

Shaped extensions of Singular Spectrum Analysis

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- 1 Introduction to Singular Spectrum Analysis (SSA)
- 2 Contribution: Shaped 2D-SSA
- 3 Contribution: Circular Shaped 2D-SSA
- 4 Contribution: Circular Shaped 2D-ESPRIT
- 5 Contribution: Efficient implementation
- 6 Conclusions

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Common scheme of Singular Spectrum Analysis (SSA)

Input: time series or array (or something else) $\mathbb{X} \in \mathcal{X} \sim \mathbb{R}^N$

Parameter: **embedding operator** $\mathcal{T}: \mathcal{X} \hookrightarrow \mathbb{R}^{L \times K}$, injective, linear

Decomposition

1 Embedding

Trajectory matrix $\mathbf{X} = \mathcal{T}(\mathbb{X})$, **structured** (i.e. $\mathbf{X} \in \mathcal{H} = \text{image } \mathcal{T}$)

2 Singular Value Decomposition (SVD)

$$\mathbf{X} = \sum_{k=1}^d \sqrt{\lambda_k} U_k V_k^T, \{U_k\}_{k=1}^d \text{ and } \{V_k\}_{k=1}^d \text{ are orthonormal}$$

Reconstruction

1 Grouping: $\{1, \dots, d\} = I_1 \sqcup \dots \sqcup I_c$

$$\mathbf{X} = \mathbf{X}_{I_1} + \dots + \mathbf{X}_{I_c}, \text{ where } \mathbf{X}_I = \sum_{k \in I} \sqrt{\lambda_k} U_k V_k^T$$

2 Projection

Matrix $\mathbf{X}_I \xrightarrow{\mathcal{H}=\Pi^{(\mathcal{H})}}$ structured matrix $\tilde{\mathbf{X}}_I \xrightarrow{\mathcal{T}^{-1}}$ reconstructed $\tilde{\mathbb{X}}^{(I)}$

Output: $\mathbb{X} = \tilde{\mathbb{X}}^{(I_1)} + \dots + \tilde{\mathbb{X}}^{(I_c)}$

Particular SSA cases: Classical 1D-SSA

Input: time series $\mathbb{X} = (x_1, x_2, \dots, x_N)$

Parameter: L , **window length**; $1 < L < N$; $K = N - L + 1$

The **embedding operator**:

$$\mathcal{T}(\mathbb{X}) = \mathcal{T}_L(\mathbb{X}) := [X_1 : \dots : X_K] = \begin{pmatrix} x_1 & x_2 & \dots & x_{K-1} & x_K \\ x_2 & \ddots & \dots & \ddots & x_{K+1} \\ \vdots & \vdots & & \vdots & \vdots \\ x_{L-1} & \ddots & \dots & \ddots & x_{N-1} \\ x_L & x_{L+1} & \dots & x_{N-1} & x_N \end{pmatrix},$$

where the columns X_j are lagged subvectors of length L :

$$X_j = (x_j, x_{j+1}, \dots, x_{j+L-1})^T$$

- $H = \text{image } \mathcal{T}_L$ is the set of **Hankel** matrices
- Projection $\Pi^{(H)}$ is **diagonal averaging** (or “**hankelization**”)

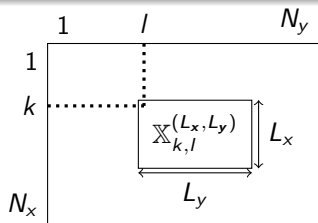
Particular SSA cases: 2D-SSA

Input: 2D array $\mathbb{X} = \begin{pmatrix} x_{1,1} & \dots & x_{1,N_y} \\ \vdots & & \vdots \\ x_{N_x,1} & \dots & x_{N_x,N_y} \end{pmatrix}$

Parameters: window sizes L_x, L_y ;

$1 < L_x < N_x$; $1 < L_y < N_y$;

$K_x = N_x - L_x + 1$; $K_y = N_y - L_y + 1$;



The **embedding operator**:

$$\mathcal{T}_{L_x, L_y}(\mathbb{X}) := \mathbf{X} = [X_1 : \dots : X_{K_x K_y}] \in \mathbb{R}^{L_x L_y \times K_x K_y},$$

where the columns X_j are vectorizations of **shifted** $L_x \times L_y$ subarrays:

$$X_{k+(l-1)K_x} = \text{vec}(\mathbb{X}_{k,l}^{(L_x, L_y)})$$

- $\mathbf{H} = \text{image } \mathcal{T}_{L_x \times L_y}$ is the set of **Hankel-block-Hankel** matrices
- Projection $\Pi^{(\mathbf{H})}$ is “**block-hankelization**” (also **averaging**)

Note: $\mathbb{R}^{L_x \times L_y} \sim \mathbb{R}^{L_x L_y}$, thus any singular vector U can be considered as $L_x \times L_y$ array $\mathbb{U}$

2D-SSA example: Gentlemen

Eigenvectors

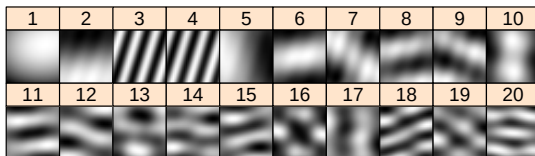


Figure: Gentlemen, 2D-SSA: Eigenarrays, $(L_x, L_y) = (25, 25)$

Reconstructions

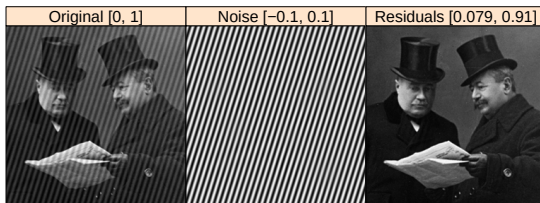


Figure: Gentlemen, 2D-SSA: Separated periodic noise, $(N_x, N_y) = (256, 256)$, $(L_x, L_y) = (25, 25)$, $I = \{3, 4\}$

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Problem: non-rectangular arrays & edge effects

Eigenvectors

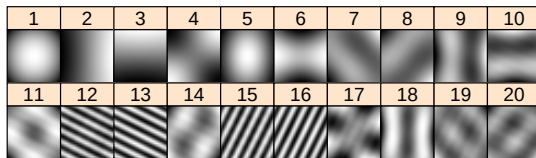


Figure: Mars, 2D-SSA: Eigenarrays, $(L_x, L_y) = (25, 25)$

Reconstructions

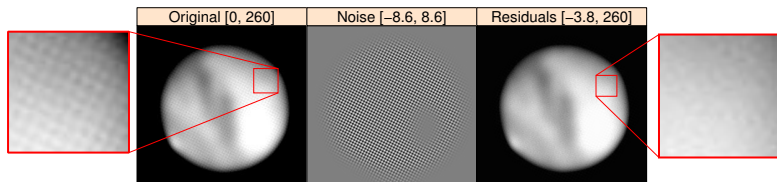


Figure: Mars, 2D-SSA: Separated periodic noise, $(N_x, N_y) = (258, 275)$, $(L_x, L_y) = (25, 25)$, $I = \{12, 13, 15, 16\}$

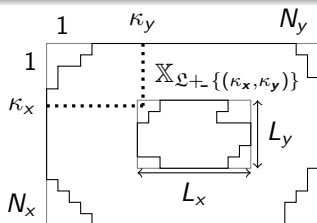
Let's develop "*shaped*" SSA, by analogy with *shaped* filters

New method: Shaped 2D-SSA

Input: $\mathfrak{N}$ -shaped array $\mathbb{X} = \mathbb{X}_{\mathfrak{N}} = (x_{\eta})_{\eta \in \mathfrak{N}}$, where

$\mathfrak{N} = \{\eta_1, \dots, \eta_N\} \subseteq \{1, \dots, N_x\} \times \{1, \dots, N_y\}$

Parameter: window shape $\mathcal{L} = \{\ell_1, \dots, \ell_L\} \subseteq \{1, \dots, L_x\} \times \{1, \dots, L_y\}$



The **embedding operator**:

$$\mathcal{T}_{\mathcal{L}}(\mathbb{X}) := \mathbf{X} = [X_1 : \dots : X_K] \in \mathbb{R}^{L \times K},$$

where columns $X_j = (x_{\ell_i + \kappa_j})_{i=1}^L$ are vectorizations of **shifted $\mathcal{L}$ -shaped arrays** $\mathbb{X}_{\mathcal{L} + \{-\kappa_j\}} = (x_{\eta})_{\eta \in \mathcal{L} + \{-\kappa_j\}}$

And $\mathfrak{K}$ is the set of all possible **origin positions** for **$\mathcal{L}$ -shaped windows**:

$$\mathfrak{K} = \{\kappa_1, \dots, \kappa_K\} = \{\kappa \in \mathbb{N}^2 \mid \mathcal{L} + \{-\kappa\} \subseteq \mathfrak{N}\},$$

where $\cdot + \eta$ means **shape shift** by vector η : $\mathfrak{A} + \eta := \{\alpha + \eta - 1 \mid \alpha \in \mathfrak{A}\}$

- $\mathbf{H} = \text{image } \mathcal{T}_{\mathcal{L}}$ is the set of **quasi-Hankel** matrices
- Projection $\Pi^{(\mathbf{H})}$ is “**quasi-hankelization**” (again, **averaging**)

Note: $\mathbb{R}^{\mathcal{L}} \sim \mathbb{R}^L$, thus any singular vector U can be considered as **$\mathcal{L}$ -shaped subarray** $\mathbb{U}$

Eigenvectors

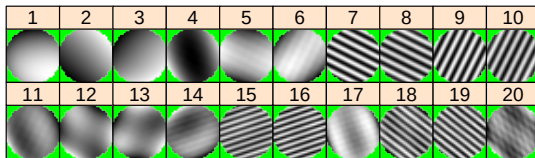


Figure: Mars, ShSSA: Eigenarrays, $\mathcal{L}$ is circle of radius 15

Reconstructions

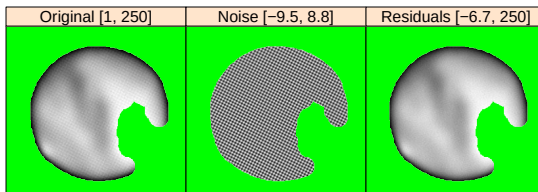


Figure: Mars: Separated period noise, $\mathcal{L}$ is circle of radius 15, $I = \{7, 8, 9, 10\}$

Comparison: shaped vs non-shaped

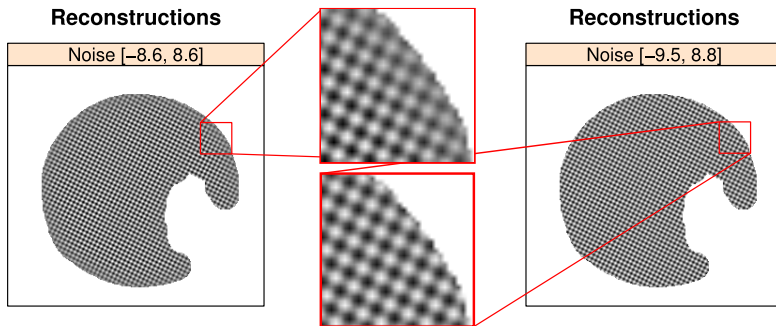


Figure: Left: cutted image of periodic noise from rectangular SSA; right: image of periodic noise from Shaped SSA

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Problem: circular topology

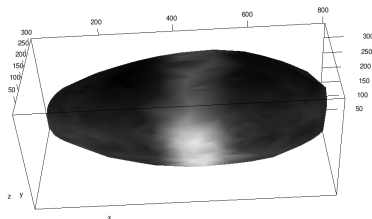
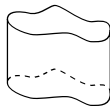


Figure: Drosophila, 3d intensity plot, “Krüppel” gene

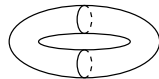
Three **topology** cases:



(a) Planar topology
($T_x = T_y = \infty$)



(b) Cylindrical topology
($T_x \neq \infty$; $T_y = \infty$)



(c) Toroidal topology
($T_x \neq \infty$; $T_y \neq \infty$)

New method: Circular Shaped 2D-SSA

Input: topology characteristics $T_x, T_y \in \mathbb{N} \cup \{\infty\}$,
 $\mathfrak{N}$ -shaped array $\mathbb{X} = \mathbb{X}_{\mathfrak{N}} = (x_{\eta})_{\eta \in \mathfrak{N}}$, where
 $\mathfrak{N} = \{\eta_1, \dots, \eta_N\} \subseteq \{1, \dots, T_x\} \times \{1, \dots, T_y\}$
Parameter: window shape $\mathfrak{L} = \{\ell_1, \dots, \ell_L\} \subseteq$
 $\subseteq \{1, \dots, L_x\} \times \{1, \dots, L_y\}$

The **embedding operator**:

$$\mathcal{T}_{\mathfrak{L}}(\mathbb{X}) := \mathbf{X} = [X_1 : \dots : X_K] \in \mathbb{R}^{L \times K},$$

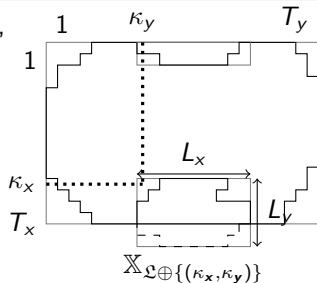
where columns $X_j = (x_{\ell_i + \kappa_j})_{i=1}^L$ are vectorizations of **shifted $\mathfrak{L}$ -shaped** arrays

And $\mathfrak{K}$ is the set of all possible **origin positions** for $\mathfrak{L}$ -shaped windows:

$$\mathfrak{K} = \{\kappa_1, \dots, \kappa_K\} = \{\kappa \in \mathbb{N}^2 \mid \mathfrak{L} \oplus \{\kappa\} \subseteq \mathfrak{N}\},$$

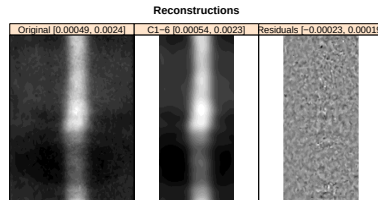
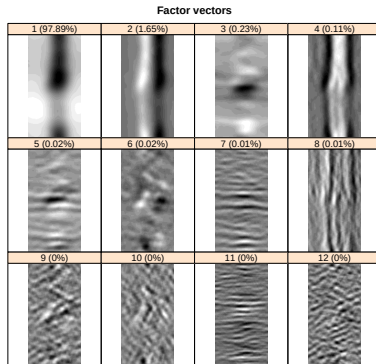
where $\cdot \oplus \eta$ means **cycled shape shift** by vector η :

$$\mathfrak{A} \oplus \eta := \{((\alpha_x + \eta_x - 2) \bmod T_x + 1, (\alpha_y + \eta_y - 2) \bmod T_y + 1) \mid \alpha \in \mathfrak{A}\}$$



- $\mathbf{H} = \text{image } \mathcal{T}_{\mathfrak{L}}$ is the set of **partially circulant quasi-Hankel** matrices
- Projection $\Pi^{(\mathbf{H})}$ is **"quasi-hankelization"** (and again, **averaging**)

Real life example: drosophila embryo



(e) Drosophila, reconstruction and residuals

(d) Drosophila, factor vectors

Figure: Drosophila, Circular SSA, cylindrical case ($T_x = N_x$ and $T_y = \infty$).

The data are downloaded from the BDTNP archive, the file “v5_s11643-28no06-04.pca”, gene “Krüppel”.

Middle part from 20% to 80% of the embryo length was processed.

Interpolation step is 0.5% and both window sizes are 10% of the embryo length

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2D-ESPRIT for shaped array

Parametric model: corrupted finite-rank array:

$$(\mathbb{S})_{l,n} = \sum_{k=1}^r A_k \mu_k^l \nu_k^n, \quad A_k, \mu_k, \nu_k \in \mathbb{C} \setminus \{0\}$$

$\mathbb{X} = \mathbb{S} + \mathbb{R}$, where $\mathbb{R}$ is noise or another signal

Problem: estimate nonlinear parameters $\{(\mu_k, \nu_k)\}_{k=1}^r$

Algorithm rectangular **2D-ESPRIT** (Rouquette and Najim, 2001):

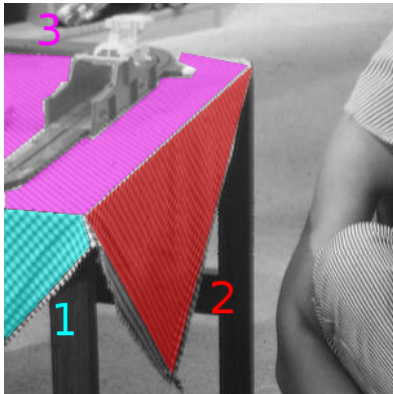
- ➊ **Estimation of the signal subspace:** Get the basis $\mathbb{U}_{i_1}, \dots, \mathbb{U}_{i_r}$ of an estimate of signal subspace (e.g. from SSA)
- ➋ **Construction of truncated matrices:**
 $\mathbf{P}_x = [\text{vec}(\underline{\mathbb{U}_{i_1}}) : \dots : \text{vec}(\underline{\mathbb{U}_{i_r}})], \mathbf{Q}_x = [\text{vec}(\overline{\mathbb{U}_{i_1}}) : \dots : \text{vec}(\overline{\mathbb{U}_{i_r}})],$
 $\mathbf{P}_y = [\text{vec}(\mathbb{U}_{i_1}|) : \dots : \text{vec}(\mathbb{U}_{i_r}|)], \mathbf{Q}_y = [\text{vec}(|\mathbb{U}_{i_1}) : \dots : \text{vec}(|\mathbb{U}_{i_r})],$
where $(\underline{\cdot})$, $(\overline{\cdot})$, $(\cdot|)$, $(|\cdot)$ denote the array without last row, first row last column and first column respectively
- ➌ **Construction of shift matrices:** Approximate solution of $\mathbf{P}_x \mathbf{M}_x \approx \mathbf{Q}_x$ and $\mathbf{P}_y \mathbf{M}_y \approx \mathbf{Q}_y$ in LS- or TLS-sense
- ➍ **Estimation of the parameters:** EVD-decomposition of $\mathbf{M}_x$ and $\mathbf{M}_y$, eigenvalues $\hat{\mu}_i$ and $\hat{\nu}_j$ is an estimates of the parameters

This talk: [from rectangular to shaped arrays](#)

Real life example: Barbara's cloth



(a) Barbara, whole image



(b) Barbara, table with enumerated subimages

Real life example: Barbara's cloth

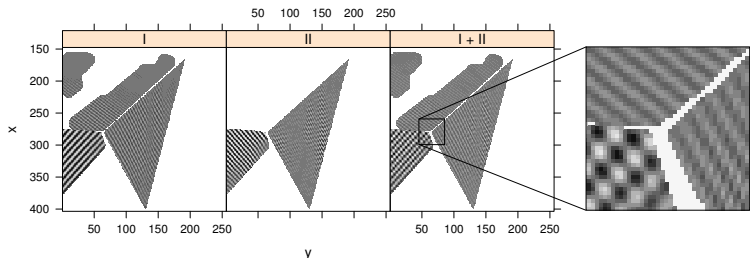


Figure: Barbara's table: reconstructions of texture.

$I_I = \{2, 3\}$, $I_{II} = \{4, 5\}$ (if present)

Table: Barbara, ESPRIT periods and rates

Part	t_x	t_y	Width	$\alpha, ^\circ$	Rate_x	Rate_y
1	8.1	6.9	5.2	50	-0.0066	-0.0061
1	11.0	10.1	7.4	47	-0.0022	-0.0064
2	9.7	5.0	4.4	63	-0.0031	-0.0075
2	7.2	2.6	2.5	70	0.0025	0.017
3	5.1	6.8	4.1	37	0.0054	-0.0046

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Main ideas:

- 1 The **truncated SVD** calculated by **Lanczos methods** is used, since only a number of leading SVD components correspond to the signal (**ν -TRLan** and **PROPACK** libraries are used)
- 2 In Lanczos methods, only multiplication of $\mathbf{X}$ (and $\mathbf{X}^T$) by a vector is needed. Due to quasi-Hankel structure it can be implemented via **FFT** (**FFTW** library is used)
- 3 At the Reconstruction step, **hankelization** or **quasi-hankelization** of a matrix of rank 1 can be also implemented via FFT

Total complexity: $\mathcal{O}(rN \log N)$

R implementation:

R package “**RSSA**”: <http://cran.r-project.org/web/packages/Rssa/>

Article:

N. Golyandina, A. Korobeynikov, A. Shlemov and K. Usevich (2014):
“*Multivariate and 2D Extensions of Singular Spectrum Analysis with the RSSA Package*”, accepted to “Journal of Statistical Software”

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- An extension of SSA (shaped SSA) was proposed for decomposition of shaped images including circular and toroidal topology
- An extension of 2D-ESPRIT method for parameter estimation in the signal-plus-noise model was described
- An efficient publicly available implementation is provided
- The algorithms and their implementation can be easily extended to higher dimensions

Thanks for your attention!