# The European Commission's science and knowledge service

Joint Research Centre



Introduction to metamodelling/emulation

M. Ratto DYNARE Summer School Paris, June 14th 2018



## **Introduction to ANOVA Models**

Consider the computational model

$$Y = f(X)$$

where x is a vector of input 'factors'  $X = (X_1, X_2, ..., X_k)$ 

$$X = (X_1, X_2, \dots, X_k)$$

and Y is the output.

 $X_i$  is the i-th element of x varying in  $0 \le X_i \le 1$ 

$$X \in \Omega \longrightarrow f(\mathbf{x}) \longrightarrow Y$$





Realistic Computer models are very costly in terms of run time. We need a way to characterize uncertainty and perform SA based on a limited number of model runs.

We aim to identify a 'simple' relationship between Xi's and Y that fits well the original model and is less computationally demanding.

A Meta-model is a simpler model that mimics the larger computational model.

Evaluations of Meta-Models are much faster.



Local approximation methods take f and its derivatives at a base point  $X_0$  and construct a function that matches the properties of f in the nearby region (Taylor series).



- 1. Linear Regression
- 2. Quadratic Response Surface Regression
- Options 1 and 2 work pretty well in some cases. However many realistic models are highly nonlinear and/or periodic and these methods fail.
- 3. Gaussian Process or Spatial Models
- 4. Nonparametric Regression Models
- Both 3 and 4 work fairly well for a small number of inputs.
- Variable selection is helpful for a modest number of inputs.

### **ANOVA Models**

ANOVA models define a decomposition of Y=f(x) into main effects and interactions

$$Y = f(x) = f_0 + \sum_{i=1}^{k} f_i(x_i) + \sum_{i} \sum_{j>i} f_{ij}(x_i, x_j) + \dots + f_{1,2,\dots,k}(x_1, x_2, \dots, x_k)$$

This is also called the High Dimensional Model Representation (HDMR)

E.g., if k=3: 
$$f(x) = f_0 + f_1(x_1) + f_2(x_2) + f_3(x_3) + f_{12}(x_1, x_2) + f_{13}(x_1, x_3) + f_{23}(x_2, x_3) + f_{123}(x_1, x_2, x_3)$$

This decomposition is non-unique for general joint pdf's of x.

The total number of summands in the ANOVA is  $2^k$ 



# Properties of the ANOVA decomposition (orthogonal case)

$$Y = f(x) = f_0 + \sum_{i=1}^{k} f_i(x_i) + \sum_{i} \sum_{j>i} f_{ij}(x_i, x_j) + \dots + f_{1,2,\dots,k}(x_1, x_2, \dots, x_k)$$

If each term is chosen with zero mean ...

$$\int_{0}^{1} f_{i}(x_{i}) dp(x_{i}) = 0, \quad \forall x_{i} \quad i = 1, 2, ..., k$$

$$\int_{0}^{1} \int_{0}^{1} f_{ij}(x_{i}, x_{j}) dp(x_{i}) dp(x_{j}) = 0, \quad \forall x_{i}, x_{j} \quad i < j$$
....
$$\int_{0}^{1} f_{12...k}(x_{1}, x_{2}, ..., x_{k}) dp(x_{1}) dp(x_{2}) ... dp(x_{k}) = 0.$$



# Properties of the ANOVA decomposition (orthogonal case)

$$Y = f(x) = f_0 + \sum_{i=1}^{k} f_i(x_i) + \sum_{i} \sum_{j>i} f_{ij}(x_i, x_j) + \dots + f_{1,2,\dots,k}(x_1, x_2, \dots, x_k)$$

... then the ANOVA decomposition has TWO properties

$$\int_{\Omega} f(x) dp(x) = f_0 = E(Y)$$

All the summands are orthogonal:

if 
$$(i_1,...,i_s) \neq (j_1,...,j_l)$$
  $\int_{\Omega^k} f_{i_1,...,i_s} f_{j_1,...,j_l} dp(x) = 0$ 

It follows that the ANOVA decomposition is **unique** and each term can be defined as ...

## **Properties of the ANOVA decomposition**

$$Y = f(X) = E(Y) + \sum_{i=1}^{k} f_i(X_i) + \sum_{i} \sum_{j>i} f_{ij}(X_i, X_j) + \dots + f_{1,2,\dots,k}(X_1, X_2, \dots, X_k)$$

$$\int_{\Omega-x_i} f(x) dp(x \mid x_i) = f_0 + f_i(x_i) = E(Y \mid x_i)$$

$$\int_{\Omega - \{x_i, x_j\}} f(x) dp(x \mid x_i x_j) = f_0 + f_i(x_i) + f_j(x_j) + f_{ij}(x_i, x_j)$$

$$= E(Y \mid x_i, x_j)$$



### **ANOVA**

$$Y - E(Y) = \sum_{i=1}^{k} f_i(x_i) + \sum_{i} \sum_{j>i} f_{ij}(x_i, x_j) + \dots + f_{1,2,\dots,k}(x_1, x_2, \dots, x_k)$$

Being the terms orthogonal, we can square and integrate the eq above over  $\Omega$  and decompose the variance of f(x) into terms of increasing dimensionality (ANOVA)

$$V(Y) = \sum_{i=1}^{k} V_i + \sum_{i} \sum_{j} V_{ij} + \sum_{i} \sum_{j} \sum_{k} V_{ijk} + \cdots + V_{1,2,\dots,k}$$

$$V_{i} = \int f_{i}^{2}(x_{i})dx_{i}$$

$$V_{i_{1},i_{2},...i_{s}} = \int f_{i_{1},...,i_{s}}^{2}dx_{i_{1}}dx_{i_{2}}...dx_{i_{s}}$$

$$1 = \sum_{i=1}^{k} S_i + \sum_{i} \sum_{j} S_{ij} + \sum_{i} \sum_{j} \sum_{k} S_{ijk} \dots + S_{1,2,\dots,k}$$



If we were to approximate f(x) with a function  $g(X_i)$  ...

$$L = E[(f(x) - g(x_i))^2]$$

$$g(x_i) = E(Y \mid x_i) \Rightarrow L = L_{\min}$$
$$L_{\min} = E[Var(Y \mid X_i)]$$

...  $f_i=E(Y|X_i)$  has the minimum loss L among univariate functions

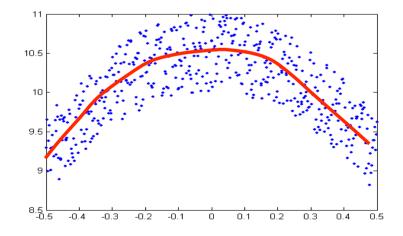
$$V_i = Var[E(Y \mid X_i)]$$
  $L = E[(f(x) - g(x))^2]$   
=  $Var(Y) - E[Var(Y \mid X_i)]$   
 $g(x) = f_0 \Rightarrow L = V(Y)$ 

when we approximate f(x) with a function  $g(x_i)$ ,  $g^*=E(Y|X_i)$  has the minimum loss L

$$g(x) = f(x) \Rightarrow L = 0$$

$$g(x_i) = E(Y \mid x_i) \Rightarrow L = L_{\min}$$

$$L_{\min} = E[Var(Y \mid X_i)]$$



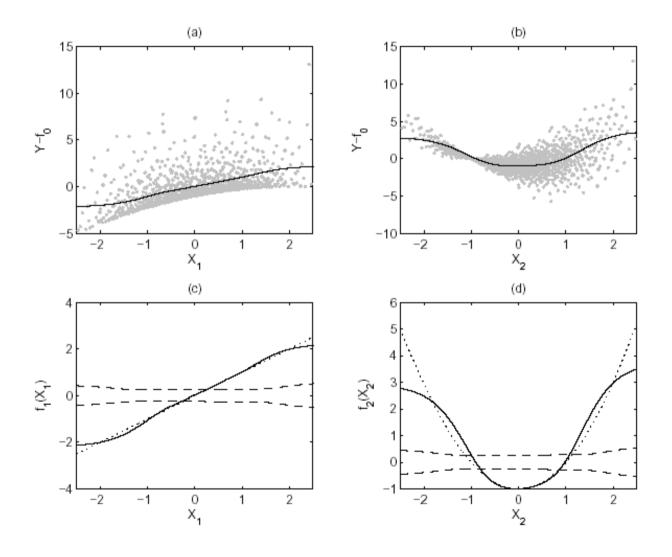


...  $f_{ij}=E(Y|X_i,X_j)$  has the minimum loss L among bivariate functions ...

... and so on ...

Var(fi)/var(Y): also called non-parameteric R-squared; Pearson's correlation ratio; ...





$$f(X_1, X_2) = X_1 + X_2^2 + X_1 \cdot X_2$$

$$X_i \sim N(0,1)$$



Tensor product decomposition (reproducing kernel Hilbert space, RKHS)

$$F = \bigotimes_{j=1}^{k} H_{j} = \{1\} \oplus \left\{ \bigoplus_{j=1}^{k} \overline{H}_{j} \right\} \oplus \left\{ \bigoplus_{j< i} (\overline{H}_{j} \otimes \overline{H}_{i}) \right\} \oplus L$$

Orthogonal functional decomposition

$$F = \{1\} \oplus \left\{ \bigoplus_{j=1}^{q} F_j \right\}$$



Smoothing methods to estimate ANOVA decompositions, truncated at the 2<sup>nd</sup> -3<sup>rd</sup> order terms:

SMOOTHING SPLINES ANOVA MODELS

Smoothing the y vs x mapping (think of an HP-filter), that provides efficient convergence properties to the true ANOVA decomposition.

[this is one possible methods, other are RBF's, kernel regressions, ...]



Denote the generic mapping as  $Y = f(\mathbf{X})$ , where  $\mathbf{X} \in [0, 1]^k$  and k is the number of parameters.

The simplest example of smoothing spline mapping estimation of z is the additive model:

$$g(\mathbf{X}) = g_0 + \sum_{i=1}^k g_i(x_i)$$



To estimate g we can use a multivariate smoothing spline minimization problem, that is, given  $\lambda_j$ , find the minimizer  $g(\mathbf{X})$  of:

$$\frac{1}{N} \sum_{n=1}^{N} (y_n - g(\mathbf{X}_n))^2 + \sum_{j=1}^{k} \lambda_j \int_0^1 [g_j''(X_j)]^2 dX_j$$

where a Monte Carlo sample of dimension N is assumed.

This minimization problem requires the estimation of the k hyperparameters  $\lambda_j$  (also denoted as smoothing parameters): GCV, GML, etc. (see e.g. Wahba, 1990; Gu, 2002).

We re-formulate the additive model for the general case with interactions as to find the minimizer g(**X**) of:

$$\left| \frac{1}{N} \sum_{n=1}^{N} (y_n - g(\mathbf{X}_n))^2 + \lambda_0 \sum_{j=1}^{q} \frac{1}{\theta_j} \| P^j g \|_F^2 \right|$$

where the q-dimensional vector of  $\theta_j$  smoothing parameters needs to be optimized 'somehow'.



```
Wahba (1990), Gu (2002), Storlie et al. (2007): en-bloc (like HP);
```

Ratto et al., 2004-2007: based on recursive filtering and smoothing estimation: SDP modelling. (like KF-smoothing version of the HP);

Liu et al. (2002-2006): polynomial basis expansion.



# **SDP** modelling

SDP modeling is one class of non-parametric smoothing approach first suggested by Young (1993).

The estimation is performed with the help of the `classical' recursive (non-numerical) Kalman filter and associated fixed interval smoothing algorithms and has been applied for sensitivity analysis in Ratto et al. (2004-2007).



## Mapping/sensitivity strategies

### OAT (Taylor):

- •truncation;
- mapping by knowing all derivatives in a base point
- decomposition with infinite terms

## GSA (ANOVA):

- non-parametric regression/smoothing
- mapping on a spacefilling MC sample
- decomposition with finite terms



#### **DSGE** models

Let us consider a generic DSGE model:

$$E_t\{g(y_{t+1}, y_t, y_{t-1}, u_t; X)\} = 0$$

y<sub>t</sub> endogenous variables, u<sub>t</sub> exogenous shocks X structural parameters.

X can be characterised by plausible ranges, expressed in terms of prior distributions, or by a posterior distribution, as a result of an estimation.



#### **DSGE** models

The model behaviour is a function of the values assumed for X within the prior or posterior space of structural parameters.

Let Y be a generic 'output' of the model: a multiplier, a measure of fit, an IRF.

Y depends on the values of X

$$Y=f(X_1, ..., X_k),$$

f non-linear analytic form is unknown



## Mapping the reduced form of RE models

Relationship between the reduced form of a rational expectation model and the structural coefficients.

let the reduced form be  $y_t=Ty_{t-1}+Bu_t$ ,

'outputs' Y of our analysis will be the entries in the transition matrix  $T(X_1,...,X_k)$  or in the matrix  $B(X_1,...,X_k)$ .



We analyse the reduced form coefficients describing the relationship between

$$\pi_t$$
 vs  $e_{R,t}$ 

We sample the structural coefficients from posterior ranges obtained after estimating the model using data for Canada.



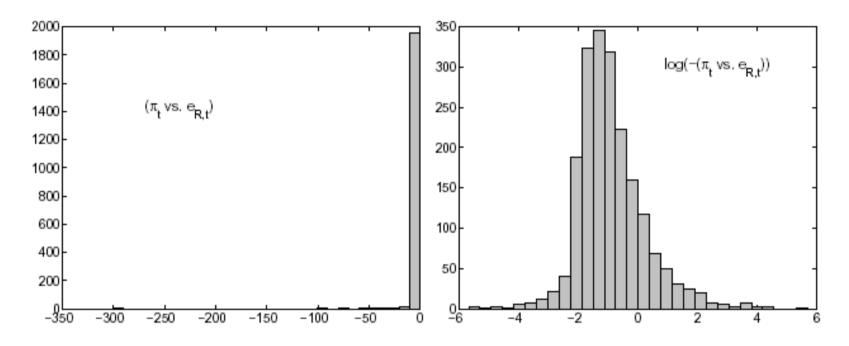


Figure 13: Lubik and Schorfheide model: histograms of the MC sample of the reduced form coefficient  $Y = (\pi_t \text{ vs } e_{R,t})$ . Left panel: actual values Y; right panel  $\log(-Y)$ .



$$\log(-(\pi_t \text{ vs } e_{R,t}))$$

$$Y = -\exp(f_0 + f_1 + ... + f_k + e)$$
  
= -\exp(e)\int\_{j=0}^k \exp(f\_j)

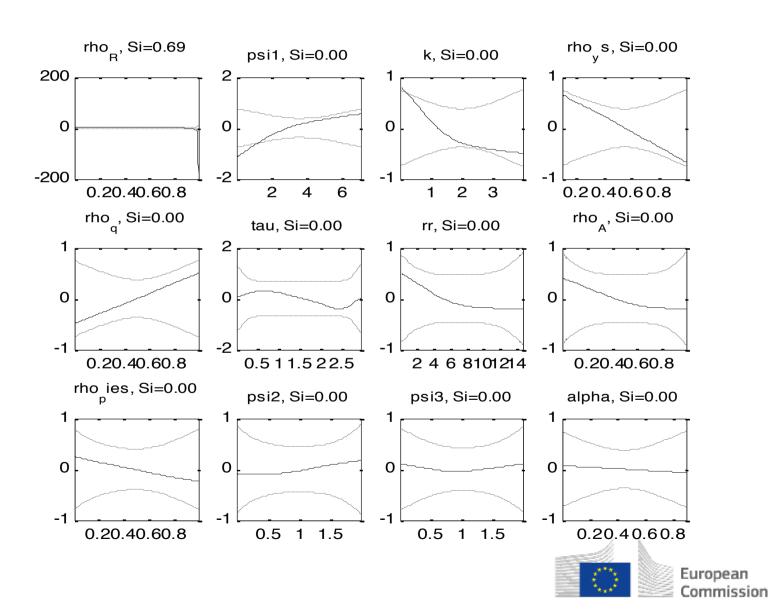
#### **Factorisation**

$$Y = -\exp(e) \prod_{j=0}^{k} \exp(f_j) + \sum_{j=0}^{k} g_j + \varepsilon$$

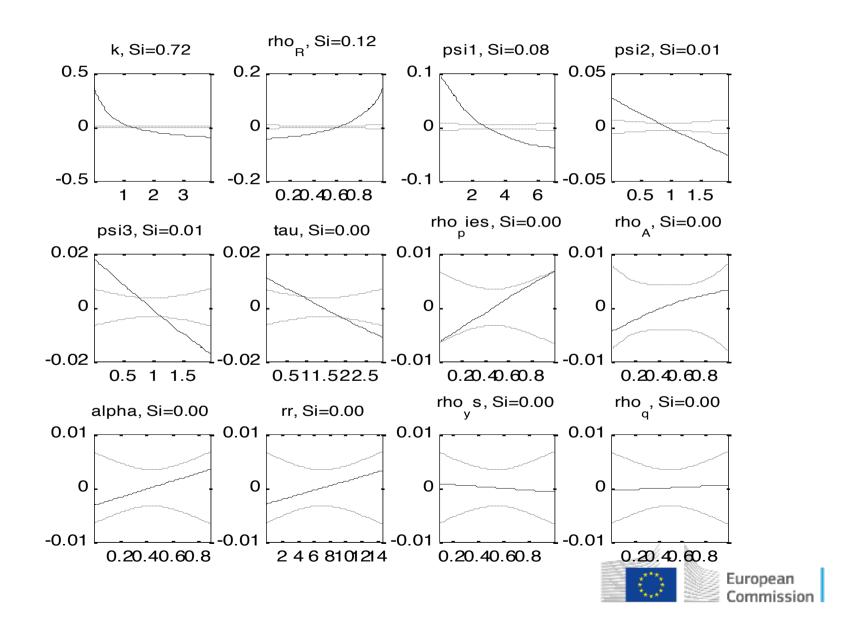
Correction [new in DYNARE 4.5]

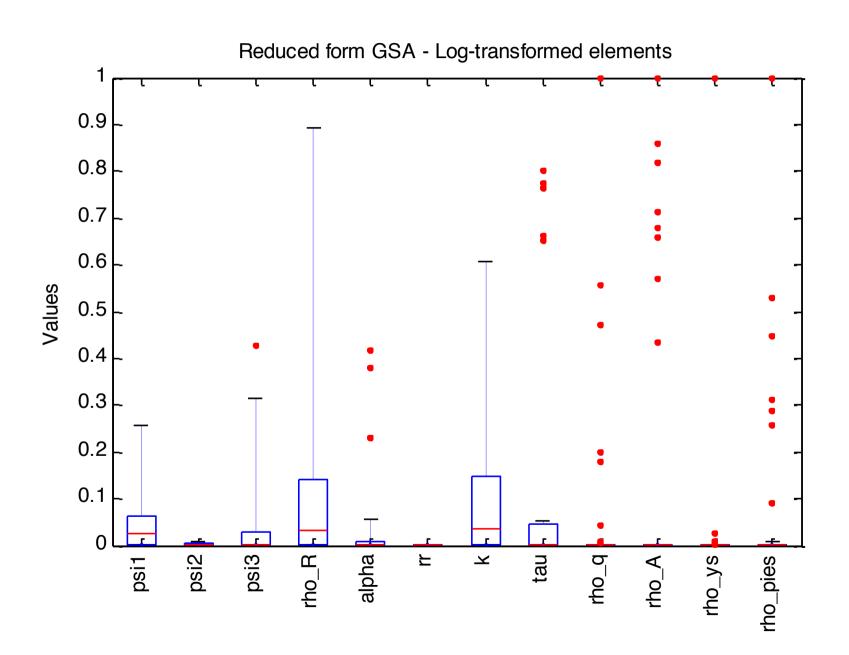


# **LS2005:** pie vs e\_R [-log(y)]



# LS2005: R vs e\_R [-log(y)]





# **Toolbox documentation**

The mapping of the reduced form soultion forces the use of samples from prior ranges or prior distributions, i.e.:

```
pprior=1 (default);
ppost=0 (default);
```

[unless neighbourhood\_width is applied]



# **Mapping reduced form solution**

option name	default	description	
redform	0	0 = don't prepare MC sample of	
		reduced form matrices	
		1 = prepare MC sample of	
		reduced form matrices	
load_redform	0	0 = estimate the mapping of	
		reduced form model	
		1 = load previously estimated mapping	
logtrans_redform	0	0 = use raw entries	
		1 = use log-transformed entries	
threshold_redform		= don't filter MC entries	
		of reduced form coefficients	
		[max max] = analyse filtered	
		entries within the range [max max]	
ksstat_redform	0.001	critical p-value for Smirnov statistics d	
		when threshold_redform is active	
		plot parameters with p-value <ksstat_redform< th=""><th></th></ksstat_redform<>	
alpha2_redform	0	critical p-value for correlation $\rho$	
		when threshold_redform is active	
		plot couples of parameters with	
		p-value <alpha2_redform< th=""><th></th></alpha2_redform<>	
namendo	()	list of endogenous variables	
	:	jolly character to indicate ALL endogenous	
namlagendo	()	list of lagged endogenous variables:	
		analyse entries [namendo×namlagendo]	
	:	jolly character to indicate ALL endogenous	
namexo	()	list of exogenous variables:	
		analyse entries [namendo×namexo]	pean
	:	jolly character to indicate ALL exogenous	ımission

# ss\_anova Toolbox

Download here the Recursive SS-ANOVA Toolbox for MATLAB from the web site of the summer school ss\_anova\_recurs.zip

