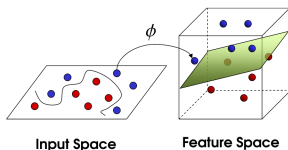
- Binary classification SVM
- Real-valued regression SVR
- kernels : popular nonparametric nonlinear methods





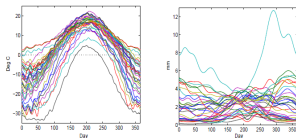
From single-valued functions ...

- Binary classification SVM
- Real-valued regression SVR
- **Scalar kernels** : popular nonparametric nonlinear methods



... to vector-valued functions

Recent interest in **operator-valued kernels**  
[Senkane and Tempel'man, 1973; Michelli & Pontil, 2005; Caponnetto *et al.*, 2008]



Development of new learning tasks:

- Multi-task learning [Evgeniou *et al.*, 2005]
- Functional regression [Kadri *et al.*, 2010]
- Structured output prediction [Brouard *et al.*, 2011]

# RKHS theory for vector-valued functions

## Notations

- Input set :  $\mathcal{X}$
- Output Hilbert space :  $\mathcal{F}_y$
- We consider functions  $h : \mathcal{X} \rightarrow \mathcal{F}_y$

# RKHS theory for vector-valued functions

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Operator-valued kernel [Senkene and Tempel'man (1973) ,  
Caponnetto *et al* (2008)]

$K : \mathcal{X} \times \mathcal{X} \rightarrow L(\mathcal{F}_y)$  is an **operator-valued kernel** if:

- $\forall (x, z) \in \mathcal{X} \times \mathcal{X}, K(x, z) = K(z, x)^*$
- $\forall m \in \mathbb{N}, \forall \{(x_i, \mathbf{y}_i)\}_{i=1}^m \subseteq \mathcal{X} \times \mathcal{F}_y, \sum_{i,j=1}^m \langle \mathbf{y}_i, K(x_i, x_j) \mathbf{y}_j \rangle_{\mathcal{F}_y} \geq 0$

## Representer theorem [Michelli and Pontil (2005)]

Let  $\lambda > 0$ ,  $\mathcal{S}_n = \{(x_1, \mathbf{y}_1), \dots, (x_n, \mathbf{y}_n)\} \subset \mathbb{R}^d \times \mathbb{R}^d$ . Then, the following optimization problem:

$$\operatorname{argmin}_{h \in \mathcal{H}} \mathcal{L}(h) = \sum_{i=1}^n \|h(x_i) - \mathbf{y}_i\|^2 + \lambda \|h\|_{\mathcal{H}}^2$$

admits a solution of the form:

$$\hat{h}(\cdot; \mathcal{S}_n) = \sum_{\ell=1}^n K(x_{\ell}, \cdot) \mathbf{c}_{\ell} \quad (2)$$

where  $\mathbf{c}_{\ell} \in \mathbb{R}^d$ ,  $\ell = \{1, \dots, n\}$  are to be learned

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## Operator-valued Kernel-based Vector AutoRegressive (OKVAR) model

Given the observed time series  $\mathcal{S}_N = \{(\mathbf{x}_0, \mathbf{x}_1), \dots, (\mathbf{x}_{N-2}, \mathbf{x}_{N-1})\} \subset \mathbb{R}^d \times \mathbb{R}^d$ , the OKVAR model  $h$  is defined as

$$h(\mathbf{x}_t; \mathcal{S}_N) = \sum_{\ell=0}^{N-2} K(\mathbf{x}_{\ell}, \mathbf{x}_t) \mathbf{c}_{\ell} \quad (3)$$

# The OKVAR model family

## Examples of matrix-valued kernels

- 1  $K_1(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z})B$  with  $k_1(\mathbf{x}, \mathbf{z}) = \exp(-\gamma_1 \|\mathbf{x} - \mathbf{z}\|^2)$  and  $B \in S_d^+(\mathbb{R})$
- 2  $\forall (p, q) \in \{1, \dots, d\}^2, K_2(\mathbf{x}, \mathbf{z})_{pq} = \exp(-\gamma_2 (x^p - z^q)^2)$
- 3  $K(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z})B \circ K_2(\mathbf{x}, \mathbf{z})$

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$$J(h_1)_{ij}(t) = \sum_{\ell=0}^{N-2} \sum_{q=1}^d b_{iq} c_\ell^q \frac{\partial k_1(\mathbf{x}_t, \mathbf{x}_\ell)}{\partial x_t^j}$$

- ②  $\forall (p, q) \in \{1, \dots, d\}^2, K_2(\mathbf{x}, \mathbf{z})_{pq} = \exp(-\gamma_2(x^p - z^q)^2)$

$$J(h_2)_{ij}(t) = 2\gamma_2(x_t^i - x_t^j) \exp\left(-\gamma_2(x_t^i - x_t^j)^2\right) c_t^j$$

- ③  $K(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z})B \circ K_2(\mathbf{x}, \mathbf{z})$

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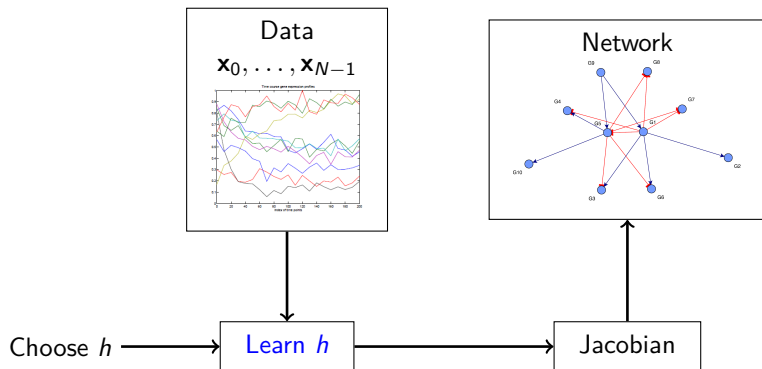
$$\begin{aligned} J_{ij}(t) &= 2\gamma_2 b_{ij} (x_t^i - x_t^j) \exp(-\gamma_2(x_t^i - x_t^j)^2) c_t^j \\ &\quad - 2\gamma_1 \sum_{\ell \neq t} k_1(\mathbf{x}_t, \mathbf{x}_\ell) (x_t^j - x_\ell^j) \sum_{p=1}^d b_{ip} \exp(-\gamma_2(x_t^i - x_\ell^p)^2) c_\ell^p \end{aligned}$$



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# Network inference chart



# Learning the OKVAR model

We aim to solve the following optimization problem :

$$\begin{aligned} & \underset{B \in M_d(\mathbb{R}), C \in M_{N-1,d}(\mathbb{R})}{\text{minimize}} & \mathcal{L}(B, C) &= \sum_{t=0}^{N-2} \|h(\mathbf{x}_t; B, C) - \mathbf{x}_{t+1}\|^2 + \Omega(B, C) \\ & \text{s.t.} & & B \in \mathcal{S}_d^+(\mathbb{R}) \end{aligned} \tag{4}$$

with  $\Omega(B, C) = \lambda_h \|h_{B,C}\|_{\mathcal{H}}^2 + \lambda_C \|C\|_{\ell_1} + \lambda_B \|B\|_{\ell_1}$

# Learning the OKVAR model

- For fixed  $B$  and for  $\mathbf{c}_\ell$  the loss function to be minimized becomes:

$$\mathcal{L}(\hat{B}, \mathbf{C}, \ell) = \sum_{t=0}^{N-2} \|h(\mathbf{x}_t; \hat{B}, \mathbf{C}) - \mathbf{x}_{t+1}\|^2 + \lambda_h \|h_{\hat{B}, \mathbf{C}}\|_{\mathcal{H}}^2 + \lambda_C \|\mathbf{C}\|_{\ell_1} \quad (5)$$

- For given  $\hat{C}$ , the loss function to be minimized is the following:

$$\mathcal{L}(B, \hat{C}) = \sum_{t=0}^{N-2} \|h(\mathbf{x}_t; B, \hat{C}) - \mathbf{x}_{t+1}\|^2 + \lambda_h \|h_{B, \hat{C}}\|_{\mathcal{H}}^2 + \lambda_B \|B\|_{\ell_1} \quad (6)$$

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- ▶ proximal gradient algorithms [Martinet (1970); Beck and Teboulle (2010)]
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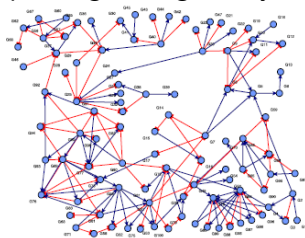
- ▶ Matrix exponentiated gradient updates [Tsuda *et al* (2005)]

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# DREAM3 data set

- DREAM = Dialogue for Reverse Engineering Assessments and Methods
- 5 size-10 and 5 size-100 networks (subgraphs of *E. coli* and *S. cerevisiae*) have been generated
  - ▶ *Ecoli1*, *Ecoli2*, *Yeast1*, *Yeast2*, *Yeast3*
- An example of gene regulatory network : *S. cerevisiae* subnetwork



→ activation  
→ inhibition

- Challenge : Reconstruct the networks from time-series data of  $N = 21$  points

# DREAM3 size-10 data sets : Results

Table 1: AUROC

	Ecoli1	Ecoli2	Yeast1	Yeast2	Yeast3
OKVAR + <i>True B</i>	0.956	0.918	0.806	0.781	0.780
<b>OKVAR</b>	<b>0.717</b>	<b>0.724</b>	0.644	<b>0.740</b>	<b>0.705</b>
LASSO	0.500	0.547	0.528	0.627	0.582
GPODE	0.607	0.516	0.494	0.613	0.571
G1DBN	0.604	0.573	0.494	0.540	0.601
Team 236	0.621	0.650	<b>0.646</b>	0.438	0.488
Team 190	0.573	0.515	0.631	0.577	0.603

Table 2: AUPR

	Ecoli1	Ecoli2	Yeast1	Yeast2	Yeast3
OKVAR + <i>True B</i>	0.752	0.677	0.473	0.523	0.586
<b>OKVAR</b>	<b>0.385</b>	<b>0.678</b>	<b>0.430</b>	<b>0.480</b>	<b>0.447</b>
LASSO	0.119	0.531	0.244	0.305	0.255
GPODE	0.180	0.146	0.089	0.377	0.341
G1DBN	0.159	0.534	0.192	0.226	0.248
Team 236	0.197	0.378	0.194	0.236	0.239
Team 190	0.152	0.181	0.167	0.371	0.373



# DREAM3 size-100 data sets : Results

**Table 1: AUROC**

	Ecoli1	Ecoli2	Yeast1	Yeast2	Yeast3
OKVAR + <i>True B</i>	0.962	0.971	0.958	0.906	0.897
<b>OKVAR</b>	<b>0.618</b>	<b>0.620</b>	<b>0.537</b>	<b>0.553</b>	<b>0.522</b>
LASSO	0.519	0.512	0.507	0.530	0.506
G1DBN	0.553	0.548	0.510	0.509	0.506
Team 236	0.527	0.546	0.532	0.508	0.508

**Table 2: AUPR**

	Ecoli1	Ecoli2	Yeast1	Yeast2	Yeast3
OKVAR + <i>True B</i>	0.432	0.516	0.279	0.407	0.364
<b>OKVAR</b>	<b>0.029</b>	<b>0.093</b>	0.024	<b>0.052</b>	0.053
LASSO	0.016	0.057	0.016	0.044	0.044
G1DBN	0.018	0.052	0.022	0.043	0.049
Team 236	0.019	0.042	<b>0.035</b>	0.046	<b>0.065</b>

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- **A key problem:** GRN inference from multivariate time-series data
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    - ★ Network inference method via the [Jacobian](#)
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  - ▶ Causality 
    - ★ Network inference method via the [Jacobian](#)
  - ▶ Nonlinear 
    - ★ A novel [operator-valued kernel](#) based vector autoregressive model
  - ▶ Scalable

# Conclusion

- **A key problem:** GRN inference from multivariate time-series data
- **Requirements:**
  - ▶ Generic
  - ▶ Causality 
    - ★ Network inference method via the [Jacobian](#)
  - ▶ Nonlinear 
    - ★ A novel [operator-valued kernel](#) based vector autoregressive model
  - ▶ Scalable 
    - ★ Learning the OKVAR model's parameters  $\sim$  minutes for size-100 data sets
- **Results:**
  - ▶ Very good performance of the OKVAR model on simulated benchmark data sets

- **Theoretical results:**

- ▶ Universality of kernels, consistency of the Jacobian estimator [Fouchet]
- ▶ Generalization error



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- **Theoretical results:**
  - ▶ Universality of kernels, consistency of the Jacobian estimator [Fouchet]
  - ▶ Generalization error
- Probabilistic framework with informative priors
- Exploit the OKVAR model for prediction

# List of recent papers

- N. Lim\*, Y. Senbabaoglu\*, G. Michailidis, F. d'Alché-Buc

**BIOINFORMATICS ORIGINAL PAPER**

2013, pages 1–8  
doi:10.1093/bioinformatics/btt167

*System biology*

Advance Access publication April 10, 2013

**OKVAR-Boost: a novel boosting algorithm to infer nonlinear dynamics and interactions in gene regulatory networks**

- N. Lim\*, F. d'Alché-Buc\*, C. Auliac, G. Michailidis, Operator-valued Kernel based Vector Autoregressive Models for Network Inference, submitted to *Machine Learning Journal* (under revision)

**Thank you for your attention !**