

HU-MATHS-IN miniproject proposal

Dimension Reduction of High Frequency and High Dimensional Data in Time and Space

Marianna Bolla, Máté Baranyi, Fatma Abdelkhalek

Department of Stochastics

Institute of Mathematics, BME

`marib@math.bme.hu`

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Background: Time Series and Random Fields

- ▶ Reduction of dimensionality in high frequency time series; for example, finding the underlying signals detected by many sensors: [M. Bolla, Factor Analysis, Dynamic. Wiley StatsRef: Statistics Reference Online \(2017\)](#) and book in preparation with T. Szabados.
- ▶ Graphical models and prediction along regression graphs (to develop artificial intelligence, e.g., for medical diagnosis): [M. Bolla, F. Abdelkhalek, M. Baranyi, Graphical models, regression graphs, and recursive linear regression in a unified way, Acta Univ. Szeged \(2019\)](#) (related SEM, PLS papers and Python program in preparation).
- ▶ Image segmentation with spectral clustering tools (tumor-segmentation, computer aided surgery): [M. Bolla, Spectral Clustering and Biclustering, Wiley \(2013\)](#).

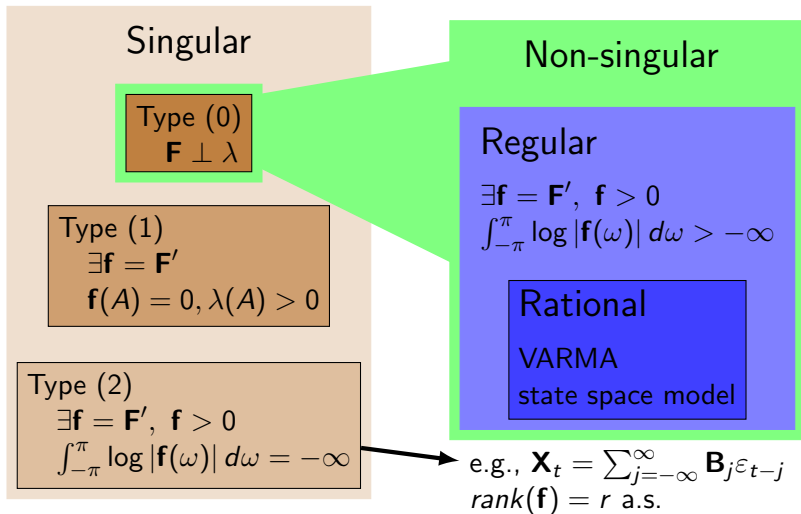
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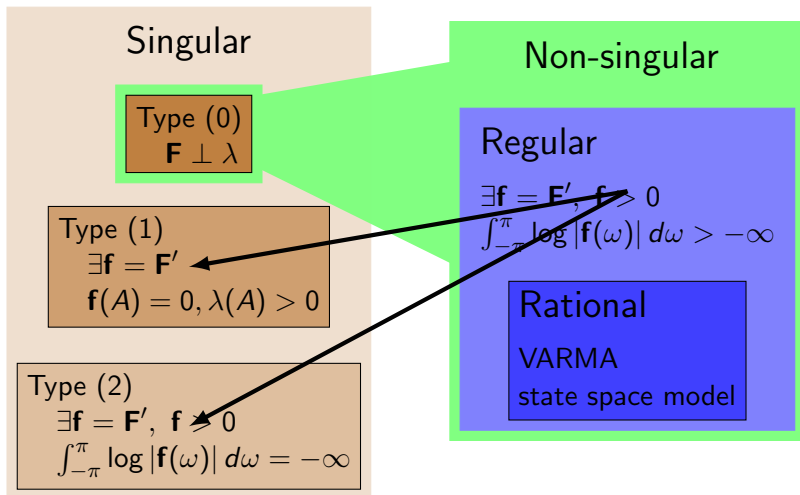
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Singular (deterministic) time series: Cramér, Wold, Wiener, Kolmogorov



Singular-Regular



Singular-Regular

Singular

Type (0)
 $\mathbf{F} \perp \lambda$

Type (1)
 $\exists \mathbf{f} = \mathbf{F}'$
 $\mathbf{f}(A) = 0, \lambda(A) > 0$

Type (2)
 $\exists \mathbf{f} = \mathbf{F}', \mathbf{f} \not\rightarrow 0$
 $\int_{-\pi}^{\pi} \log |\mathbf{f}(\omega)| d\omega = -\infty$

Non-singular

Regular

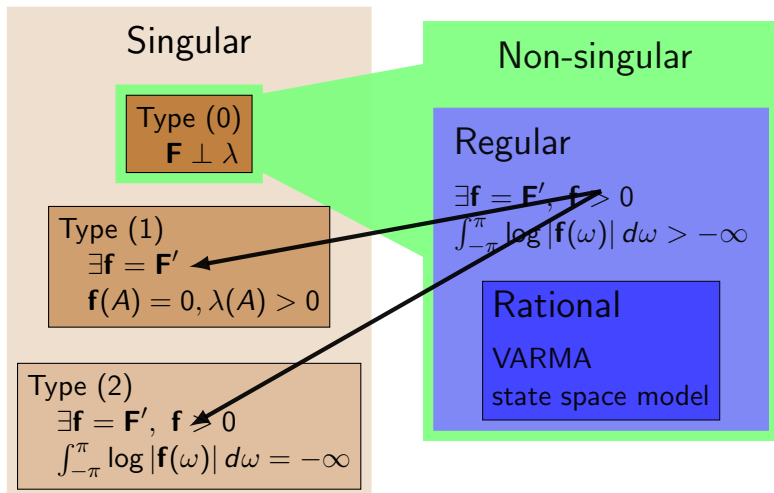
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Rational

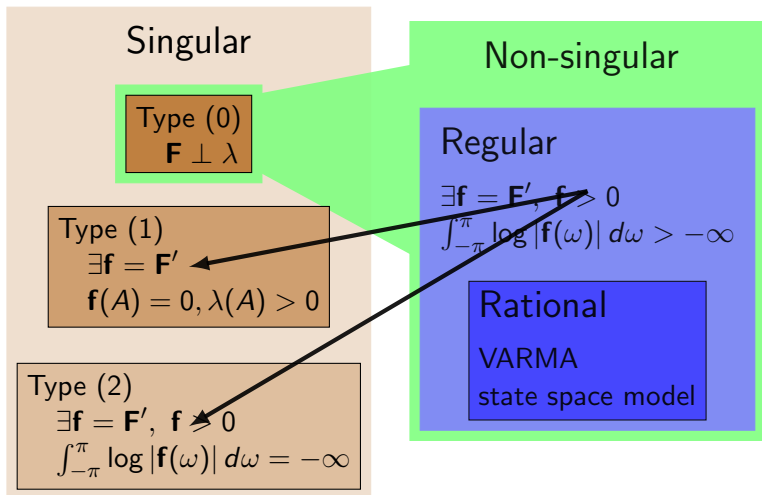
VARMA

state space model

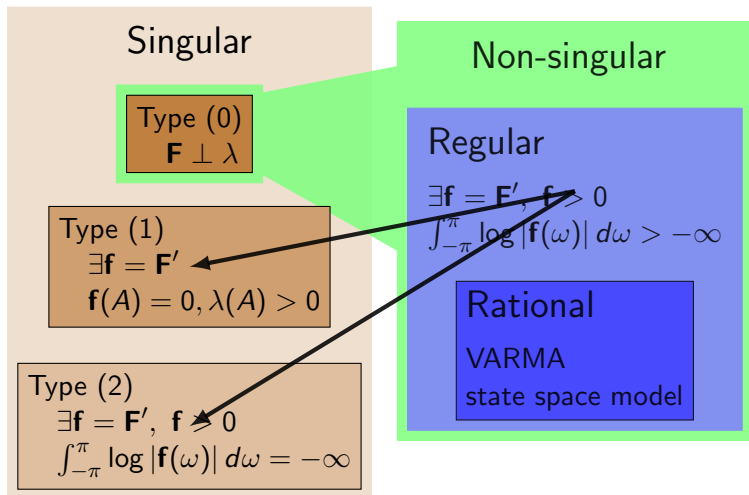
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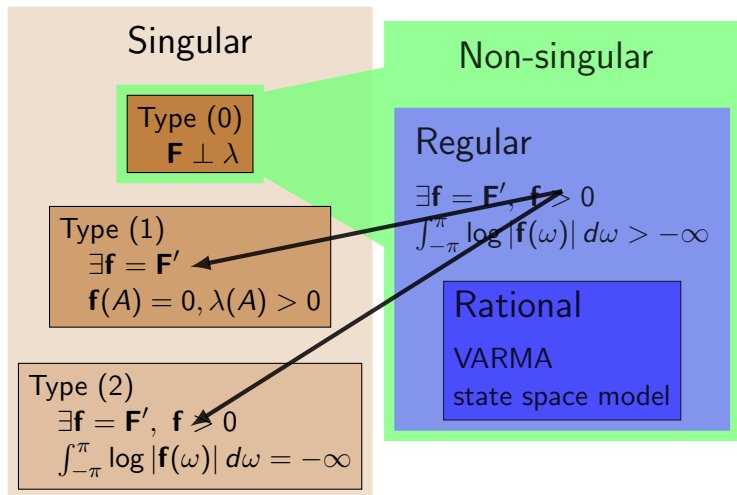
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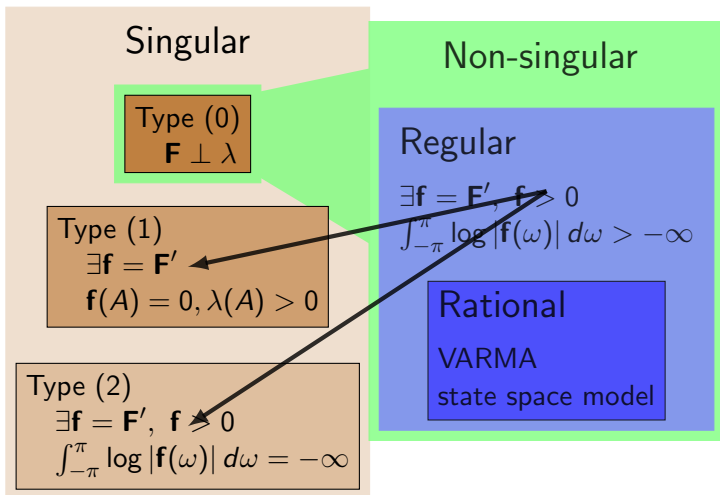
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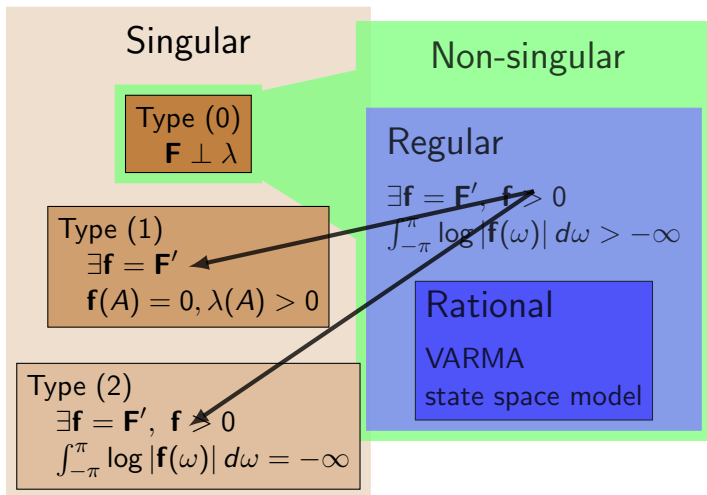
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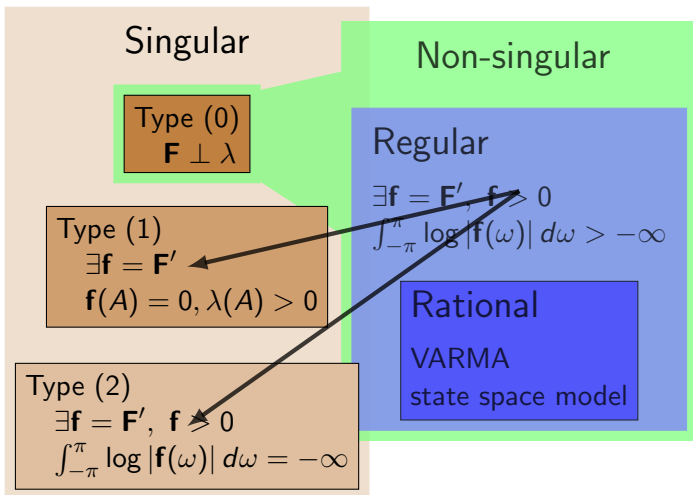
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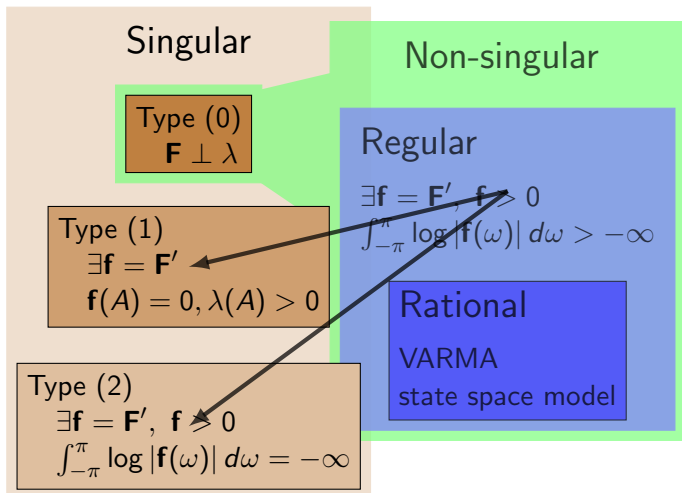
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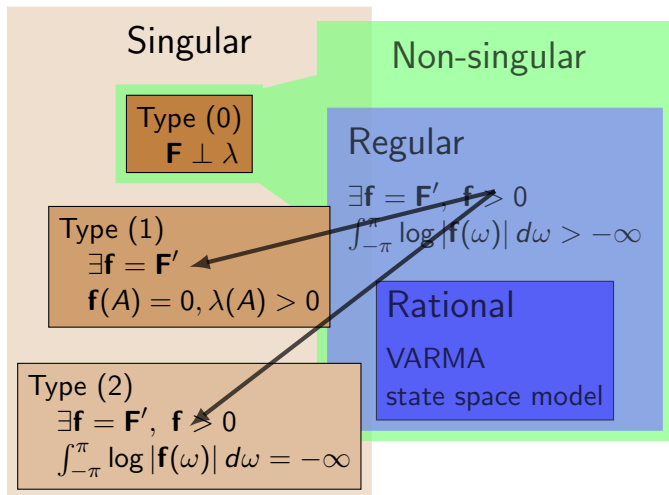
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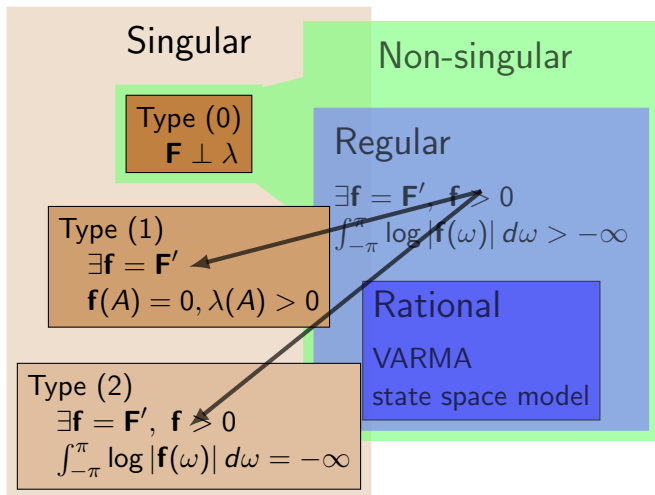
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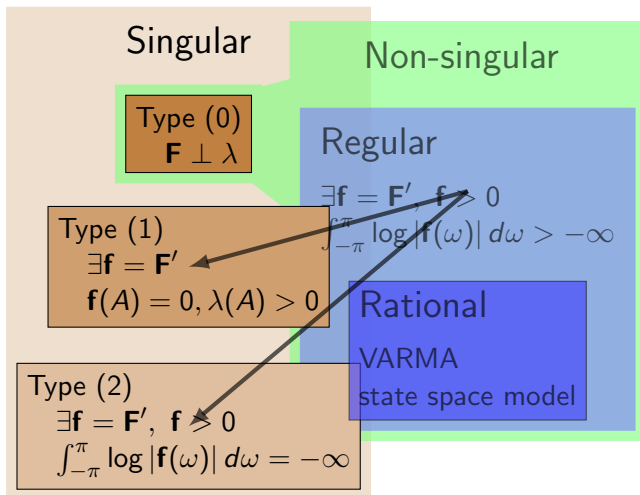
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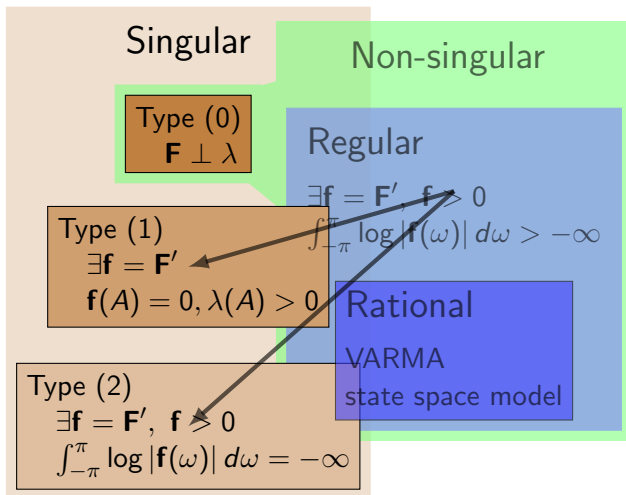
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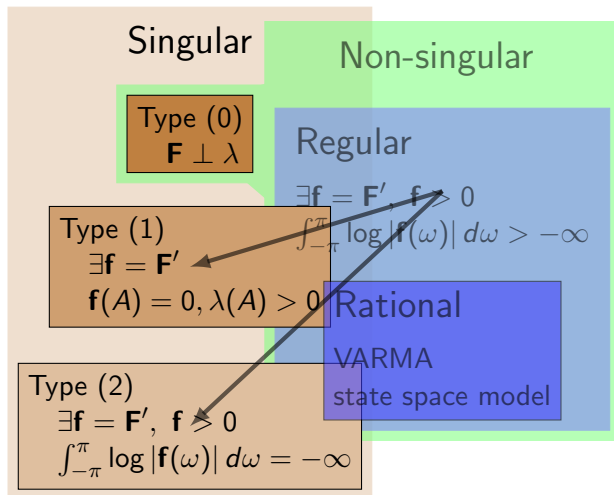
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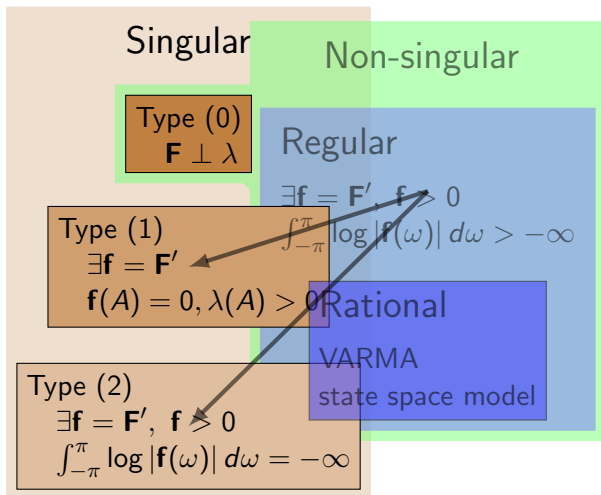
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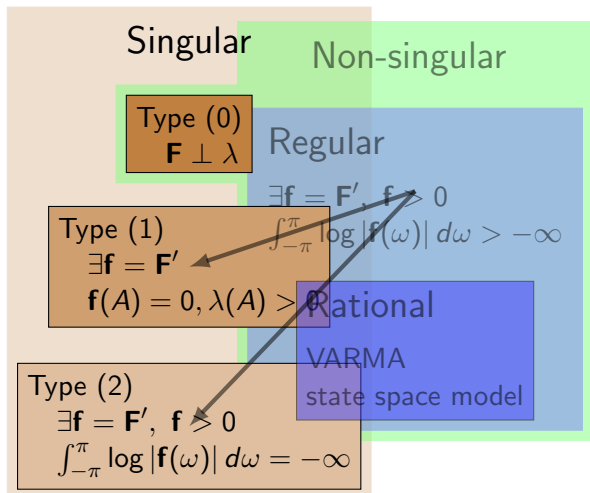
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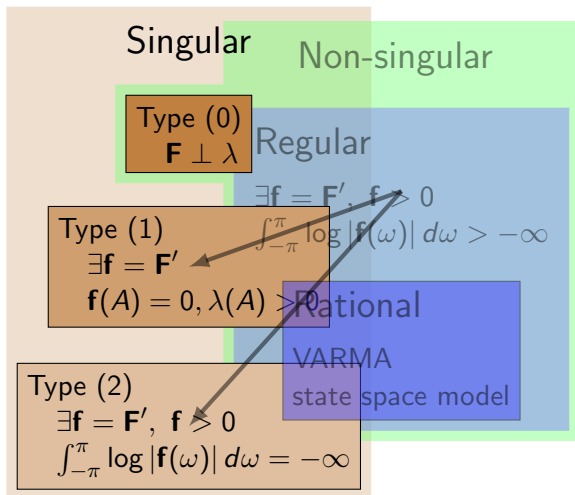
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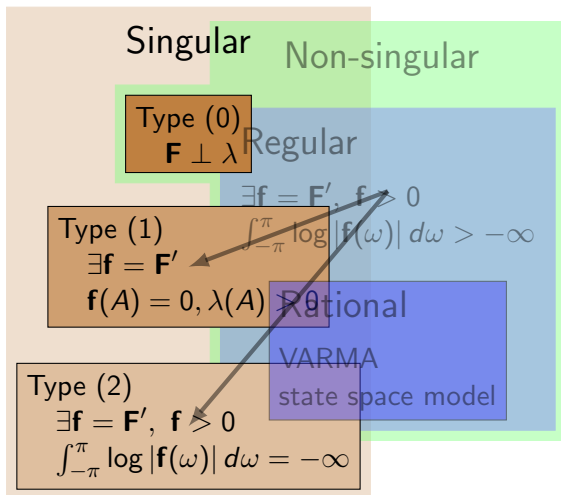
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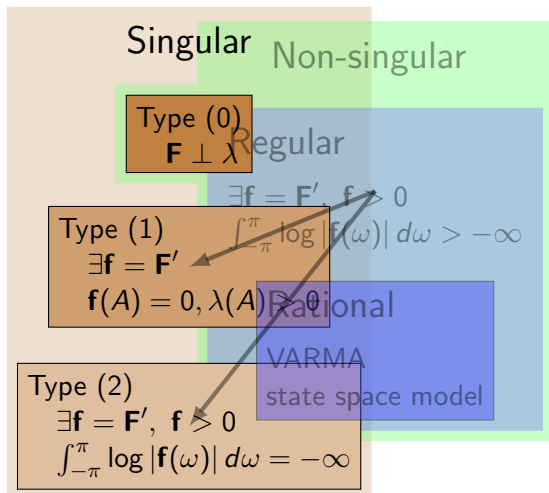
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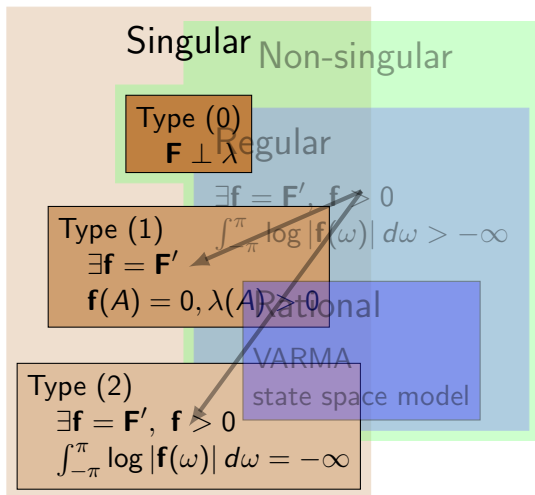
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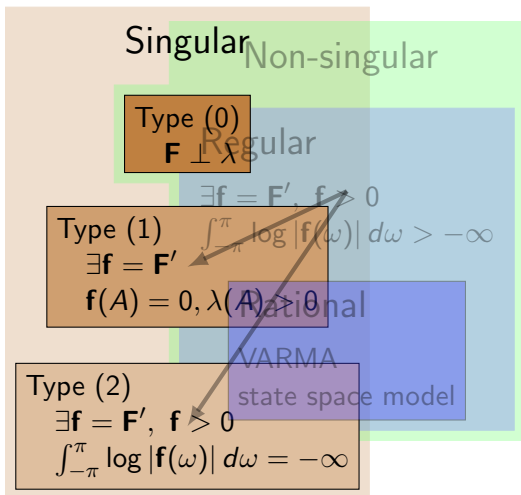
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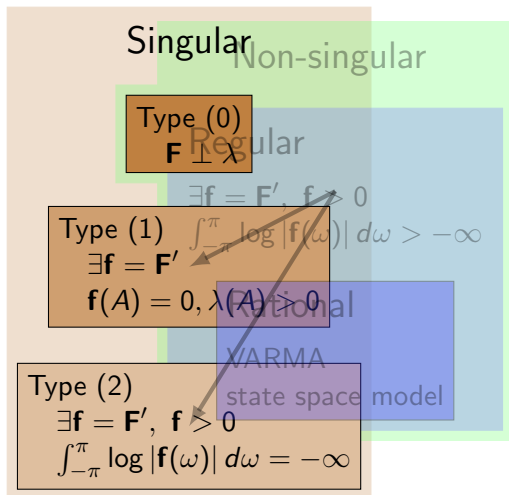
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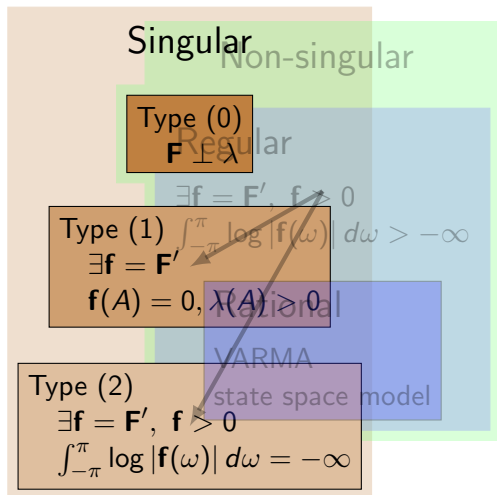
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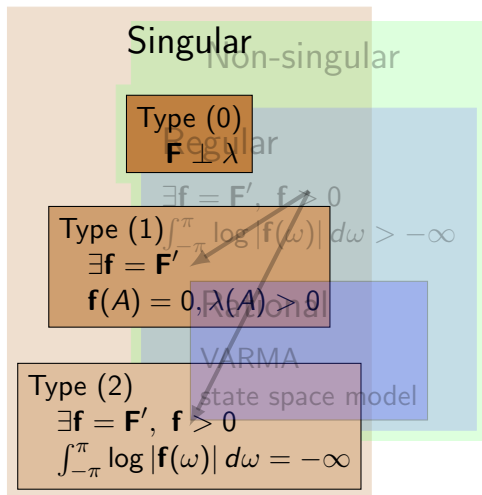
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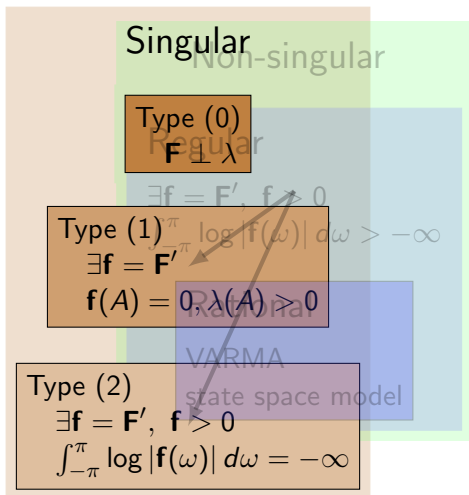
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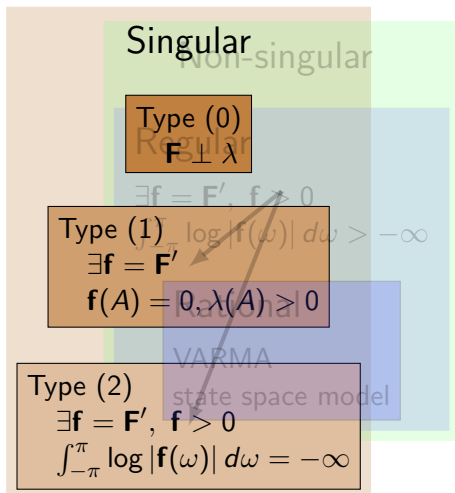
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◀ ...back to the original picture

Non-singular time series: Wold decomposition

H. Wold's Theorem:

$$\mathbf{X}_t = \sum_{j=0}^{\infty} \mathbf{B}_j \varepsilon_{t-j} + \text{Type (0) singular},$$

where \mathbf{B}_j : $n \times r$ matrix (with square-summable entries), and $\{\varepsilon_t\}$ r -dimensional *white noise* with covariance matrix Σ .
If $\{\mathbf{X}_t\}$ is **regular** with $n \times n$ symmetric, positive semidefinite *spectral density matrix* \mathbf{f} , $\text{rang } \mathbf{f}(\omega) = r \leq n$ (a.s.) and

$$\mathbf{f}(z) \iff (\mathbf{B}(z), \Sigma) : \quad \mathbf{f}(z) = \frac{1}{2\pi} \mathbf{B}(z) \Sigma \mathbf{B}^*(z),$$

where

$$\mathbf{B}(z) = \sum_{j=0}^{\infty} \mathbf{B}_j z^j, \quad |z| \leq 1$$

transfer-function.

General Dynamic Factor Model (GDFM)

If $r = n$: $\mathbf{X}_t = \widehat{\mathbf{X}}_{t|t-1} + \varepsilon_t$, then

$\widehat{\mathbf{X}}_{t|t-1}$: *one-step ahead prediction* (based on infinite past) and ε_t : **innovation**.

If $r < n$: only the innovation subspace is unique, and

dim (innovation subspace) = rank (\mathbf{f}) = r .

GDFM (Forni, Lippi, Deistler): $n, t \rightarrow \infty$ (from a starting time).

Number of dynamic factors: essential rank of \mathbf{f} (number of structural eigenvalues in the frequency domain).

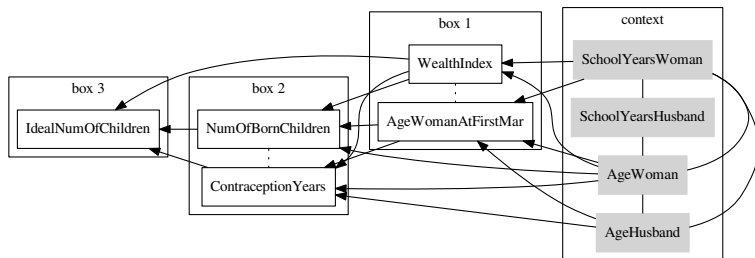
Dynamic factors: standardized innovations.

After eliminating the idiosyncratic part, we predict in the time domain (based on finite past, block Cholesky decomposition, then static factor analysis in the innovation subspaces).

Application: finding underlying signals detected by many sensors. Telecommunication systems (NOKIA), EEG, biomedical sensors.

Graphical models

Chain graph model, EDHS 2014 data



Application: artificial intelligence, medical diagnosis, health, and demography planning in developing countries.

Non-parametric regression: iteration along a DAG

X_1, \dots, X_d : random variables (vertices), $\mathbf{X}_{>k} := (X_{k+1}, \dots, X_d)$: context variables.

$$P_j(\mathbf{x}_{>j}) := \mathbb{E}(X_j | \mathbf{X}_{>j} = \mathbf{x}_{>j}), \quad j = k, k-1, \dots, 1;$$

$$S_j^{(n)}(\mathbf{x}_{>j}) := \sum_{i=1}^n x_j^{(i)} W^{(n,i)}(\mathbf{x}_{>j}), \quad j = k, k-1, \dots, 1$$

smoother (local averaging), e.g., with product kernels.

Consistency Theorem: If

$$\mathbb{E}[S_j^{(n)}(\mathbf{X}_{>j}) - P_j(\mathbf{X}_{>j})]^2 \rightarrow 0, \quad n \rightarrow \infty, \quad j = k, k-1, \dots, 1$$

then

$$\mathbb{E}[S_j^{(n)}(S_{j+1}^{(n)}(S_{j+2}^{(n)}(\dots(S_k^{(n)}(\mathbf{X}_{>k}), \dots), \mathbf{X}_{>k}), \mathbf{X}_{>k}), \mathbf{X}_{>k}) - P_j(P_{j+1}(P_{j+2}(\dots(P_k(\mathbf{X}_{>k}), \dots), \mathbf{X}_{>k}), \mathbf{X}_{>k}), \mathbf{X}_{>k})]^2 \rightarrow 0, \quad n \rightarrow \infty$$

$\forall j = k, k-1, \dots, 1.$

Using Kálmán's filtering in SEM

Structural Equation Model (SEM) as a linear dynamical system:

$$\begin{aligned}\mathbf{B}\eta_t &= \mathbf{A}\xi_t + \zeta_t \\ \mathbf{X}_t &= \mathbf{C}\xi_t + \varepsilon_t \\ \mathbf{Y}_t &= \mathbf{D}\eta_t + \delta_t,\end{aligned}$$

where η_t is m - and ξ_t is n -dimensional latent vector; \mathbf{B} and \mathbf{A} are $m \times m$ and $m \times n$ coefficient matrices; ζ_t is a random vector of residuals. It is uncorrelated with ξ_t , and \mathbf{B} is nonsingular. In the recursive models, \mathbf{B} is upper triangular.

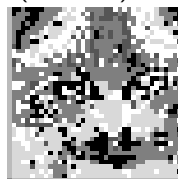
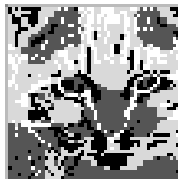
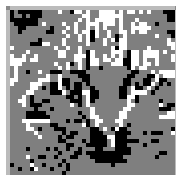
\mathbf{X}_t and \mathbf{Y}_t are q - and p -dimensional observable variables, $q \geq n$ and $p \geq m$. \mathbf{C} is $q \times n$, \mathbf{D} is $p \times m$; \mathbf{A} , \mathbf{B} , \mathbf{C} , \mathbf{D} are estimated from a learning sample (LISLER or block Cholesky decomposition).

ζ_t is an orthogonal process, $\mathbb{E}\xi_s^T \zeta_t = \mathbf{0}$ for $s \leq t$; ε_t is independent of ξ_t , δ_t is independent of η_t , they are also independent of each other and of ζ_t .

Prediction: $\mathbf{X}_t \rightarrow \hat{\xi}_t \rightarrow \hat{\eta}_t \rightarrow \hat{\mathbf{Y}}_t$.

Spectral clustering

Original picture and pixels colored with 3, 4, 5 colors (clusters)



(48 × 48 pixels)

Structural eigenvalues of the normalized modularity matrix:

0.137259, 0.014255, 0.000925,

-0.0006707, -0.0006706, ...

Reproducing Kernel Hilbert Spaces, product Gaussian kernels

image segmentation

Normalized contingency table (e.g., microarray), directed graphs:

$$\mathbf{C}_D = \mathbf{D}_{row}^{-1/2} \mathbf{C} \mathbf{D}_{col}^{-1/2} \quad (\text{correspondence analysis})$$

Application: image segmentation, microarrays (forensic data);

e.g., tumor segmentation, computer aided surgery.

Thank you for your attention.