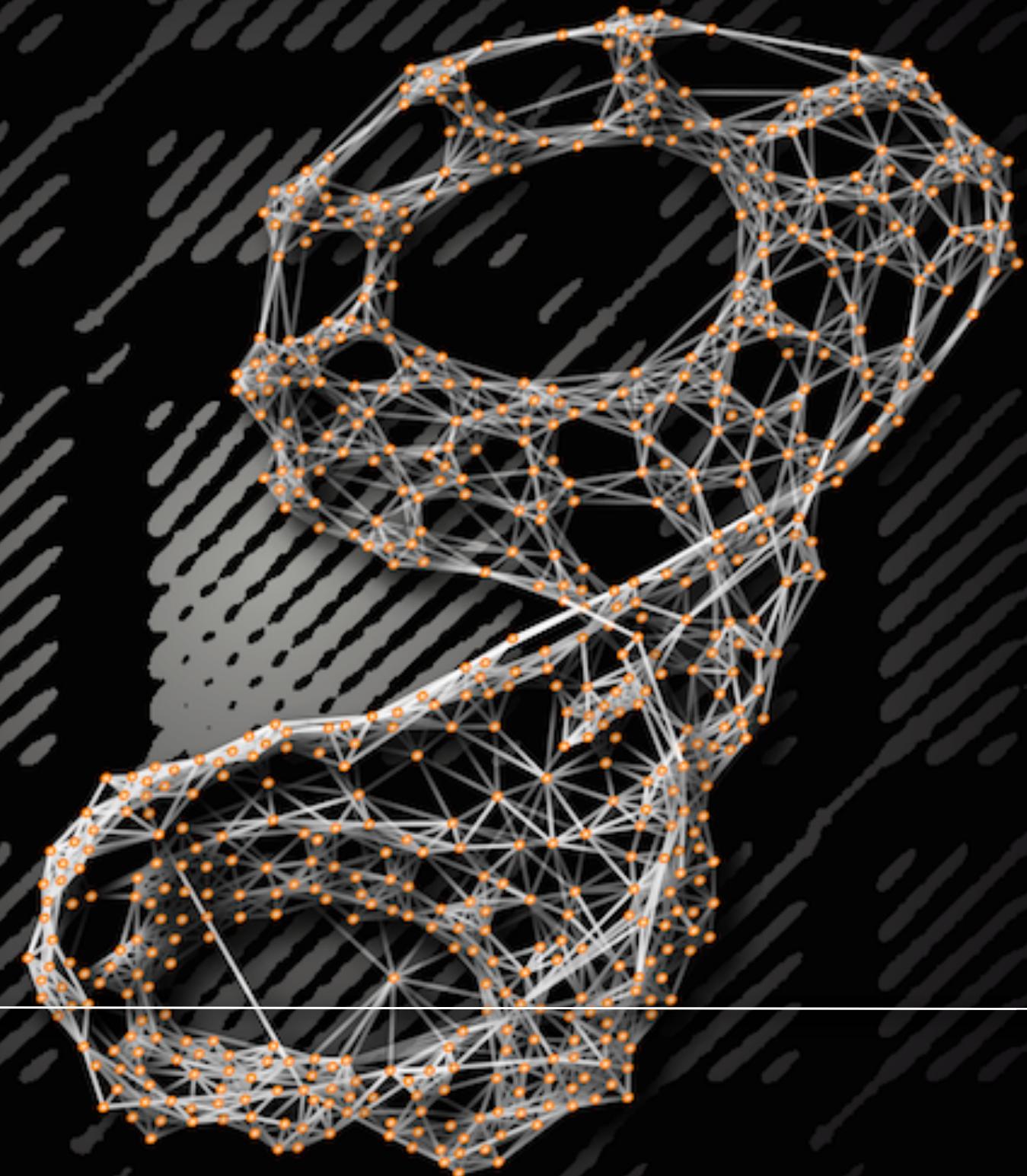




POTSDAM INSTITUTE FOR  
CLIMATE IMPACT RESEARCH

NORBERT MARWAN  
HAUKE KRÄMER

# RECURRENCE PLOTS WORKSHOP

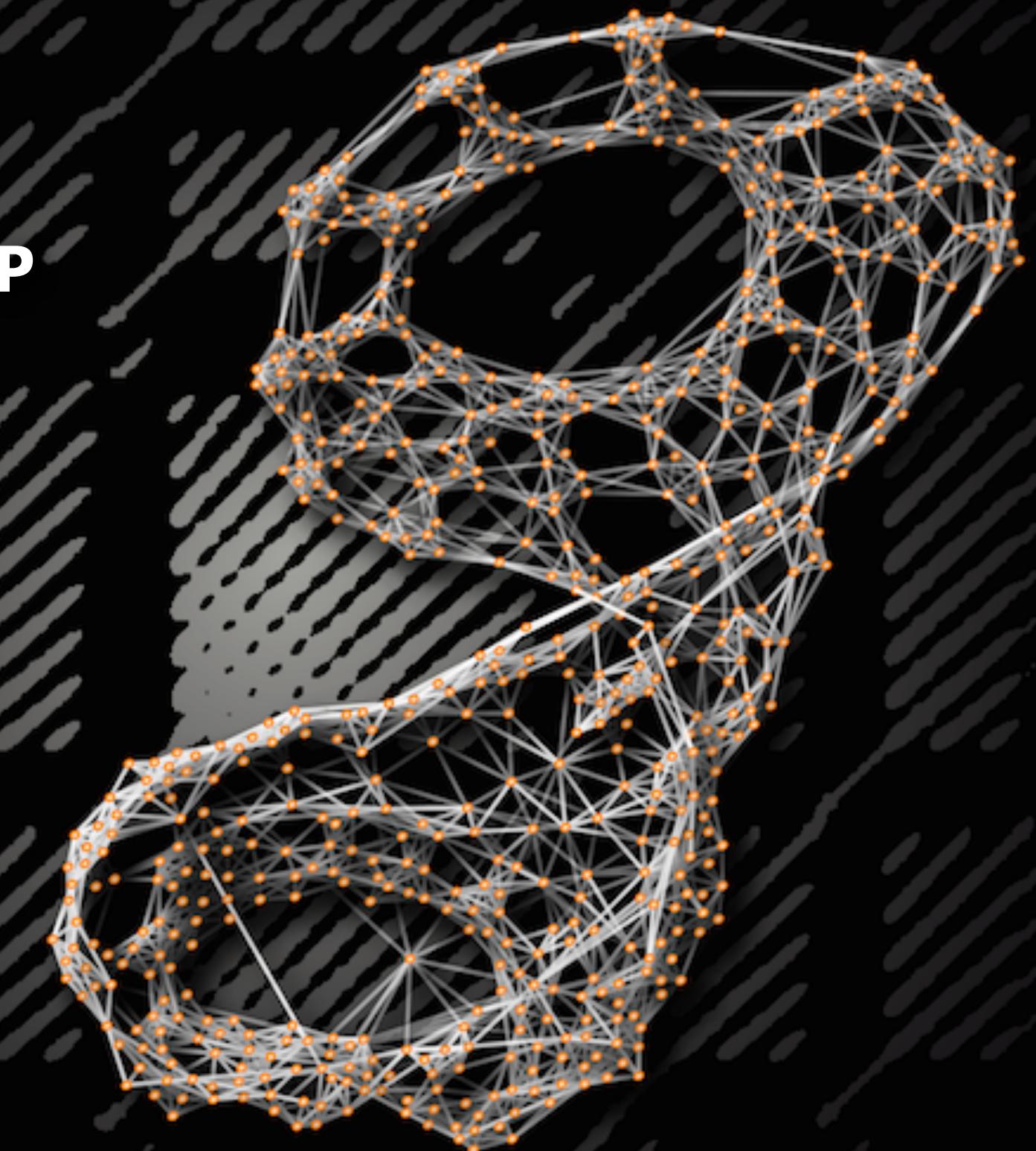




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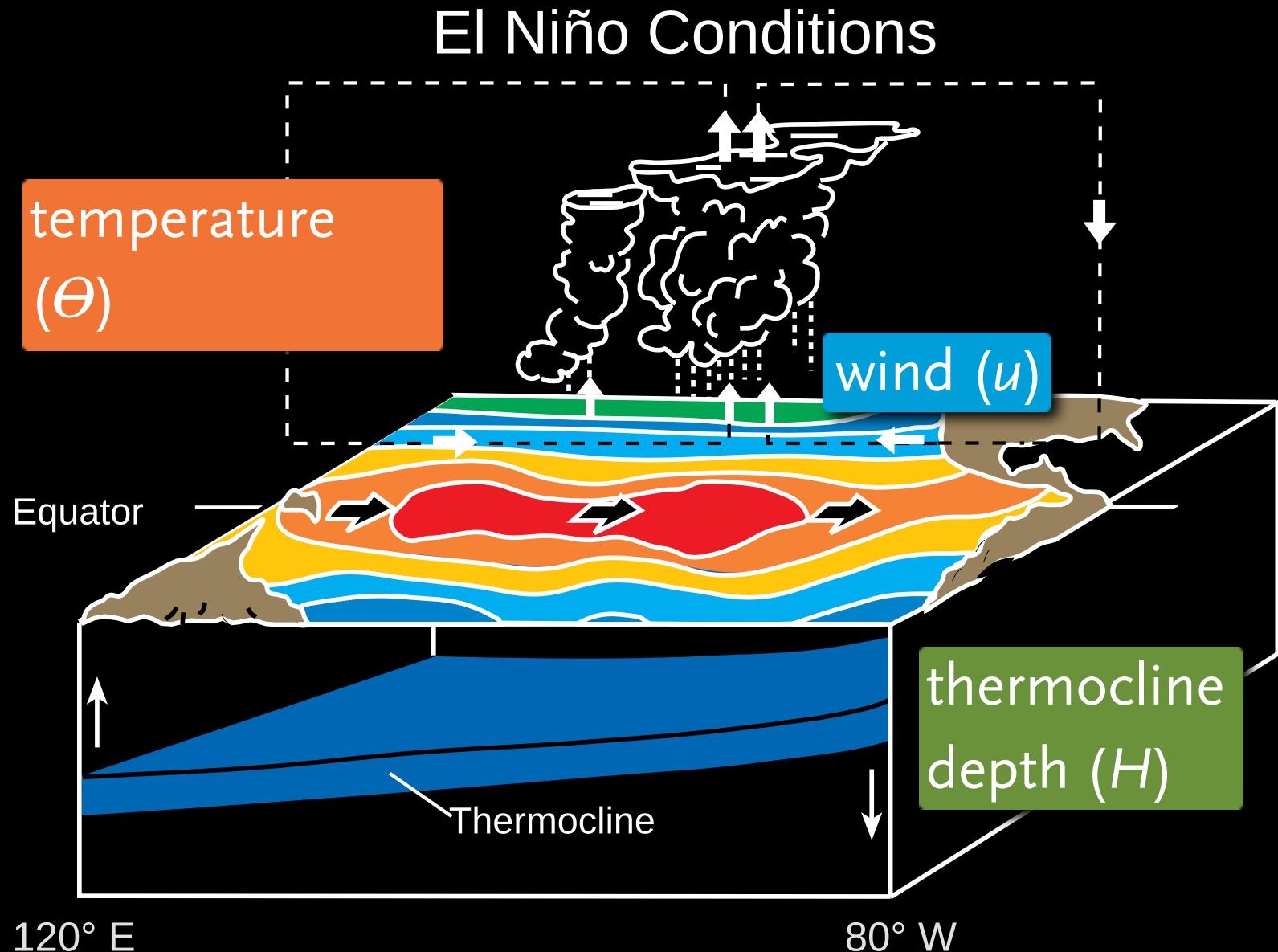
# RECURRENCE PLOTS WORKSHOP

- ① Phase space
- ② Visual interpretation
- ③ Sensitivity on Parameters
- ④ Recurrence quantification analysis
- ⑤ Coupling analysis



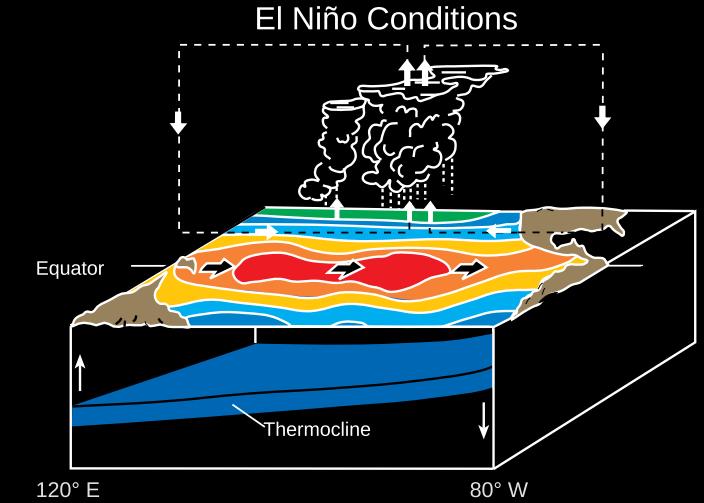
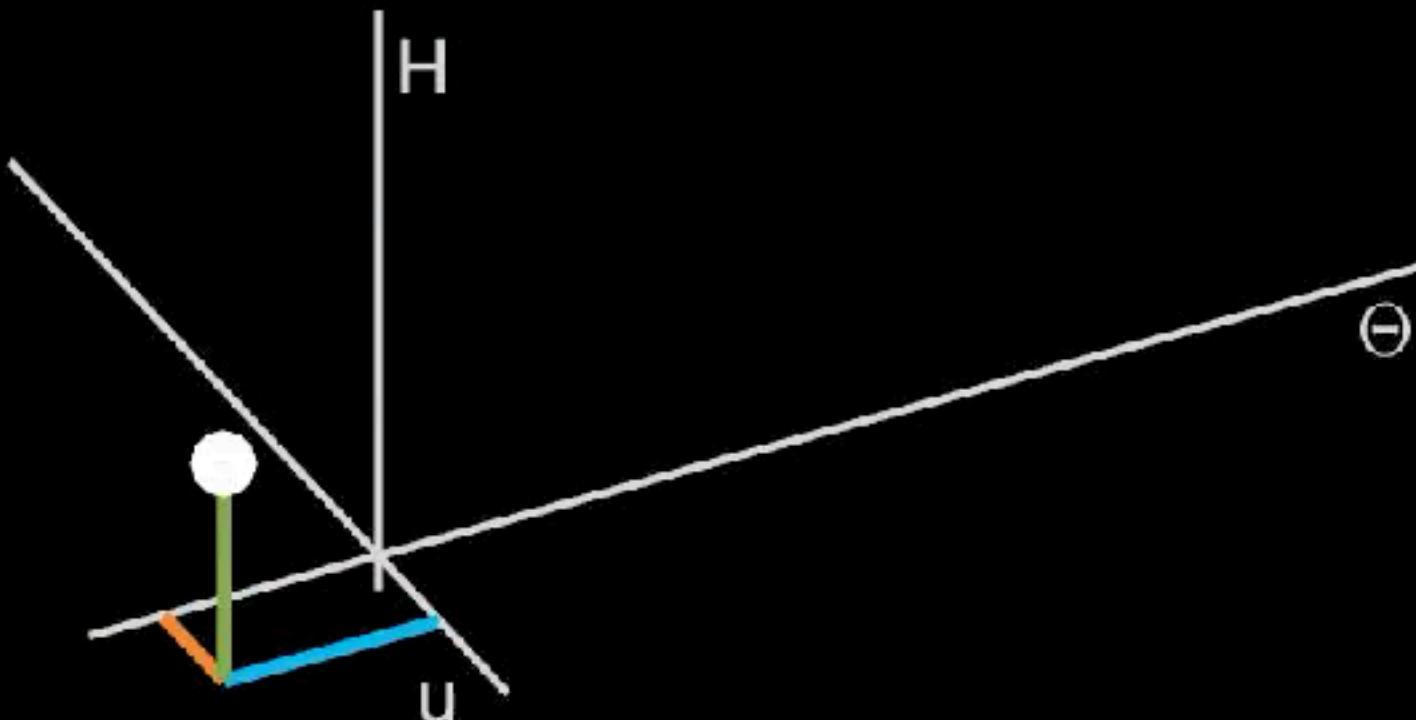
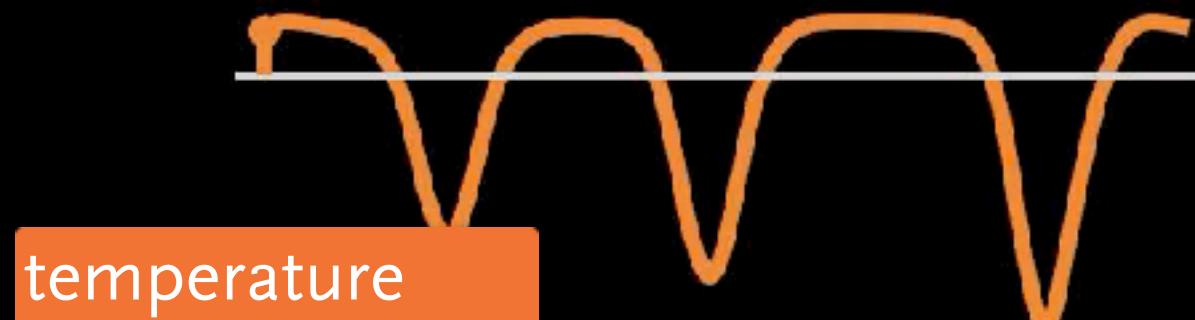
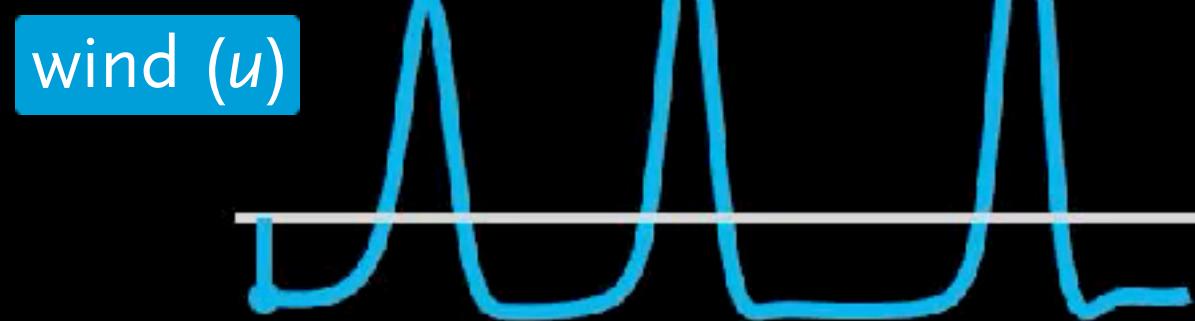
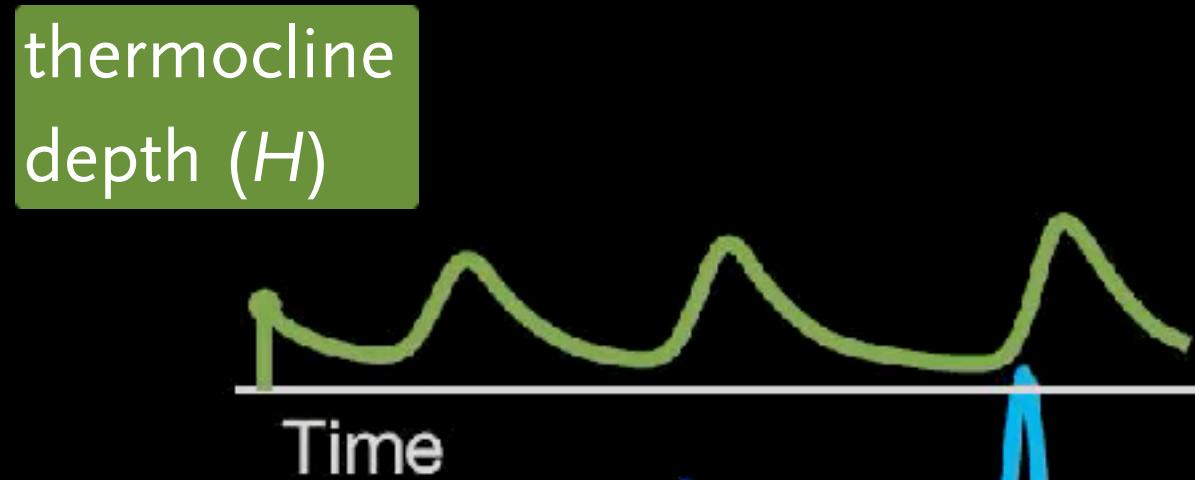
# PHASE SPACE

# PHASE SPACE

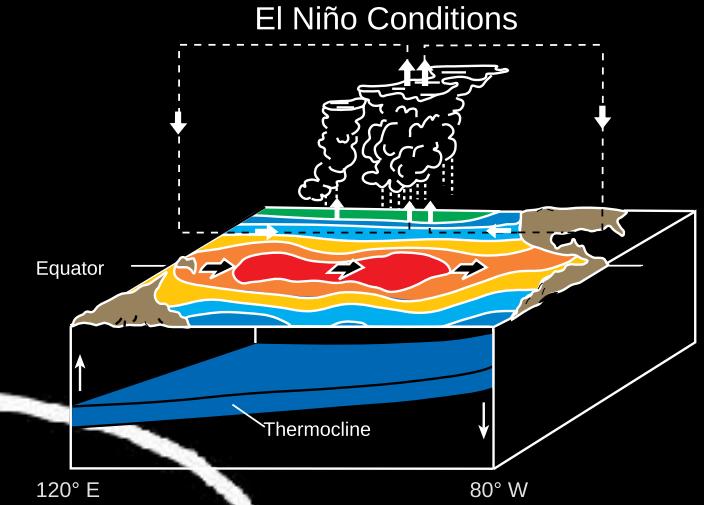
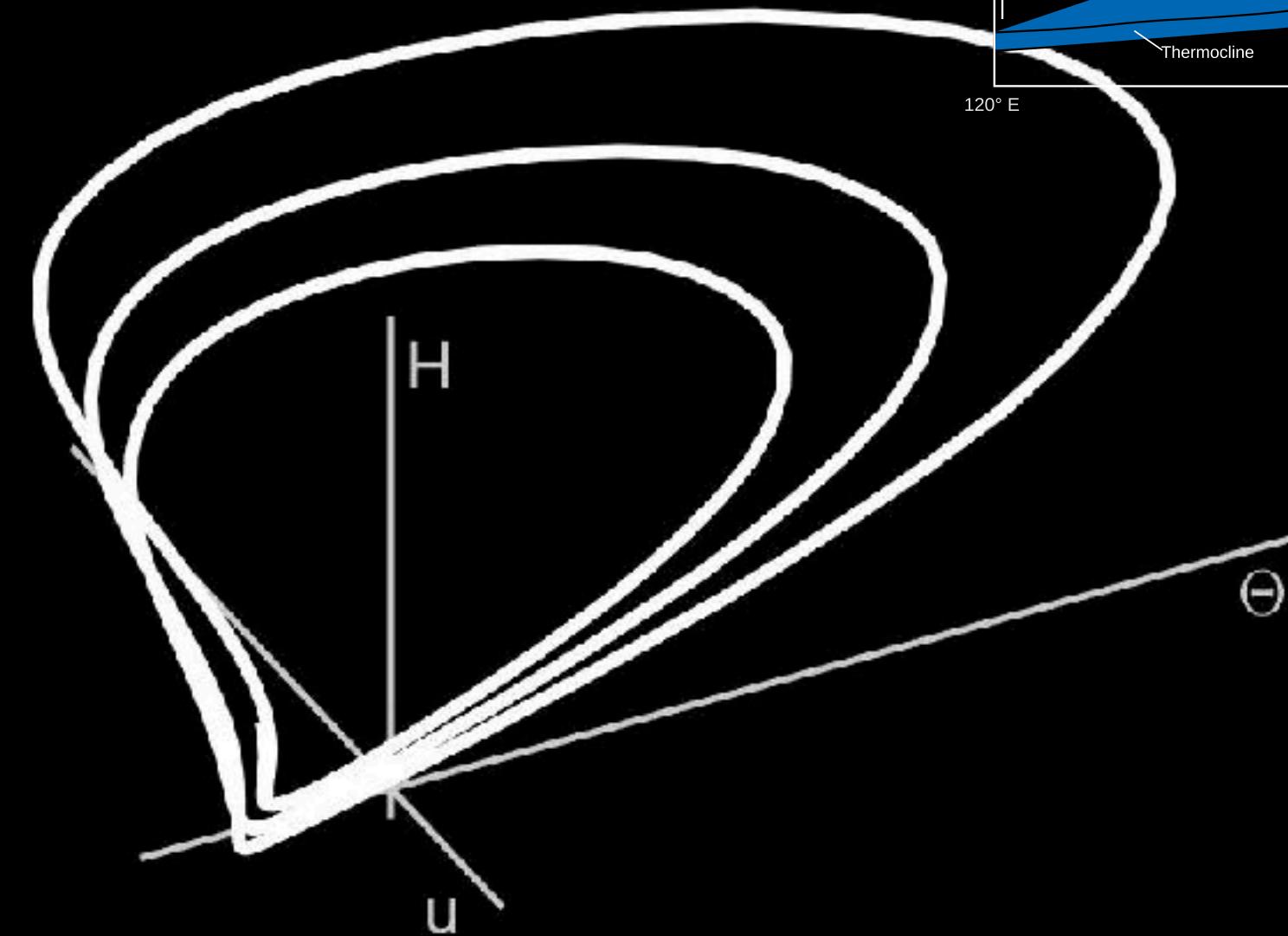
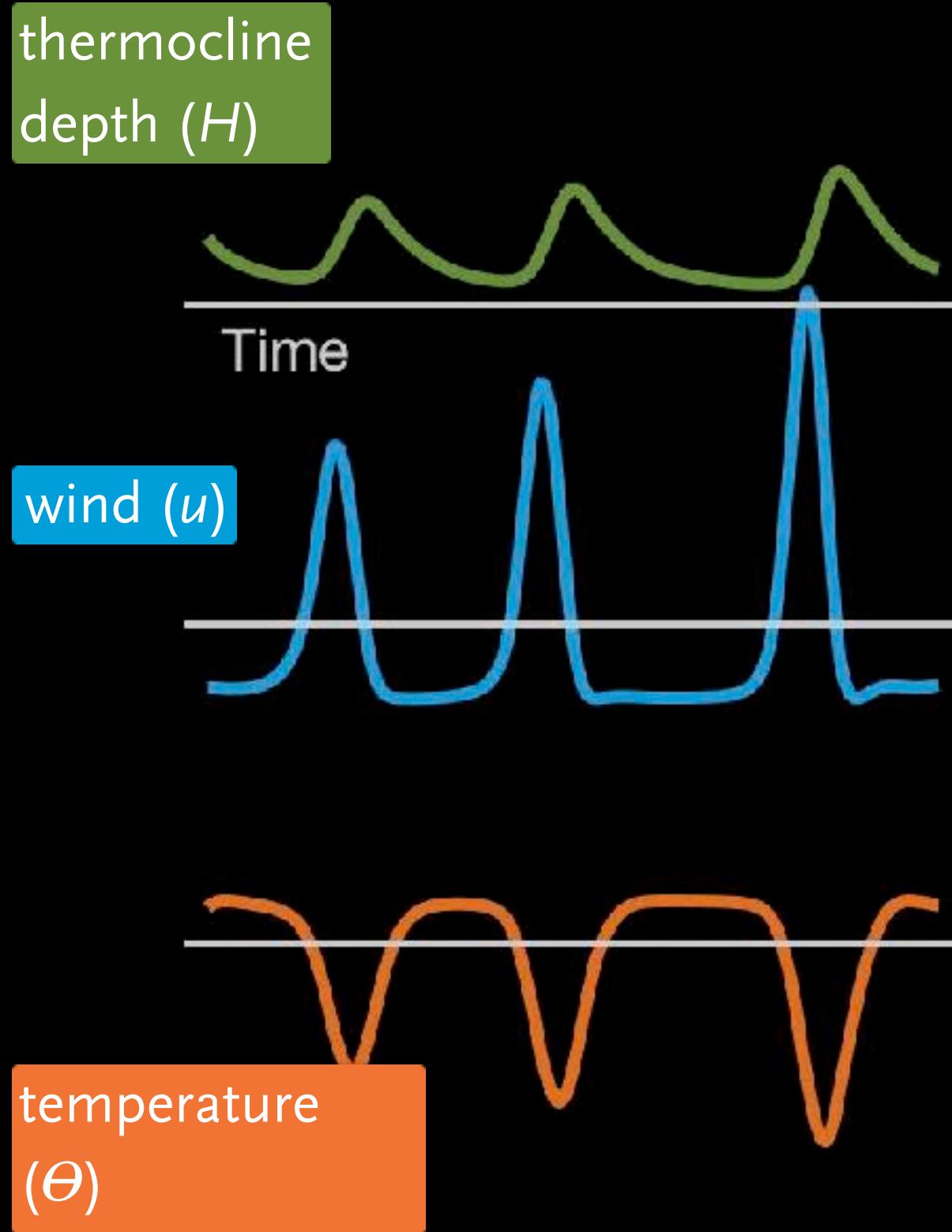


- State of the system:  
3 state variables (simplified)  
 $\Theta, H, u$
- Describe the system state at each time point
- Used to model the dynamics

# PHASE SPACE



# PHASE SPACE

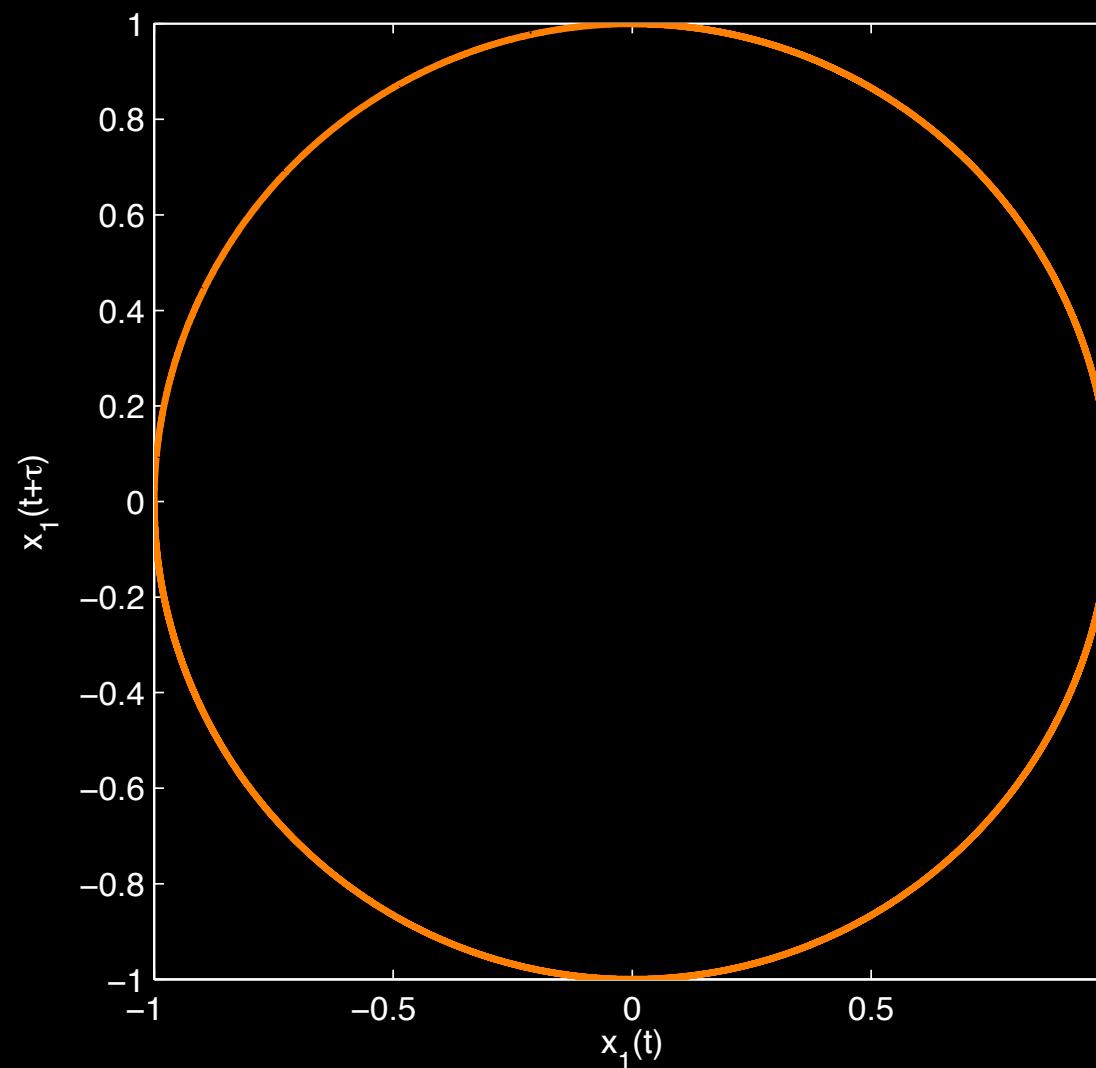
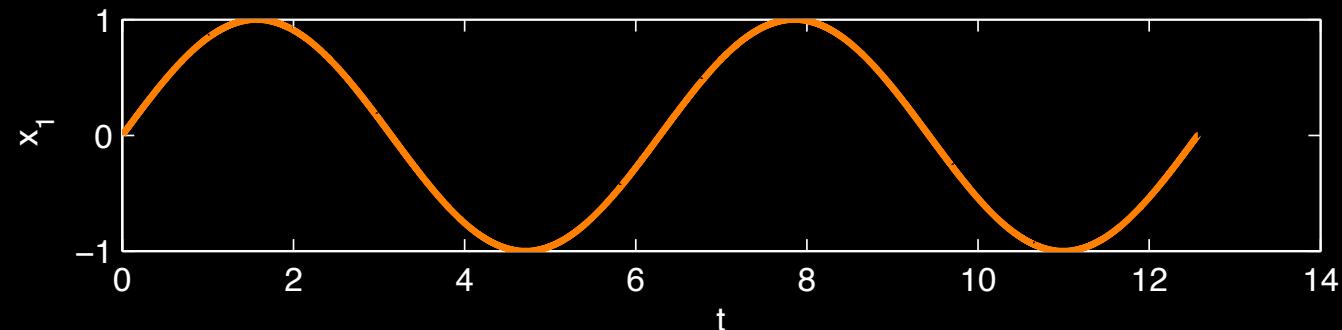


# PHASE SPACE RECONSTRUCTION

- However:  
often only one variable measured (or available)
- Because:  
not accessible, not known, too expensive to measure it ...

# METHODS FOR PHASE SPACE RECONSTRUCTION

- Auto-correlation, mutual information (time delay)
- False nearest neighbours (embedding dimension)
- More advanced:  
Cao, Pecora, Nichkawde,  
PECUZAL, ...





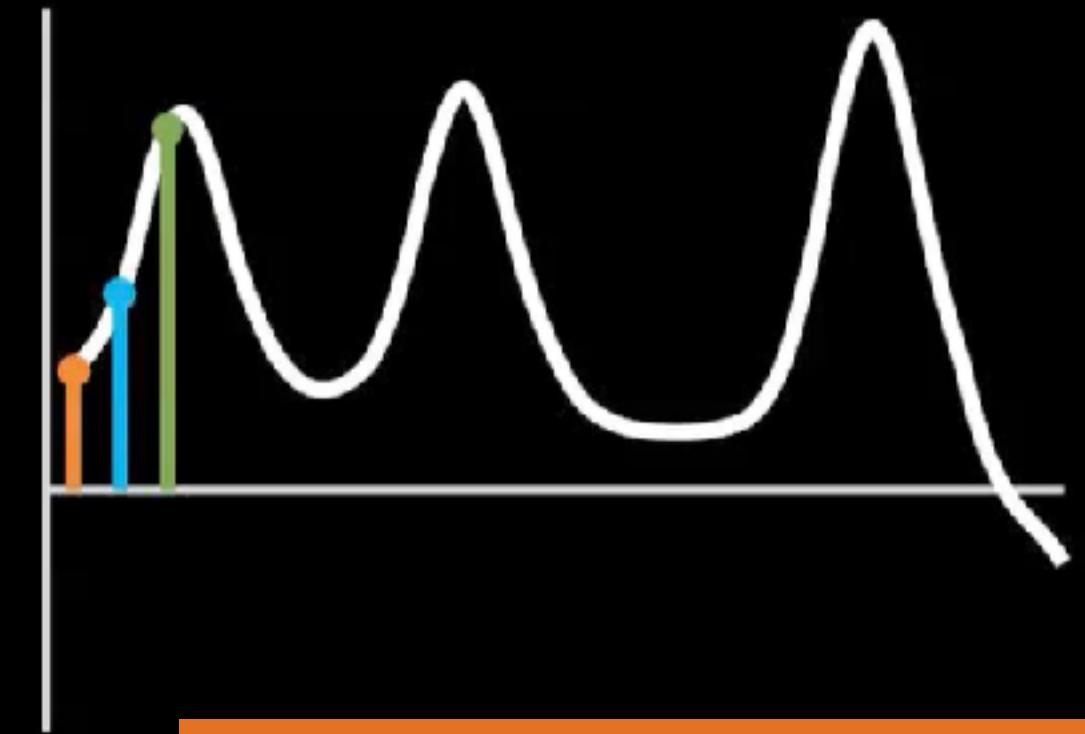
# TIME DELAY EMBEDDING

- Packard et al, 1980; Takens, 1981
- Use time shifted copies of the time series

$$\vec{x}(i) = (u_i, u_{i+\tau}, u_{i+2\tau}, \dots, u_{i+(m-1)\tau})$$

↑  
time delay      ↑  
embedding dimension

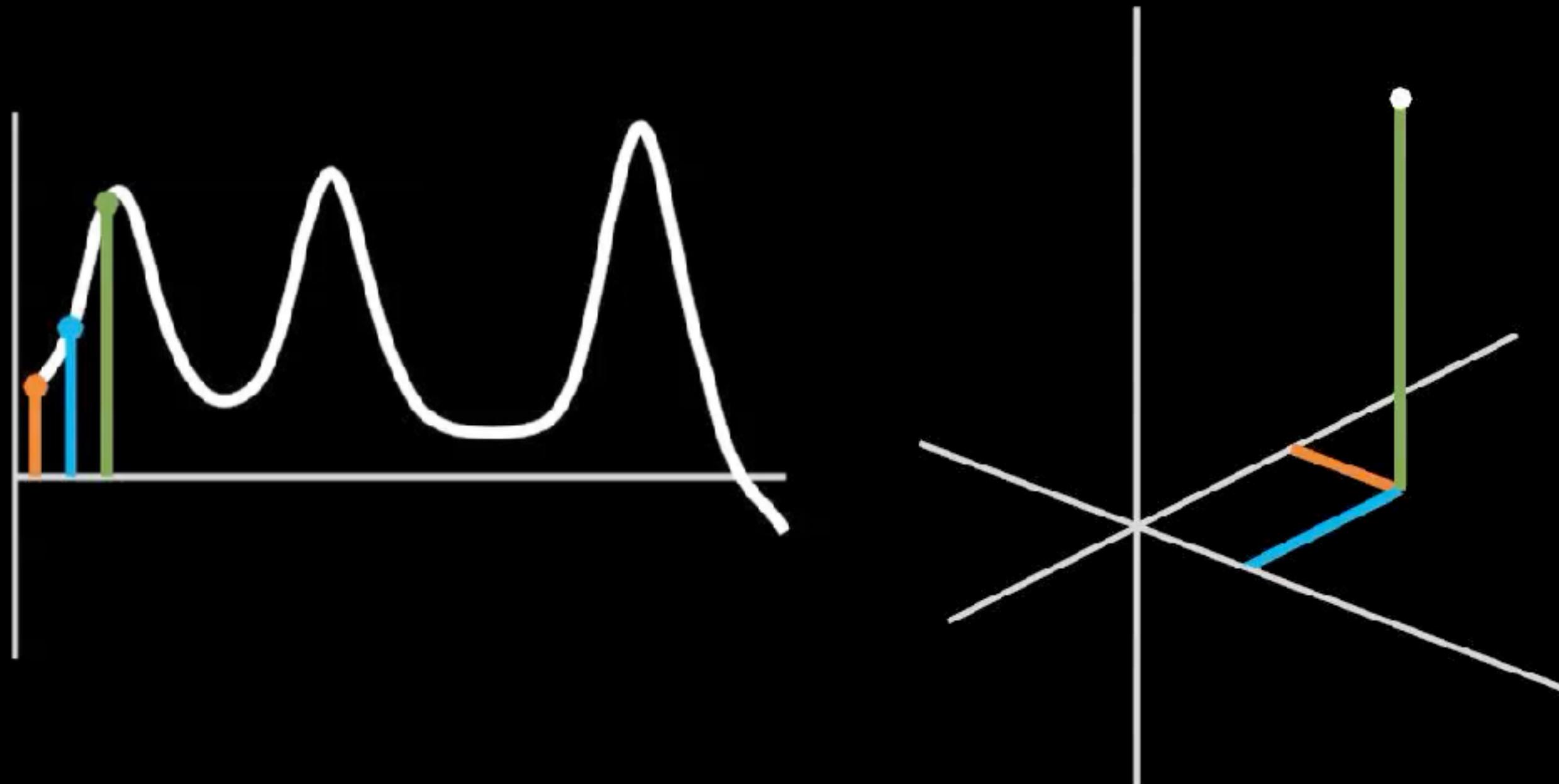
Note: Phase space trajectory is shorter by  $(m-1)\tau$



positive/ negative delay:  
check carefully with your  
research question

# TIME DELAY EMBEDDING

$$\vec{x}(i) = (u_i, u_{i+\tau}, u_{i+2\tau}, \dots, u_{i+(m-1)\tau})$$



# EMBEDDING DELAY

$$\vec{x}(i) = (u_i, u_{i+\tau}, \downarrow u_{i+2\tau}, \dots, u_{i+(m-1)\tau})$$

Independent components

## AUTOCORRELATION

$$C(\tau) = \frac{\sum_{t=1}^{N-\tau} (x_t - \bar{x})(x_{t+\tau} - \bar{x})}{(N - \tau)\sigma^2}$$

clear cycles:

first zero crossing

no clear cycles:

decorrelation time

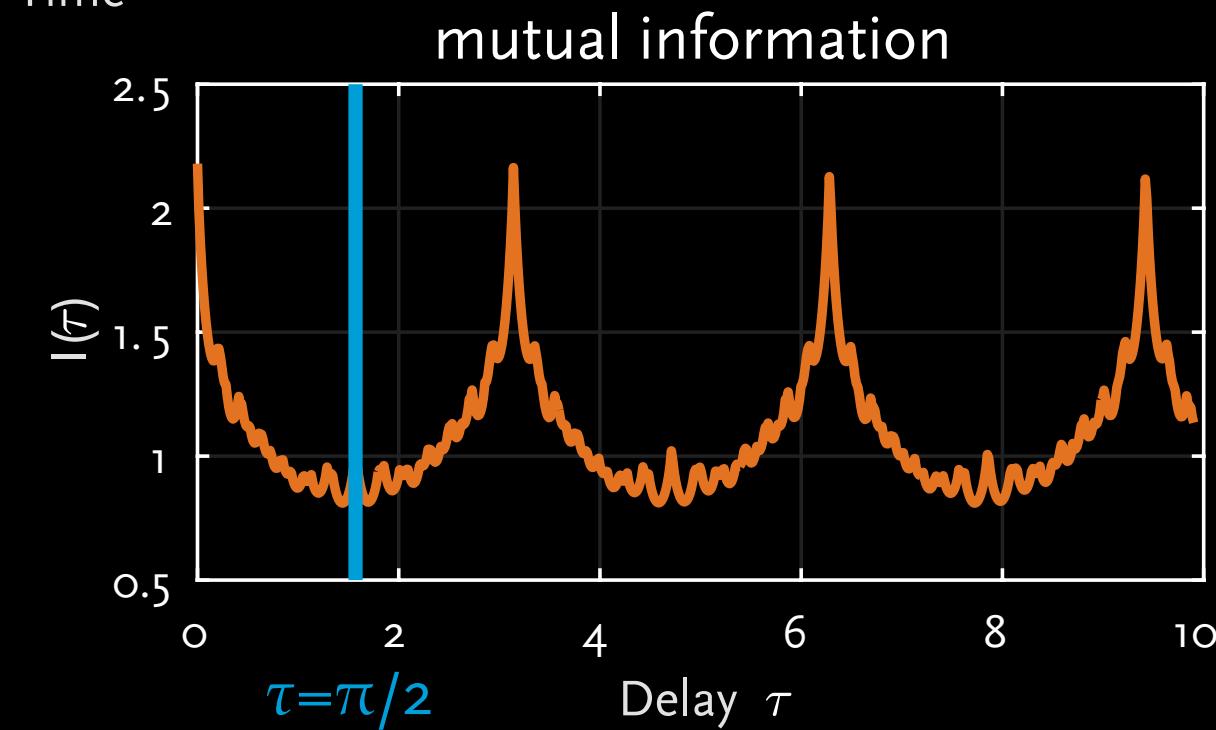
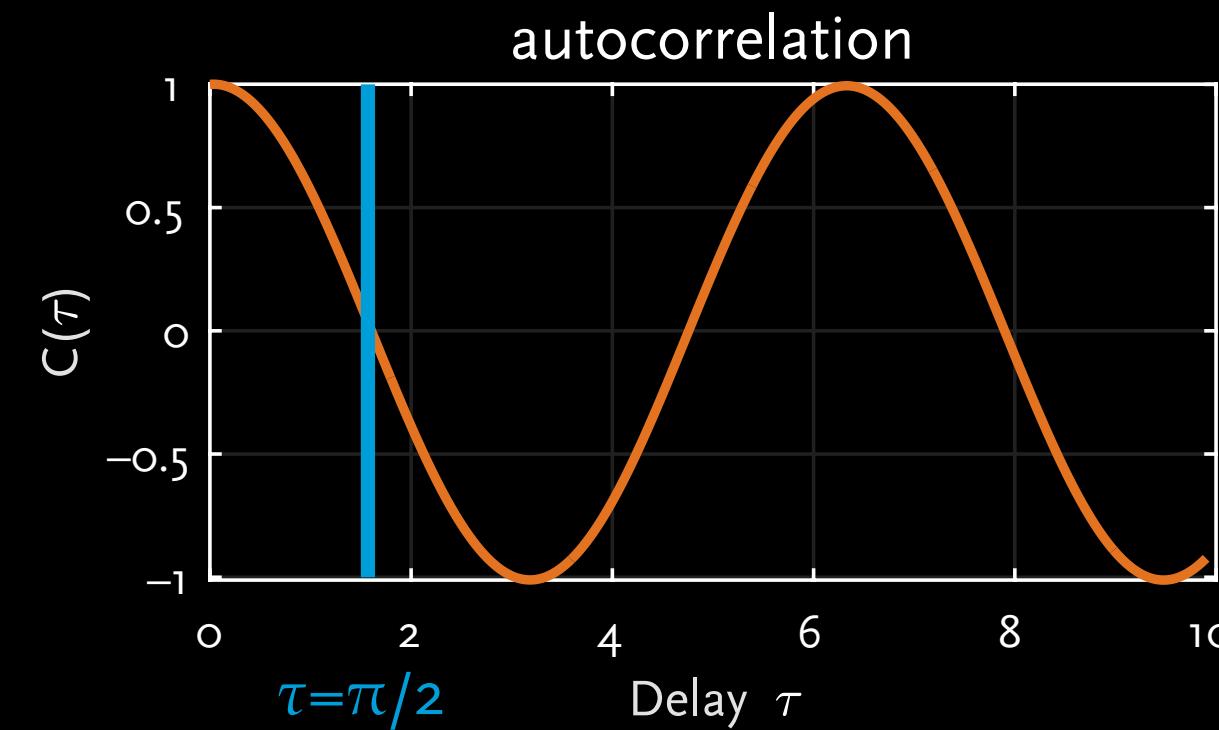
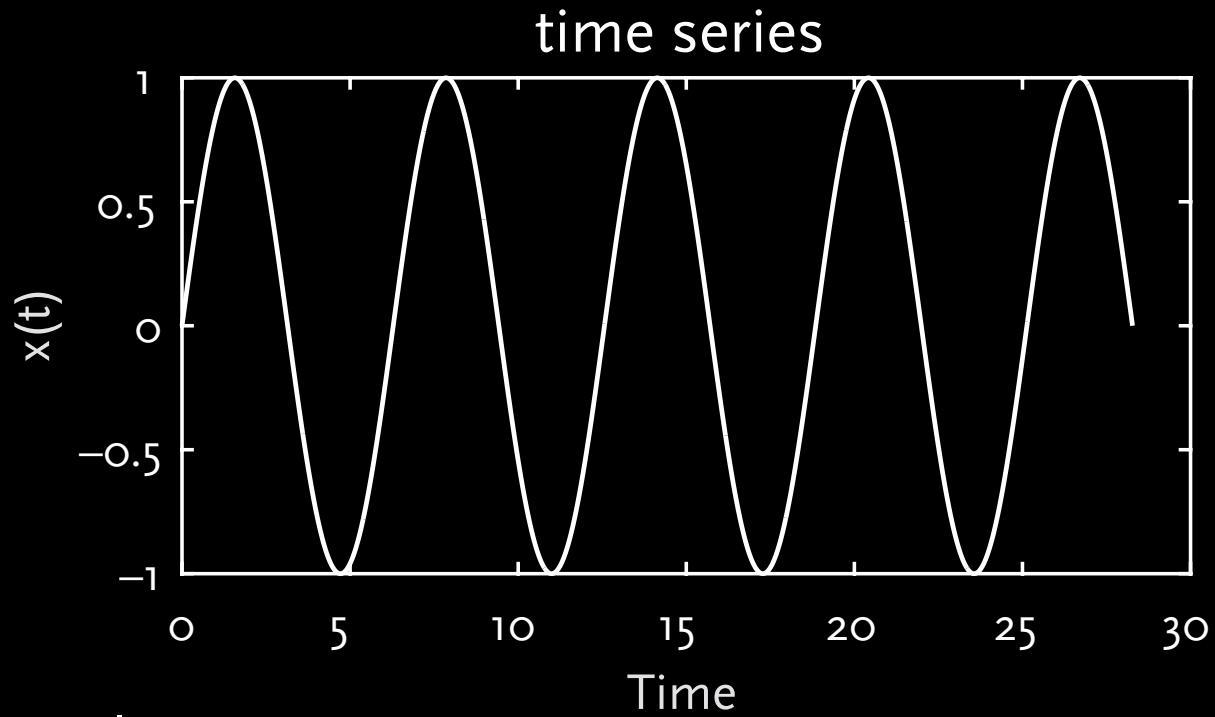
## MUTUAL INFORMATION

$$I(\tau) = \sum_x \sum_x p(x_t, x_{t+\tau}) \log \left( \frac{p(x_t, x_{t+\tau})}{p(x_t)p(x_{t+\tau})} \right)$$

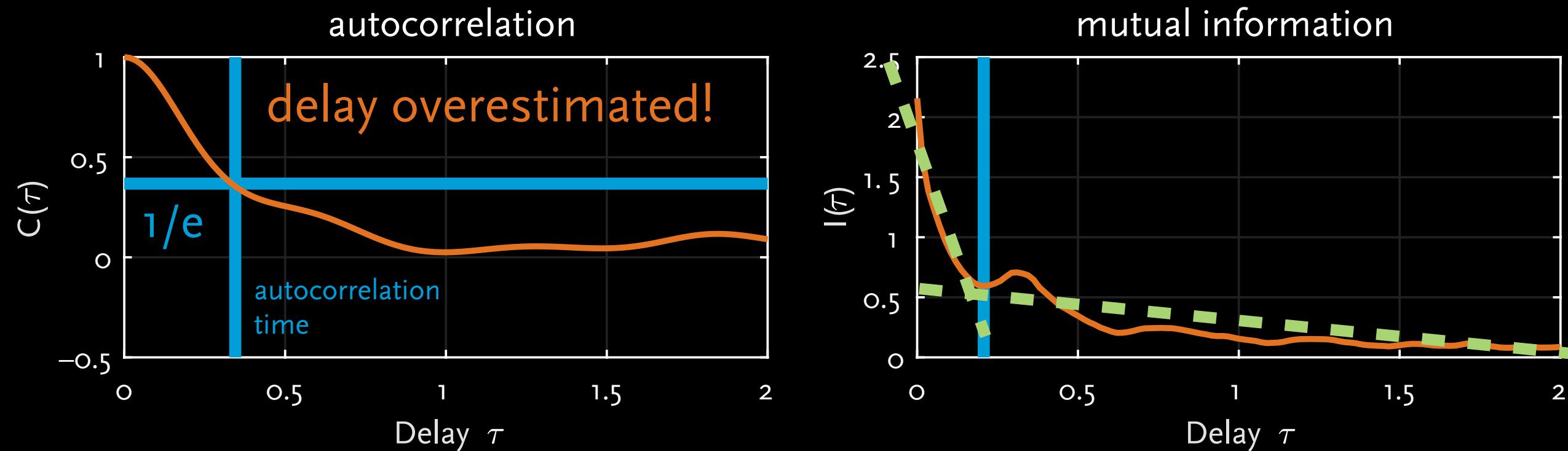
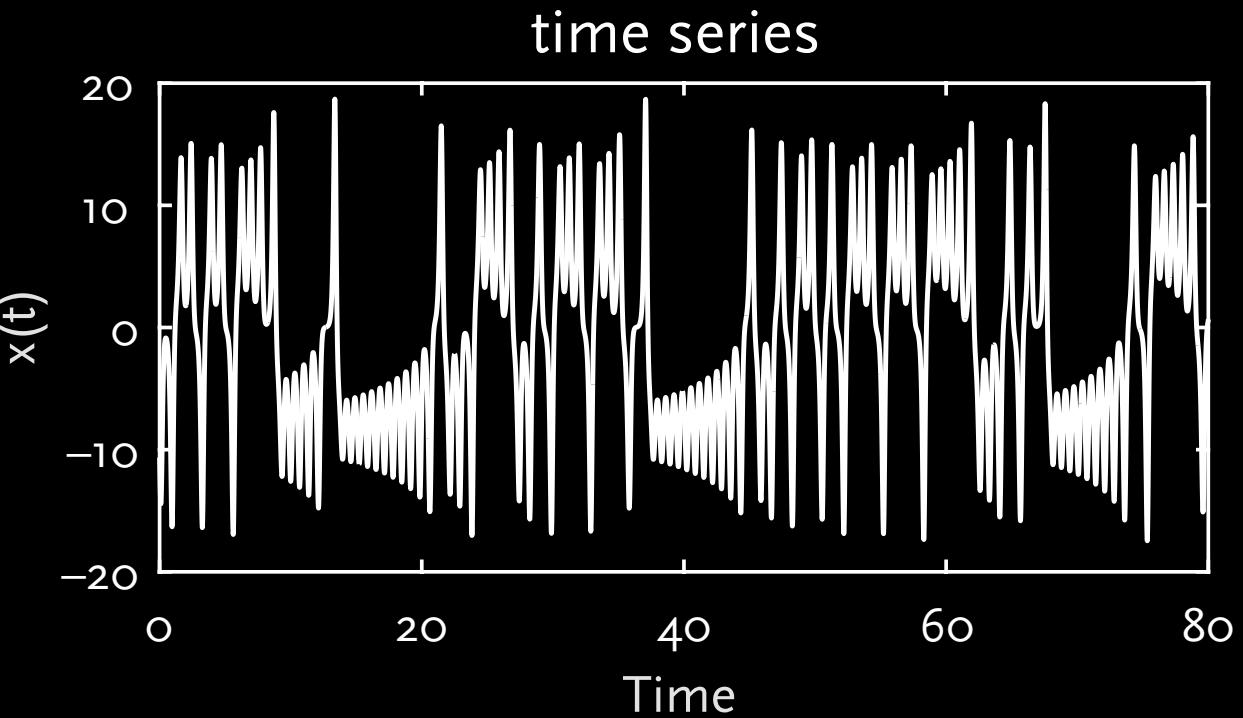
first minimum

point where steep decrease  
changes to flat decrease

# EMBEDDING DELAY



# EMBEDDING DELAY



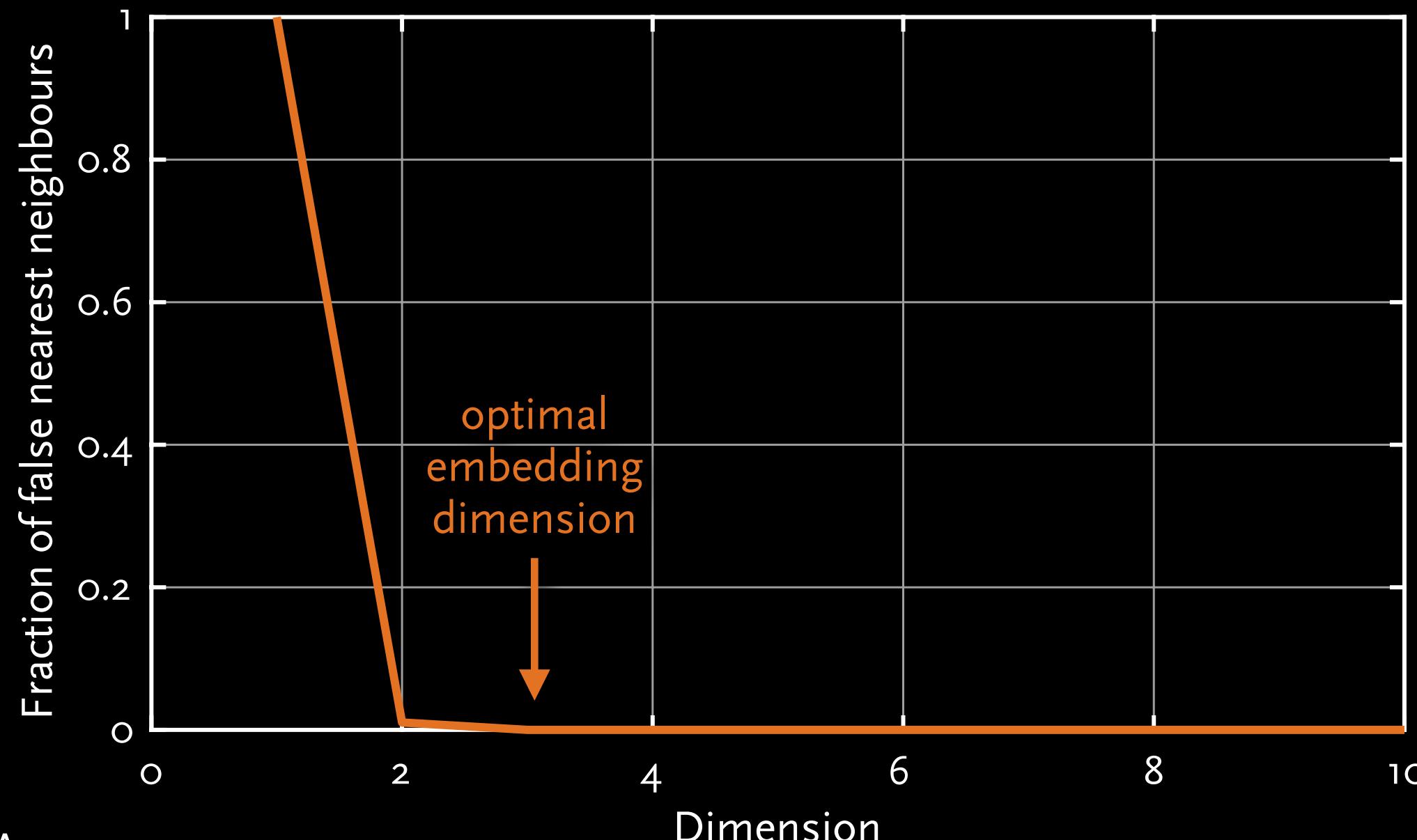
# EMBEDDING DIMENSION

$$\vec{x}(i) = (u_i, u_{i+\tau}, u_{i+2\tau}, \dots, u_{i+(m-1)\tau})$$

- Embedding dimension = number of components of phase space vector
- Aim: unfold the attractor
- Too high = state space becomes more sparse; increases correlations between phase space vectors
- False nearest neighbours (Kennel et al., 1992)

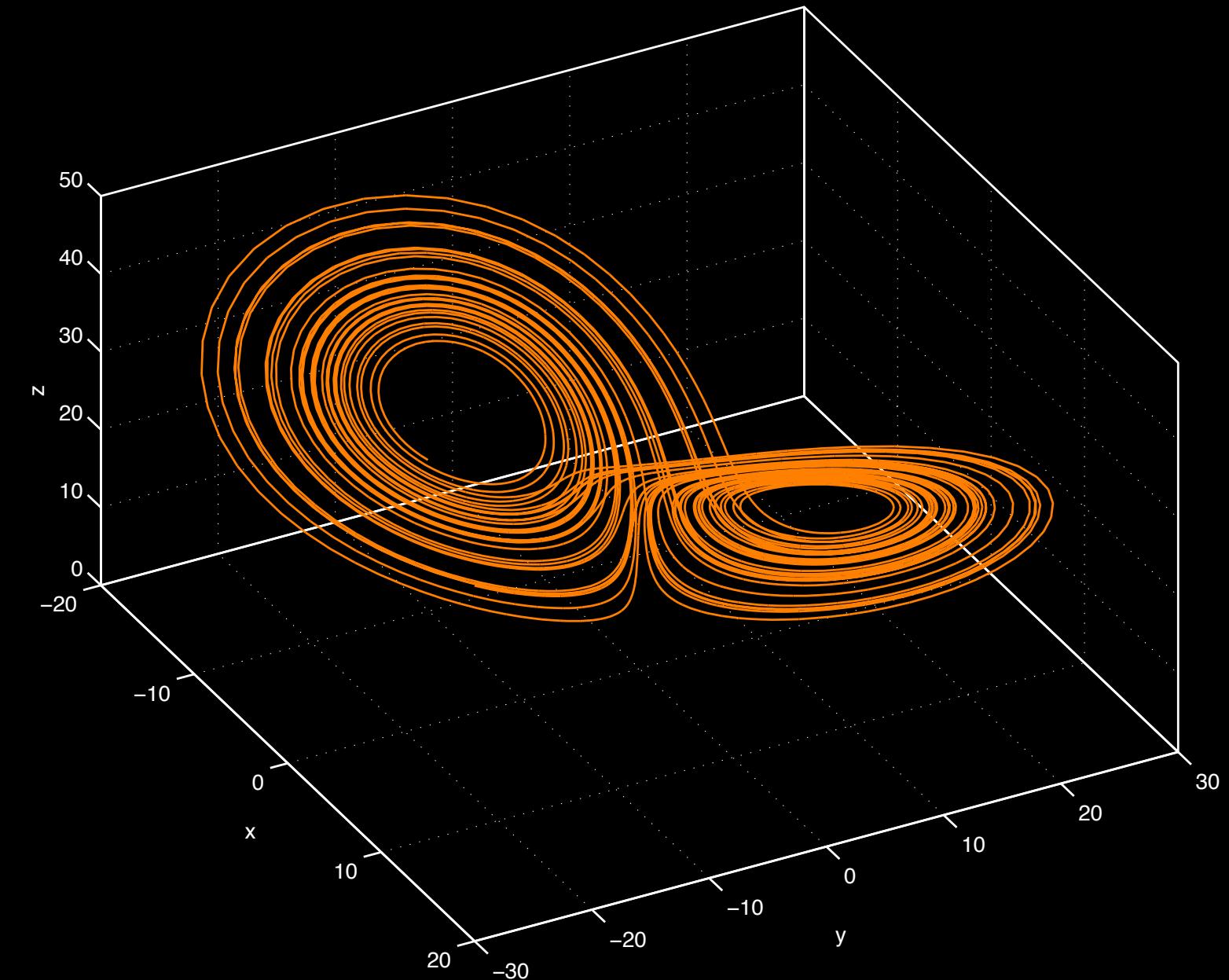
# FALSE NEAREST NEIGHBOURS

- Embedding dimension:  $m$  where fraction of false neighbours vanishes



# (RECONSTRUCTED) PHASE SPACE

- Representation of the system's variables (= dimension)
- Trajectory represents the dynamics of the system
- Many properties of the dynamical system can be derived from the phase space trajectory



# VISUAL INTERPRETATION

# RECURRENCE PLOT

- To visualise the phase space trajectory by its recurrences
- Recurrence matrix:
  - ▶ binary
  - ▶ symmetric

$$R_{i,j} = \begin{cases} 1 : \vec{x}_i \approx \vec{x}_j \\ 0 : \vec{x}_i \not\approx \vec{x}_j \end{cases}$$

$$R_{i,j} = \Theta(\varepsilon - \|\vec{x}_i - \vec{x}_j\|)$$

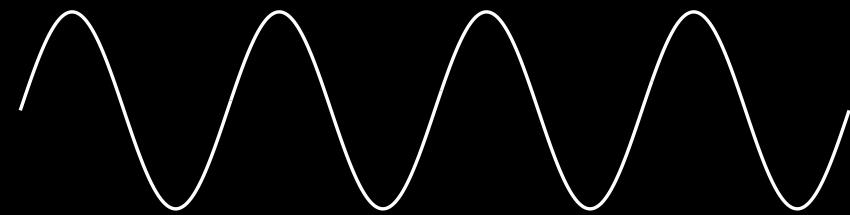
$$\Theta(x \geq 0) = 1; \Theta(x < 0) = 0$$

$$\mathbf{R} =$$

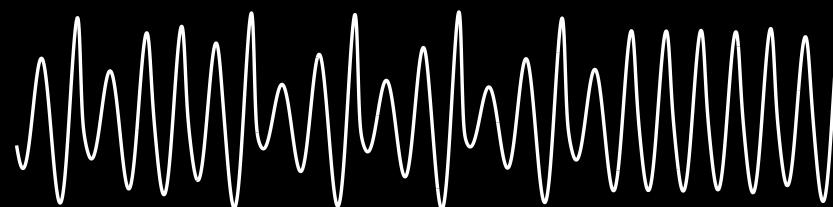
1	1	0	0	1
1	1	1	0	1
0	1	1	0	0
0	0	0	1	1
1	1	0	1	1

# DIFFERENT DYNAMICS: DIFFERENT PATTERNS

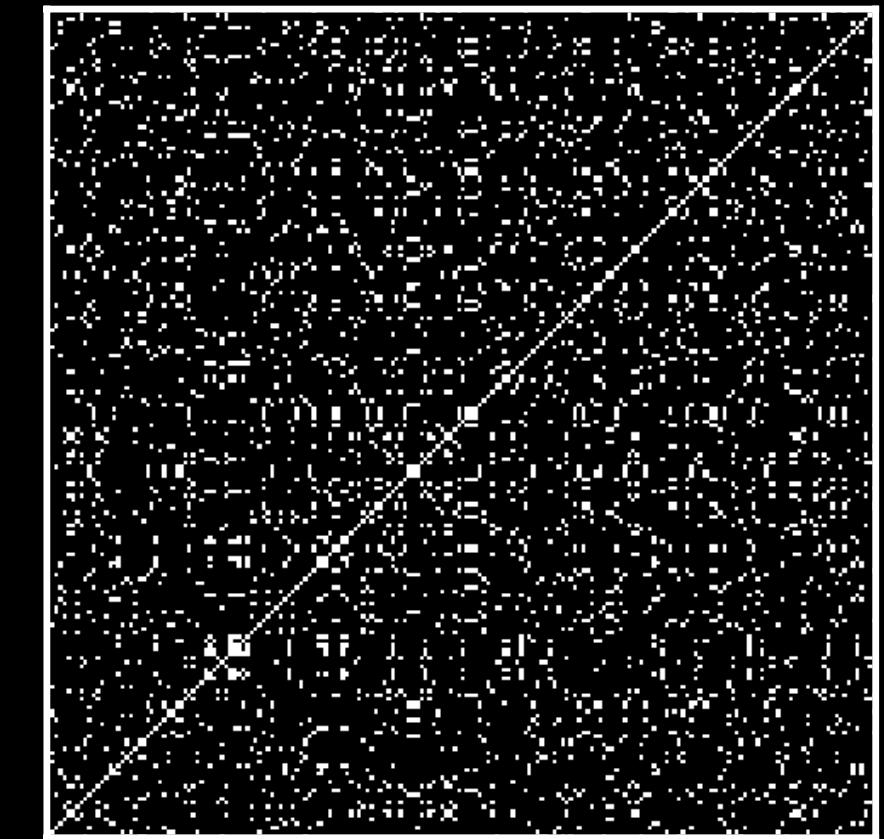
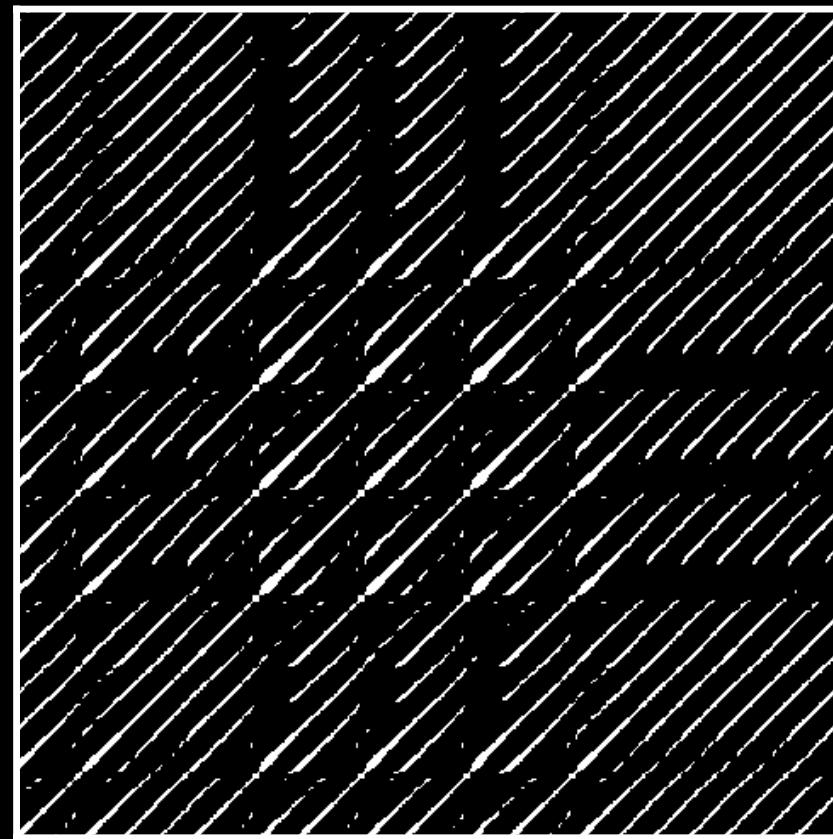
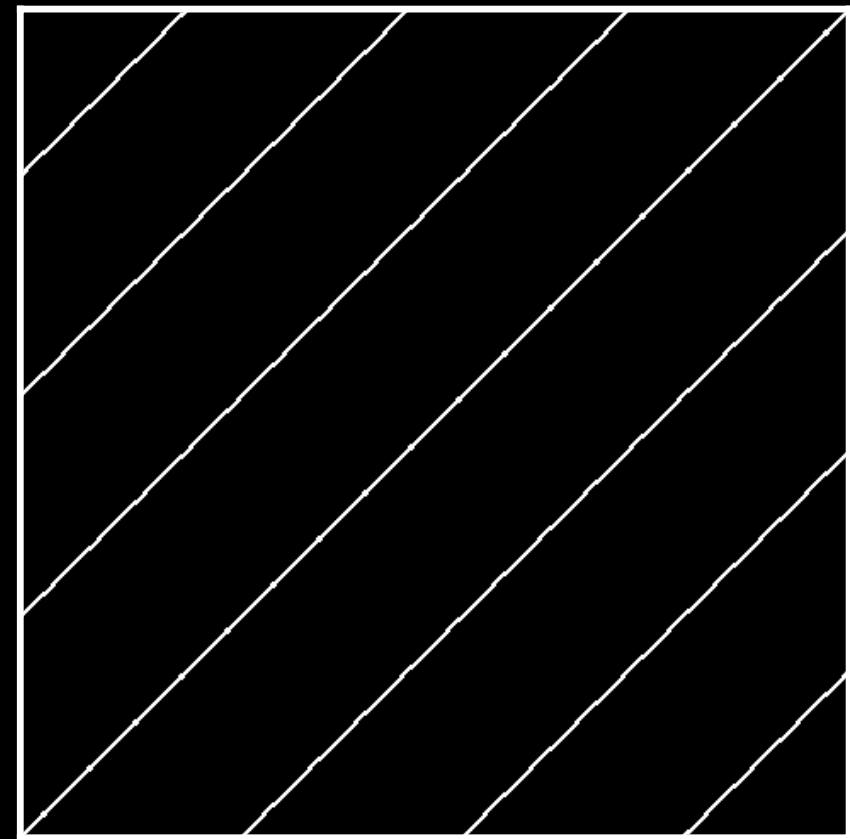
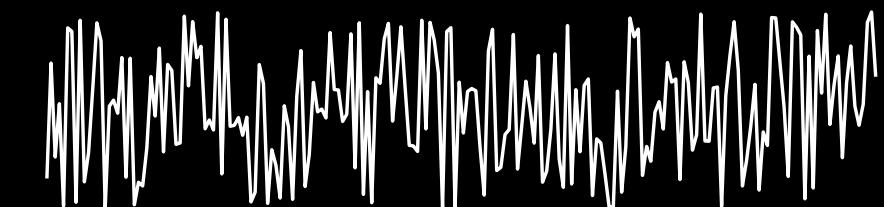
Periodic



Chaotic



Random



# VISUAL INTERPRETATION

Typology (large scale appearance)	homogeneous	periodic	trend	disrupted, separated into blocks
Texture (small scale structures)	only single points	mainly diagonal lines	vertical lines, black blocks	curved pattern
Characteristic pattern	noninterrupted diagonal lines, periodic separation of diagonal lines	sparingly occupied columns/ bars	missing recurrences	local changes in recurrence point density

# VISUAL INTERPRETATION

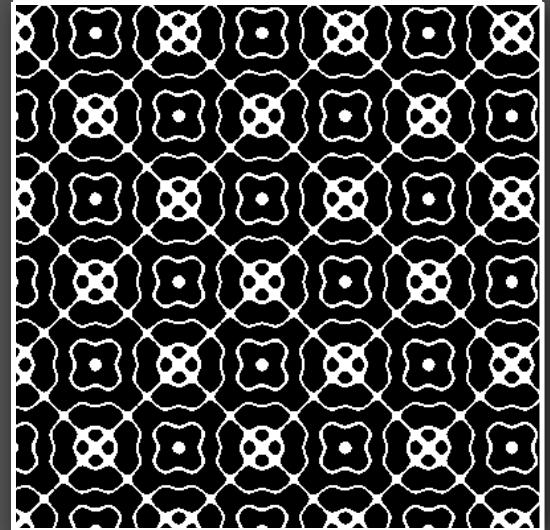
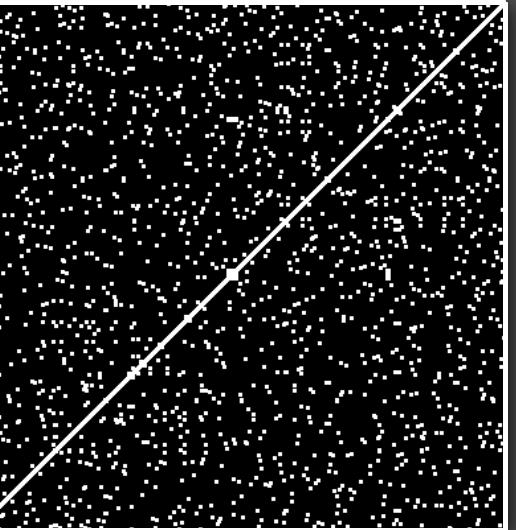
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# HOMOGENEOUS TYPOLOGY

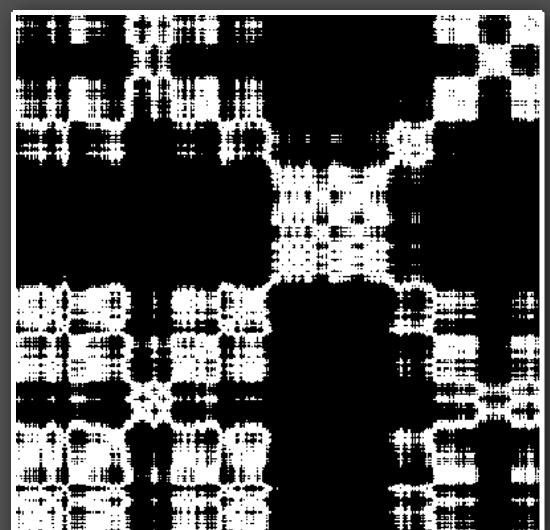
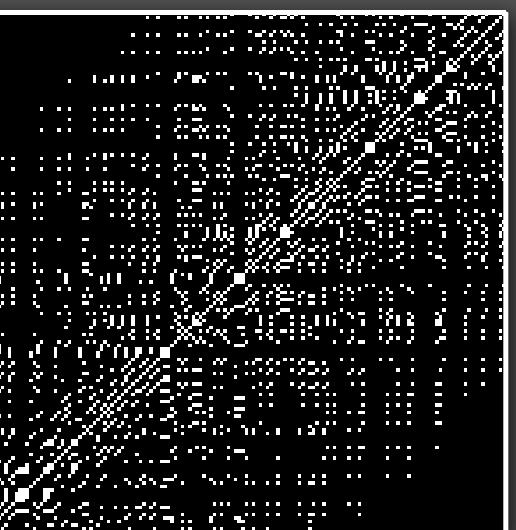
- Homogeneous:  
time series is stationary
- Heterogeneous:  
time series is not stationary  
(yet, process could be stationary!)

We distinguish:  
stationarity:  
nonstationarity:  
measures of distribution and autocorrelation change with  
changes in shifts ( $x_{t+s}$ )

homogeneous

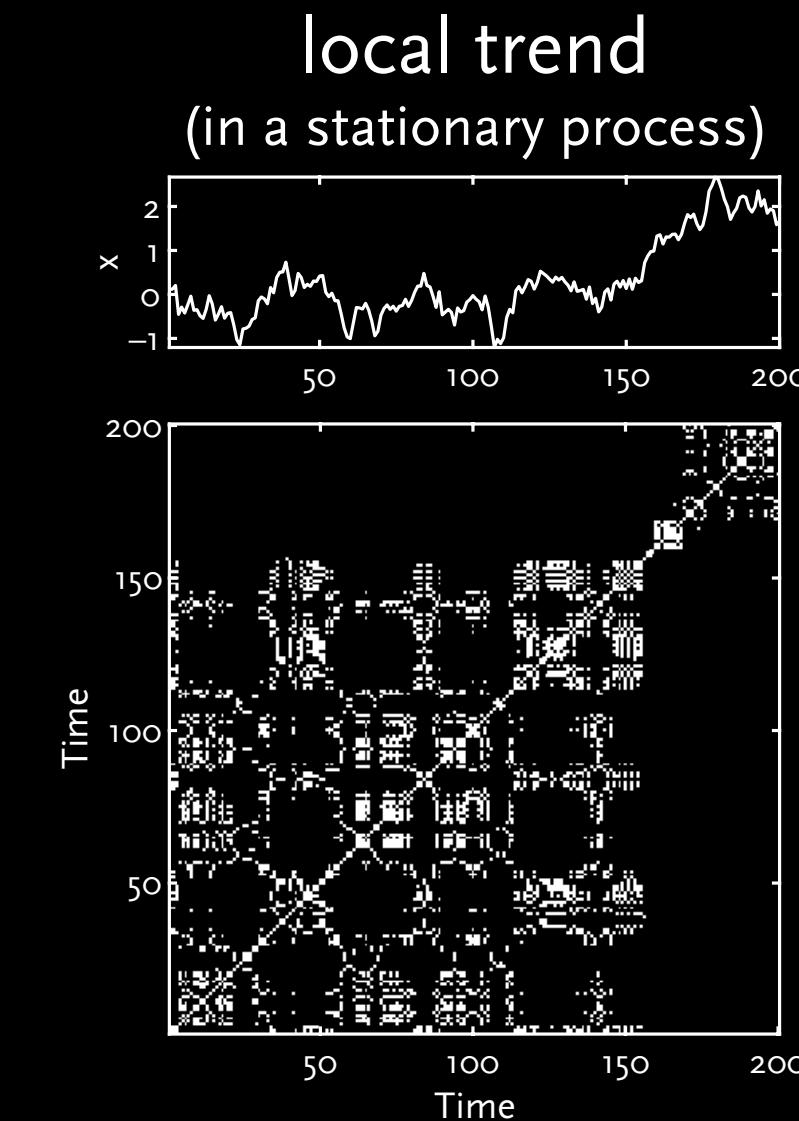
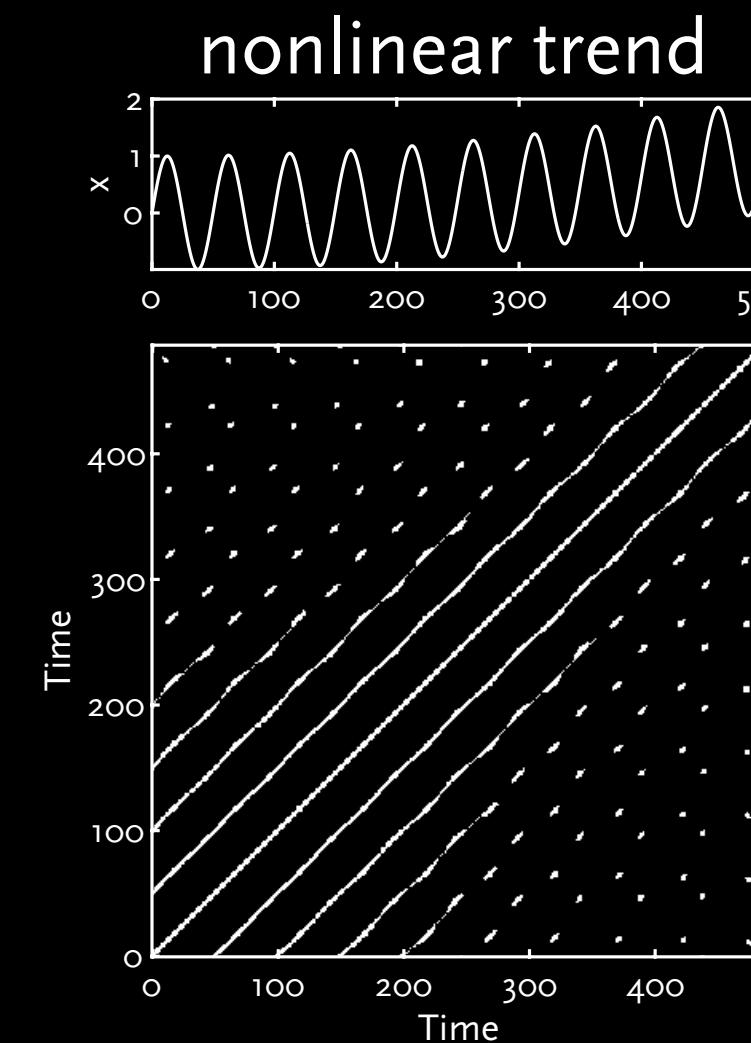
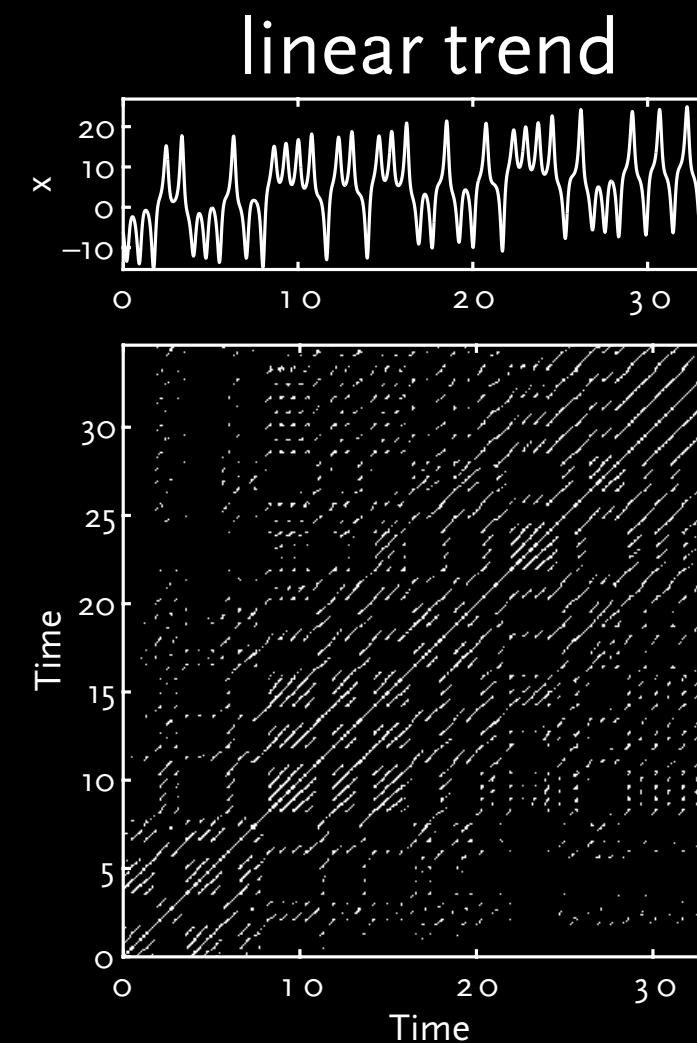


disrupted



# FADING TO THE UPPER LEFT AND LOWER RIGHT CORNERS

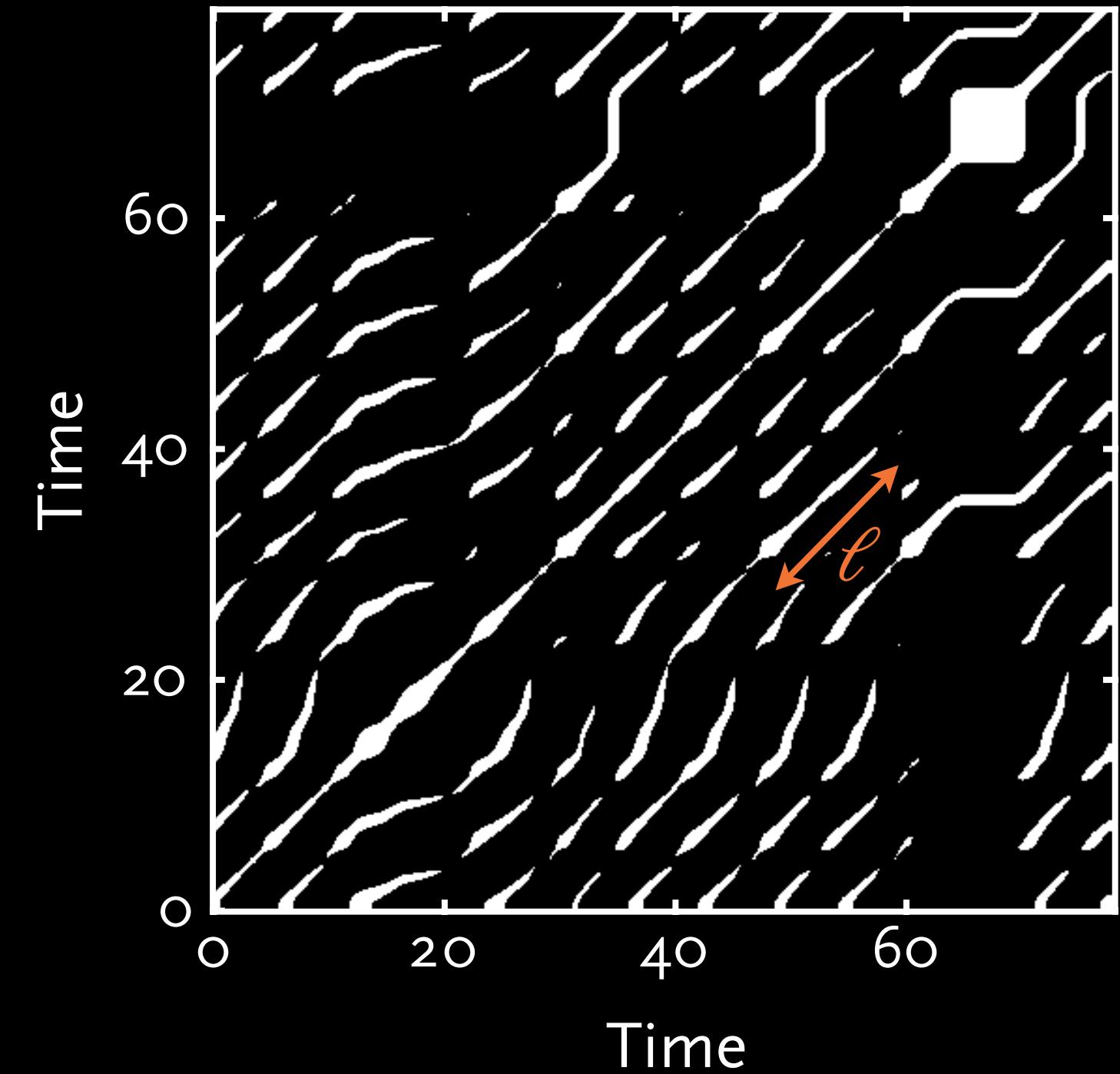
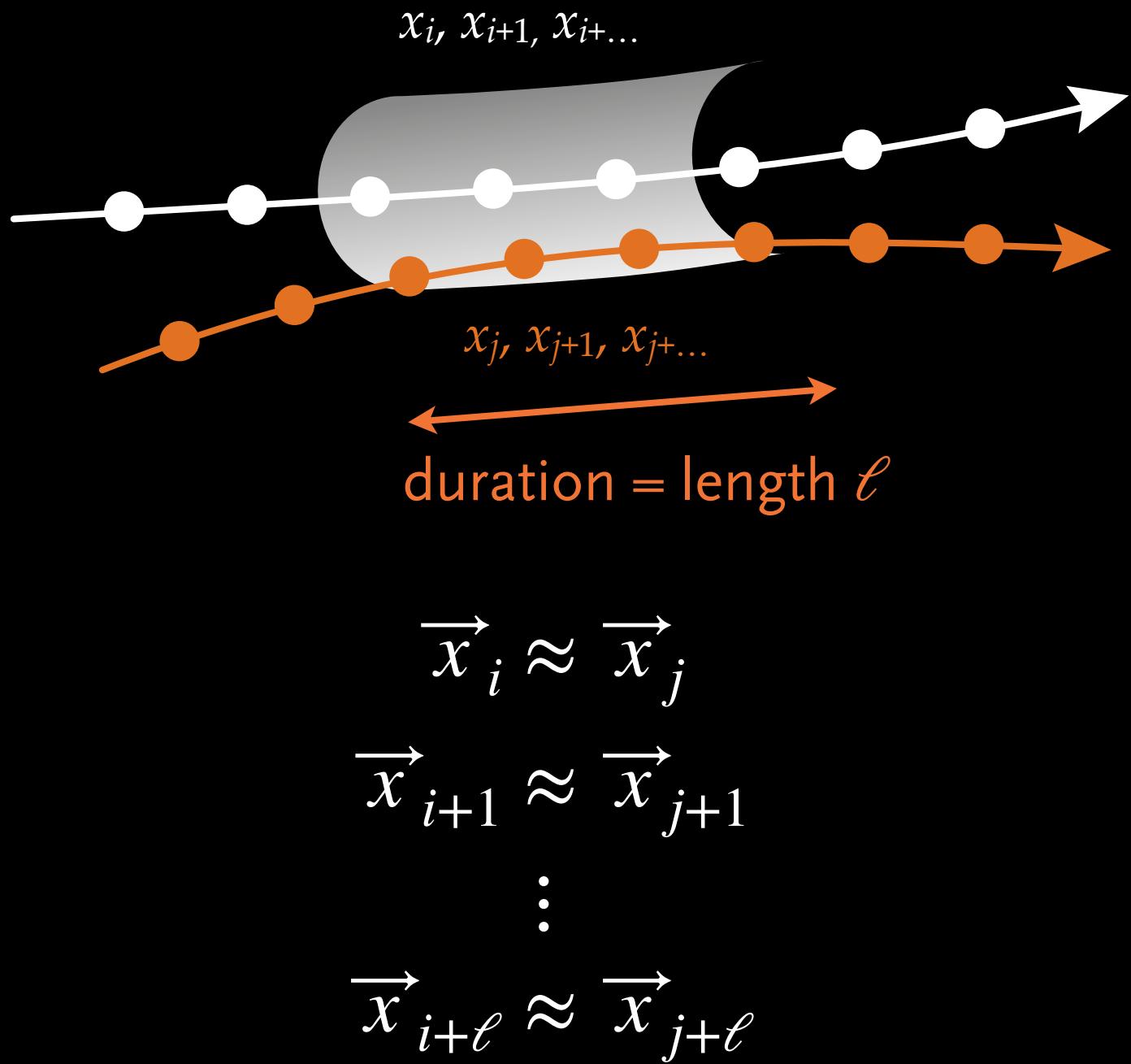
- Non-stationary data
- Process contains (at least a local) trend or shift



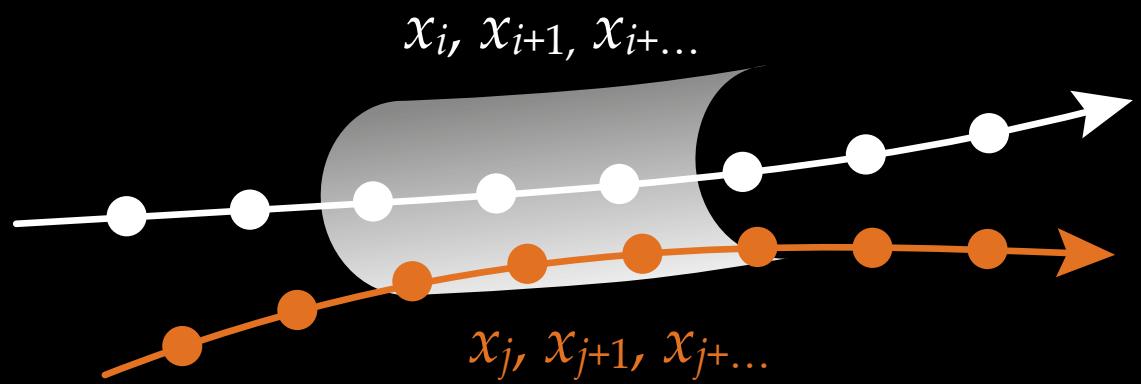
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# DIAGONAL LINES PARALLEL TO MAIN DIAGONAL



# DIAGONAL LINES PARALLEL TO MAIN DIAGONAL

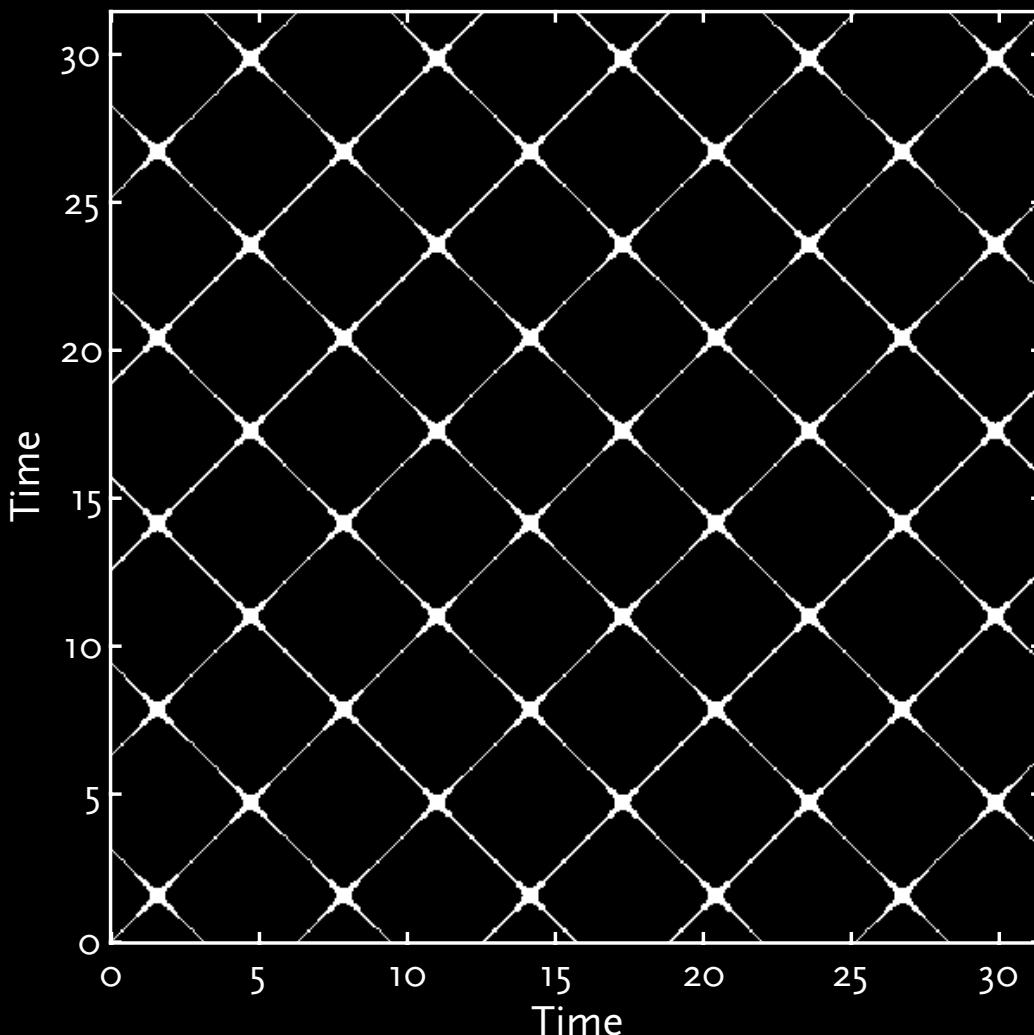


- Evolution of states is similar at different epochs
- Process could be deterministic
- Diagonal lines beside single isolated points: process could be chaotic
- Separation of diagonal lines periodic: unstable periodic orbits

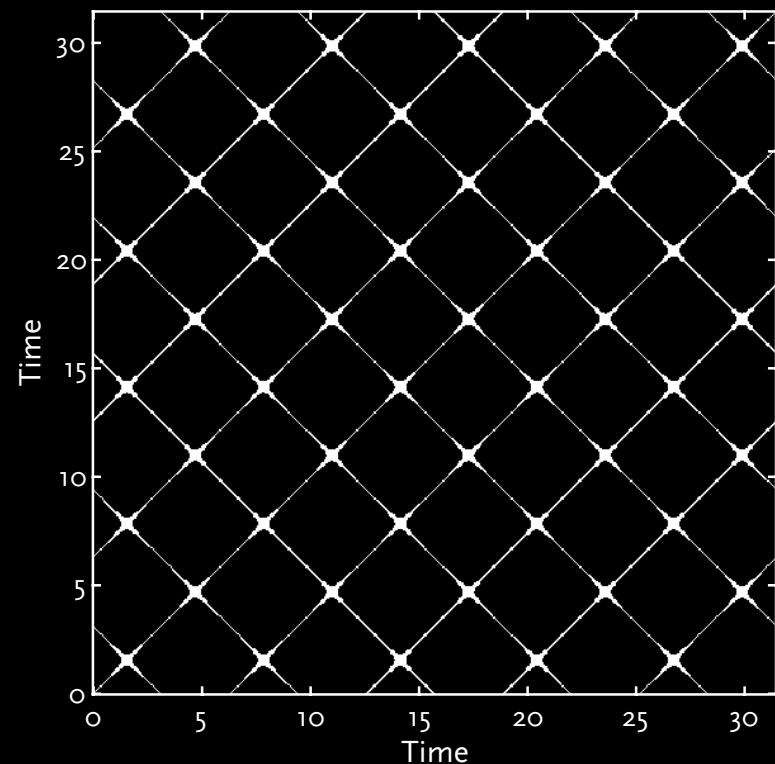
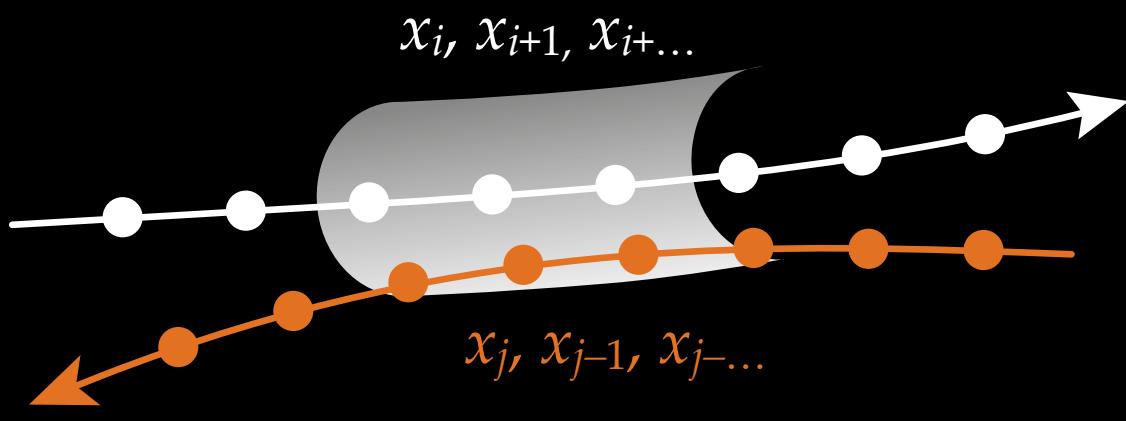
Q u i z Z

# Quiz

- What does the perpendicular diagonals mean?



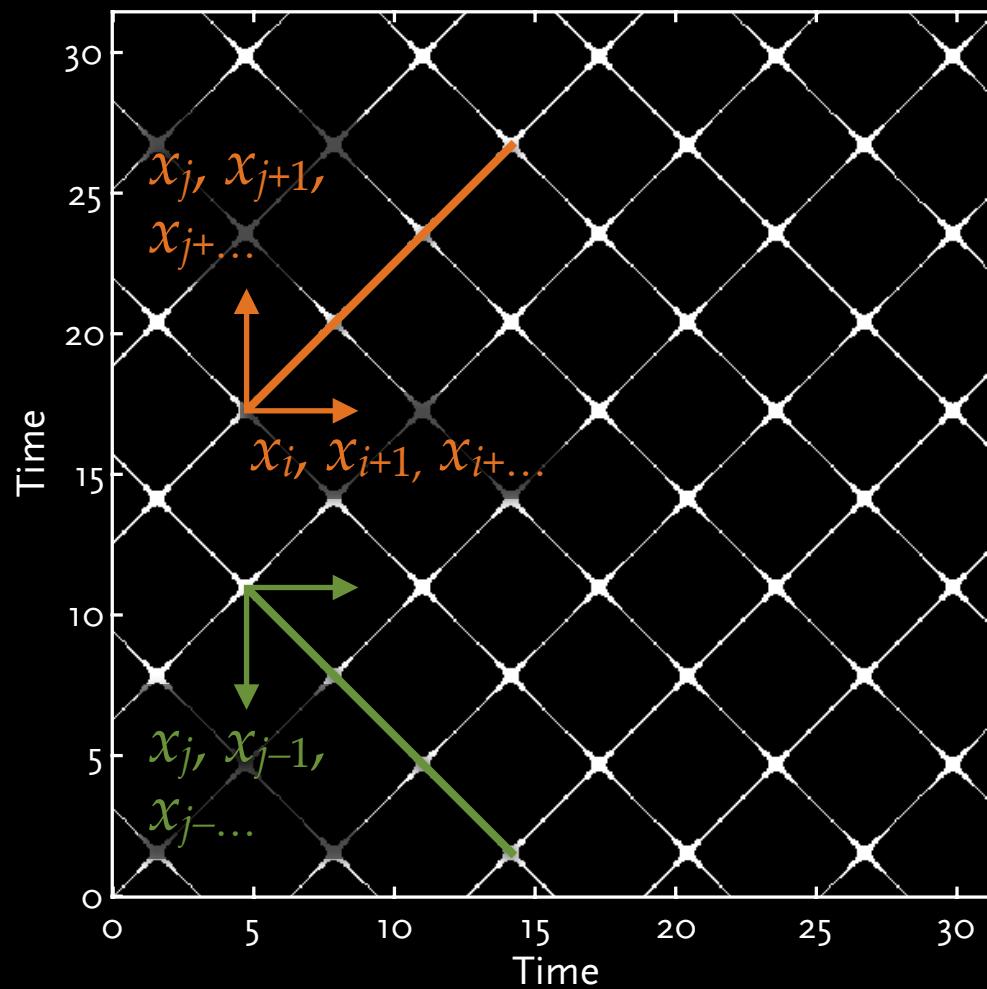
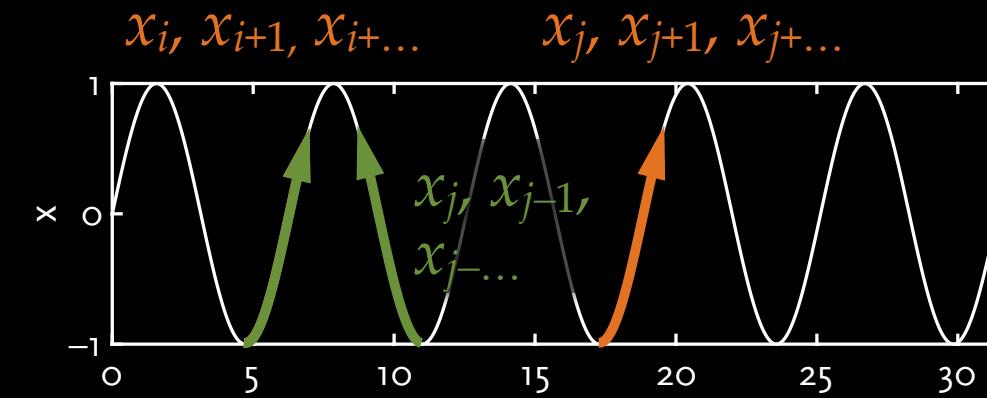
# DIAGONAL LINES ORTHOGONAL TO MAIN DIAGONAL



- Evolution of states is similar at different times but with reverse temporal order
- Indication for an insufficient embedding



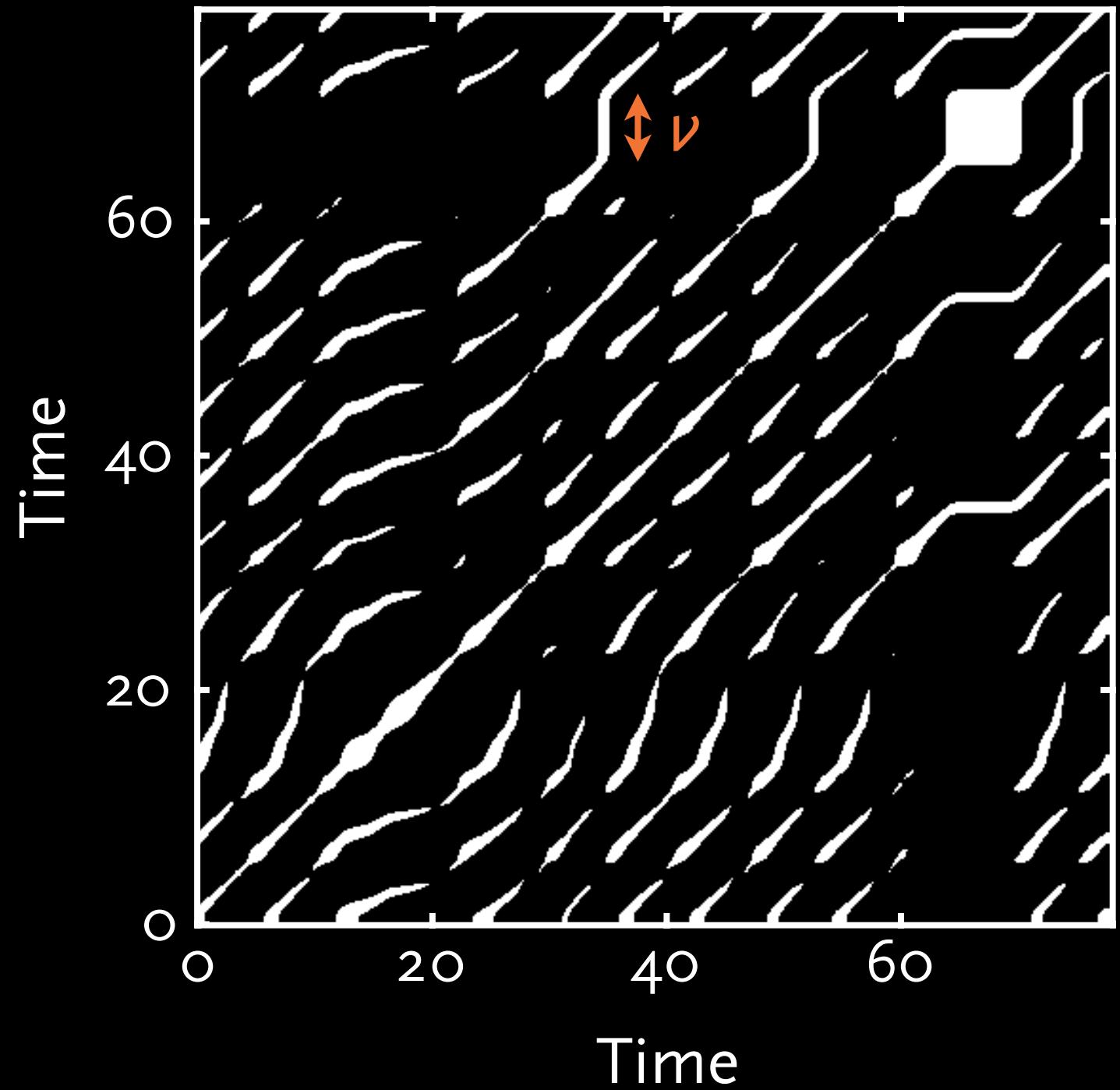
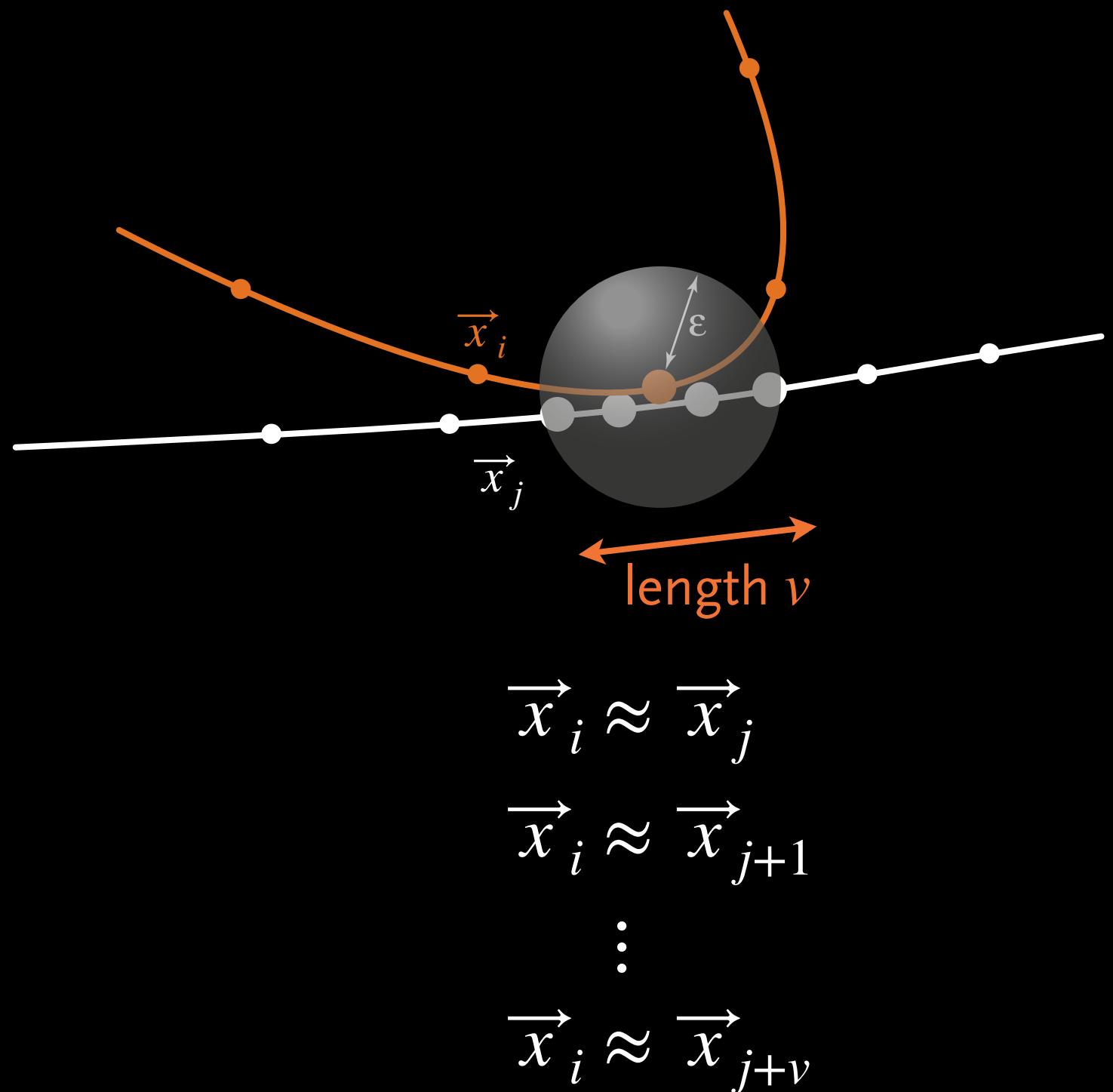
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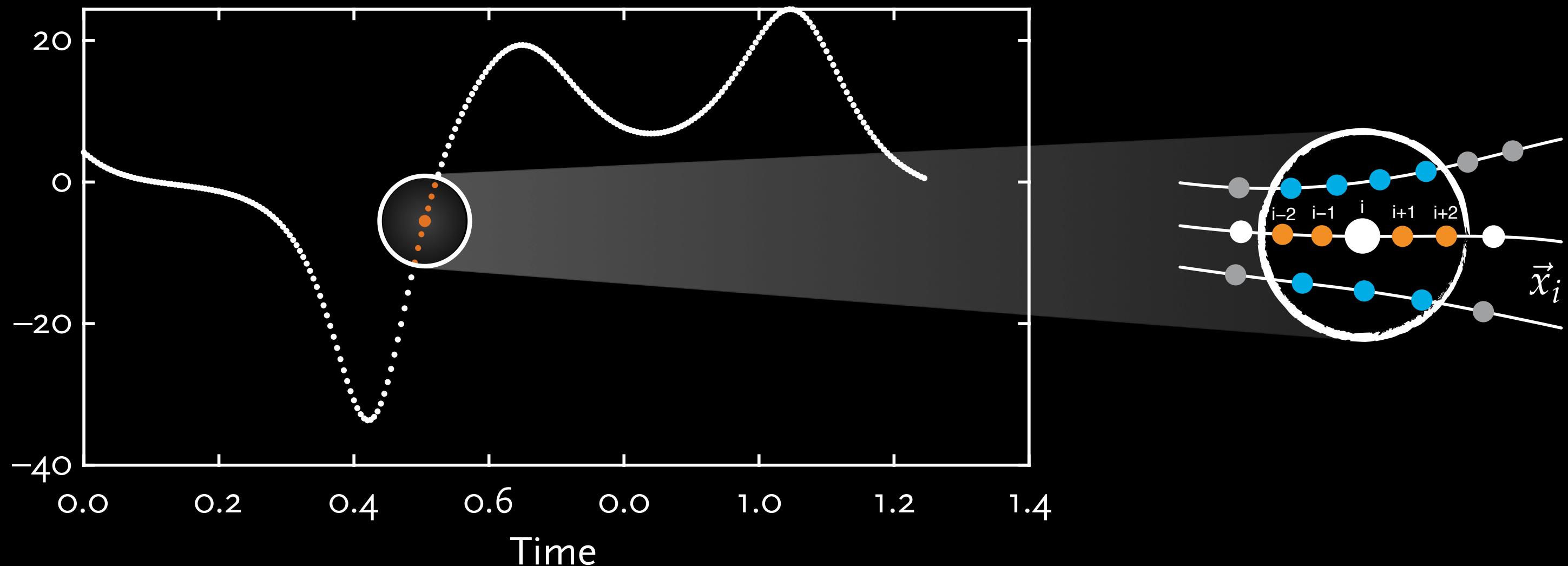


# VERTICAL LINES, BLOCKS



# VERTICAL LINES, BLOCKS

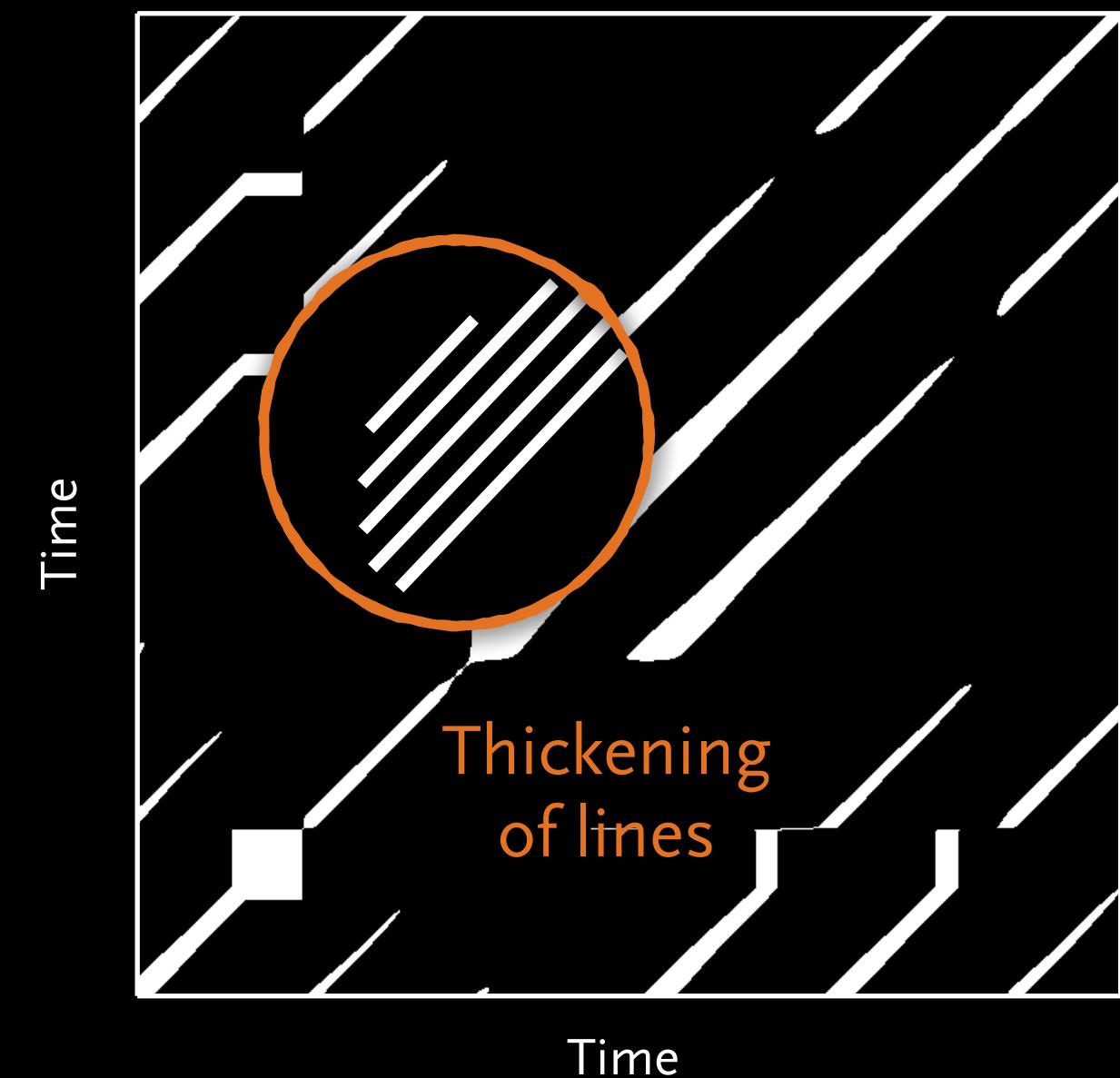
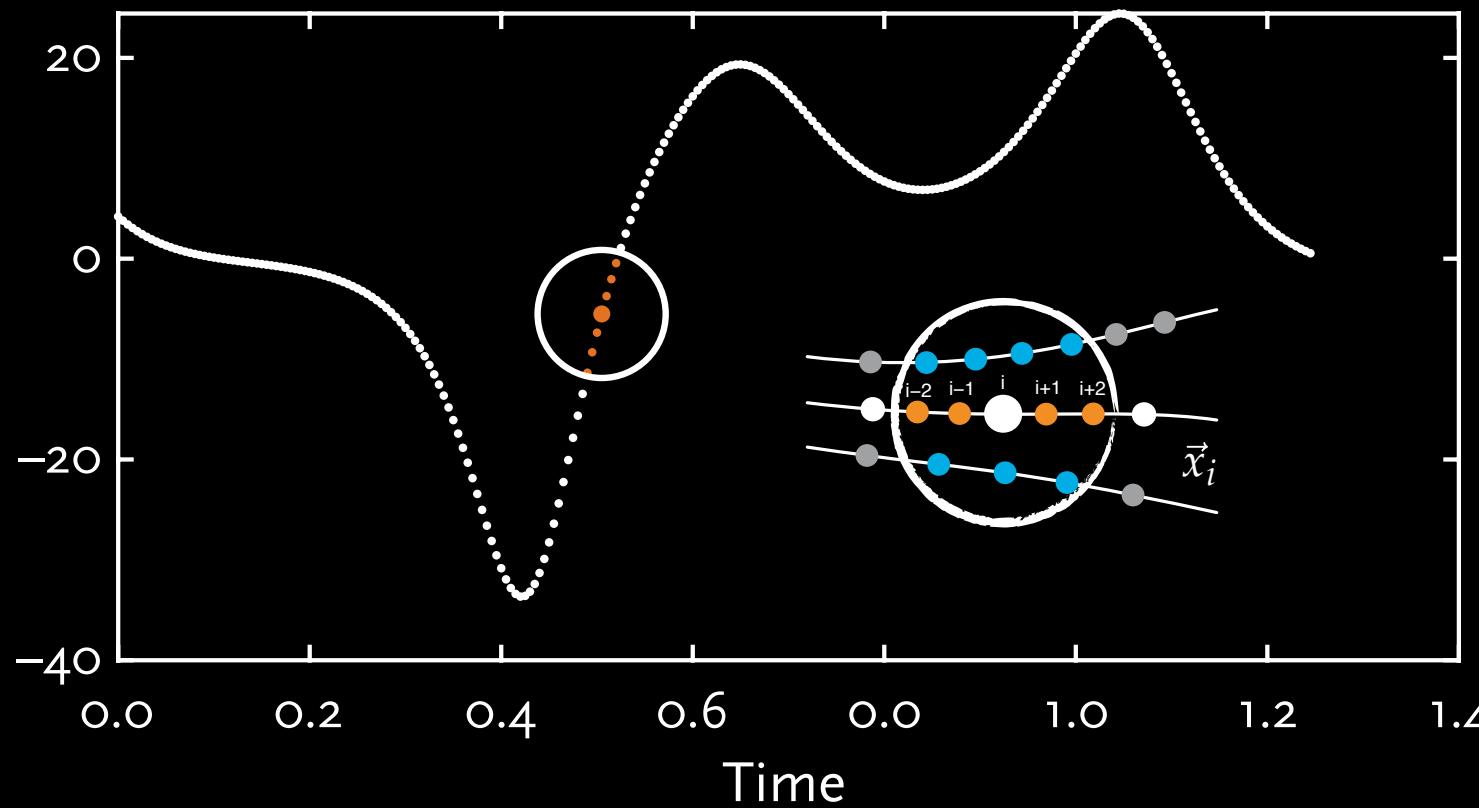
- Close in time in discretized time series or phase space



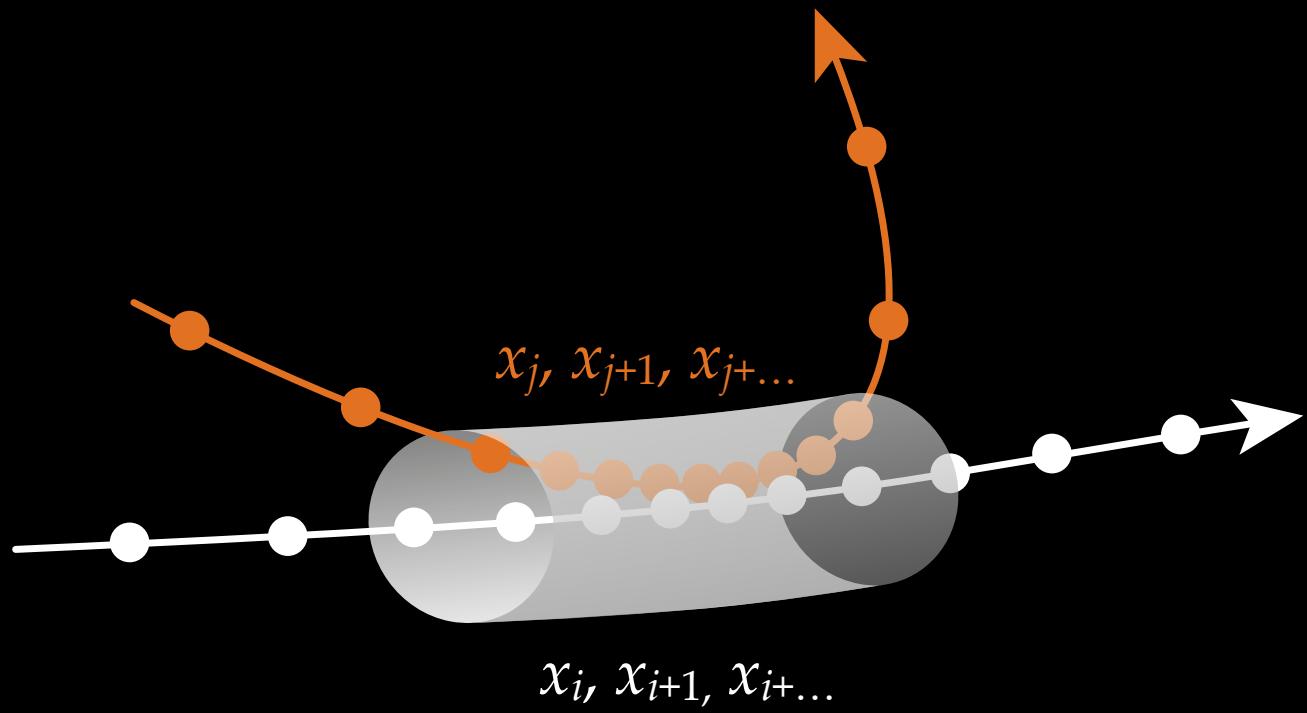


# VERTICAL LINES, BLOCKS

- Sojourn points (“tangential motion”)
- Affected by sampling time and “smoothness” of the signal



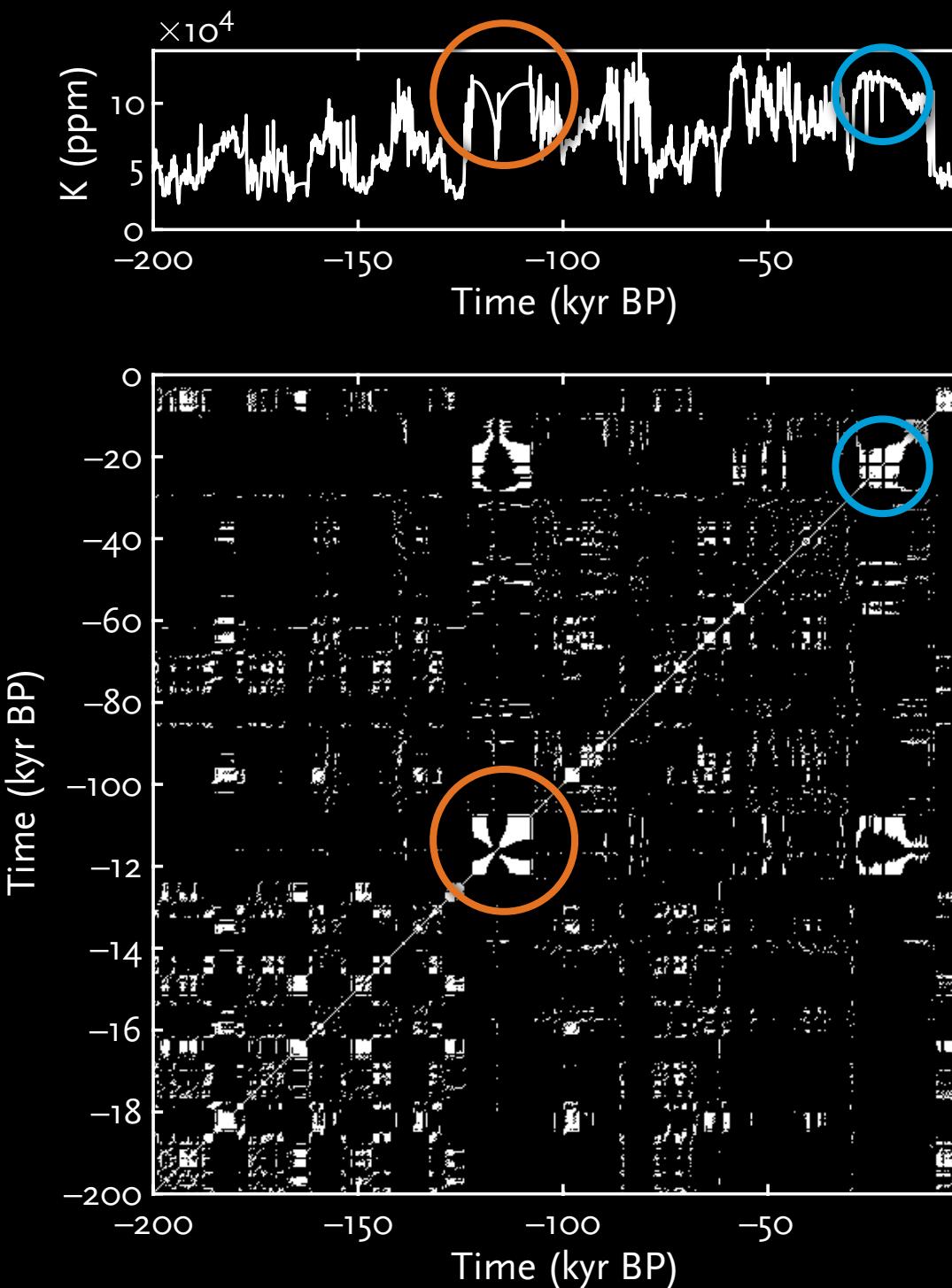
# VERTICAL LINES, BLOCKS



- System is trapped for some time, persistence
- Intermittency/ transition from one regime to another one
- Sojourn points

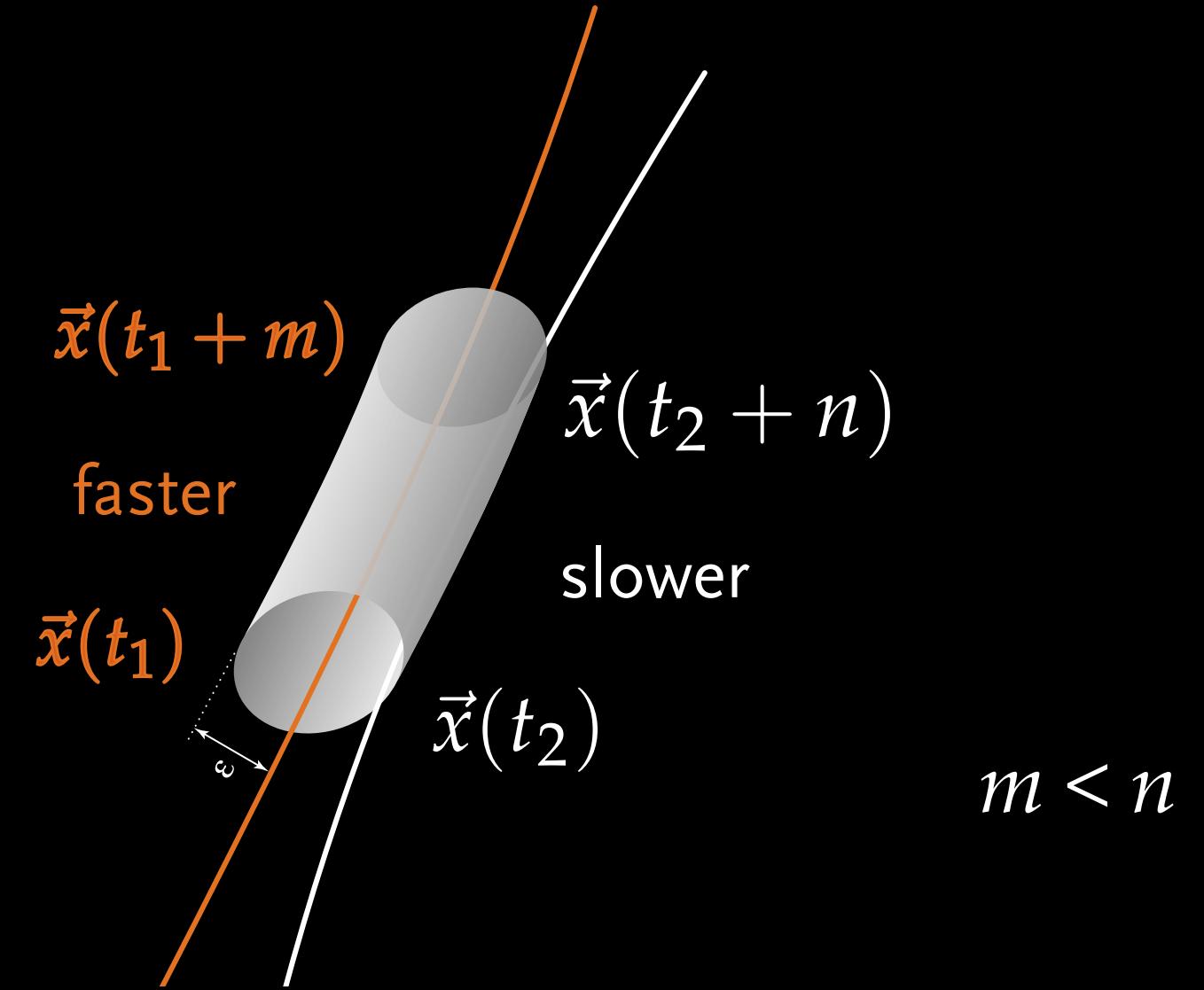
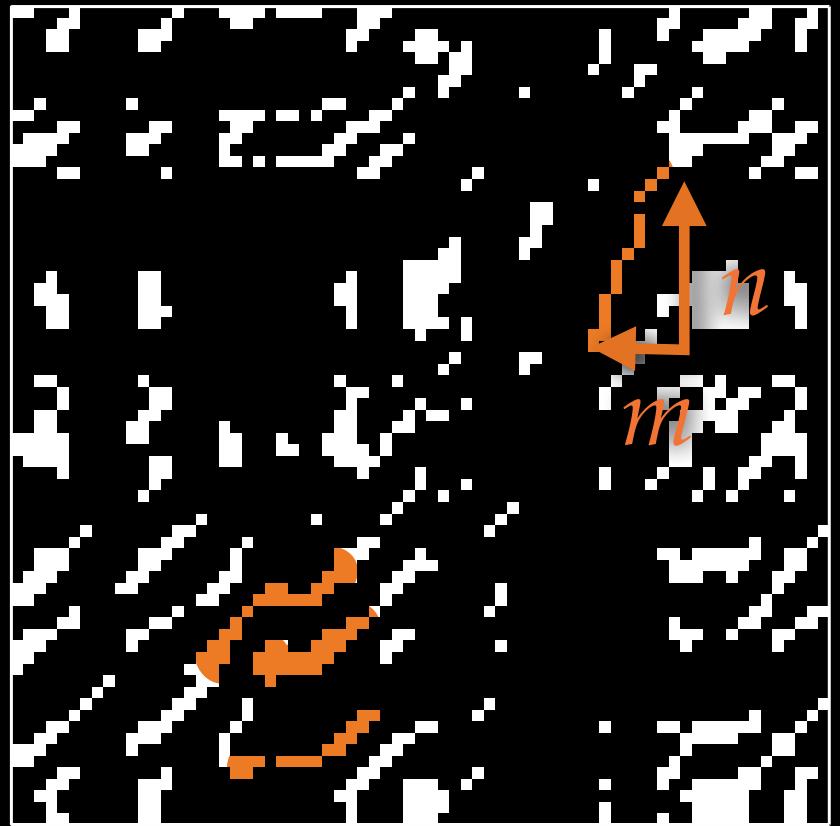


# VERTICAL LINES, BLOCKS



- Example: (lake sediment) proxy indicating wet climate in East Africa
- Persistence of states
- Check carefully when interpolation is used

# NON-DIAOGONAL AND BOWED LINES



- Recurrent epoch:

$$\vec{x}(t_1, t_1 + \Delta t_1, \dots, t_1 + m) \approx \vec{x}(t_2, t_2 + \Delta t_2, \dots, t_2 + n)$$

# NON-DIAOGONAL AND BOWED LINES

- Time transfer function:

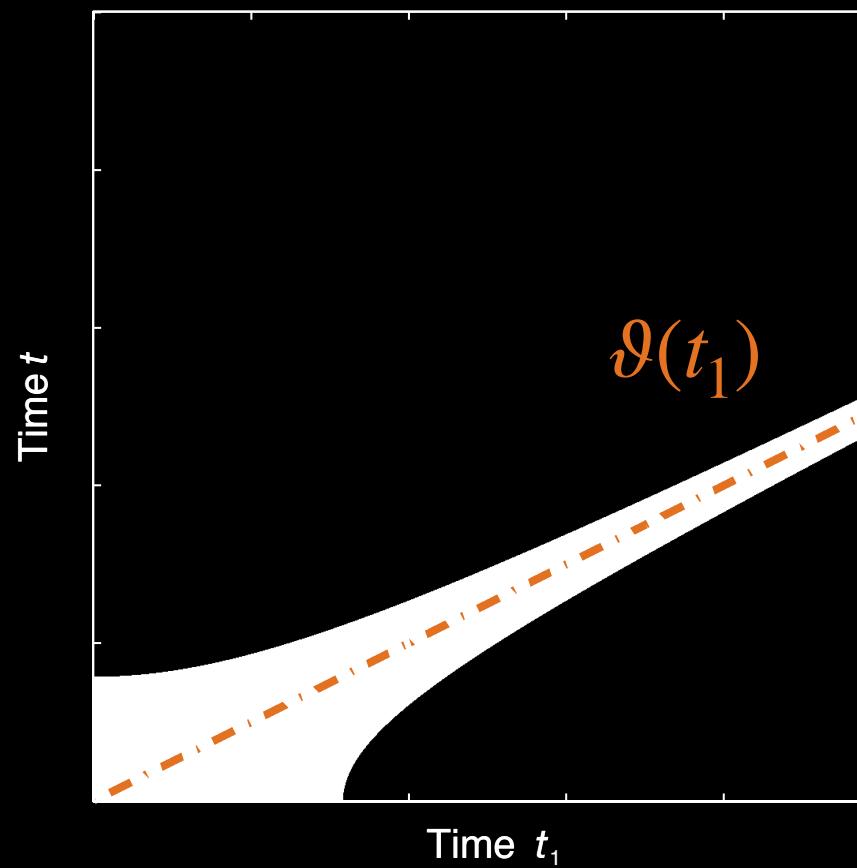
$$t_1 = T_2(t)$$

$$t = T_2^{-1}(t_1) = \vartheta(t_1)$$

- More general:

$$\vartheta(t_1) = T_2^{-1}(t_1)$$

$$\vartheta(t) = T_2^{-1} (T_1(t))$$



# NON-DIAOGONAL AND BOWED LINES

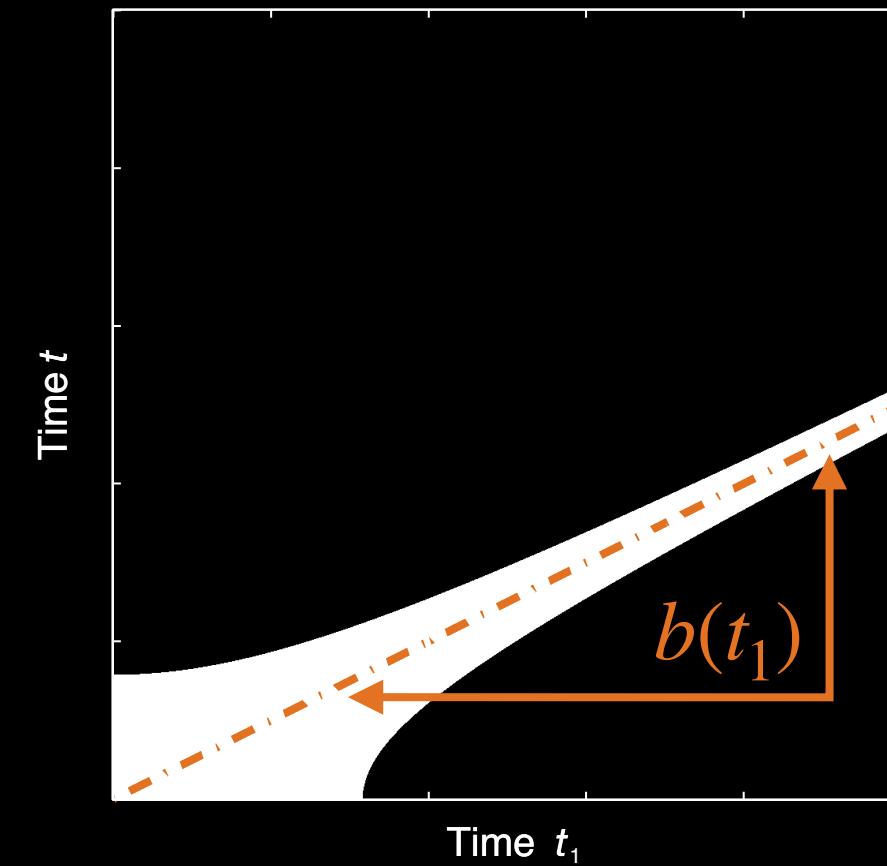
- Time transfer function:

$$t_1 = T_2(t)$$

$$t = T_2^{-1}(t_1) = \vartheta(t_1)$$

- Slope:

$$b(t_1) = \frac{d}{dt} T_2^{-1}(t_1)$$



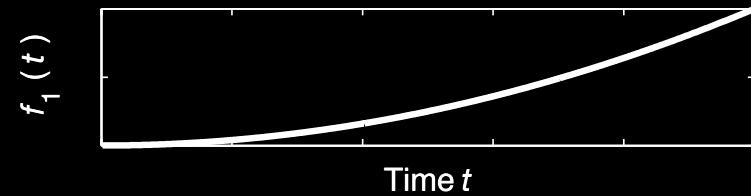
Slope depends only on local transformation of time scale!

i.e. change in velocity of trajectory, but not on the attractor itself

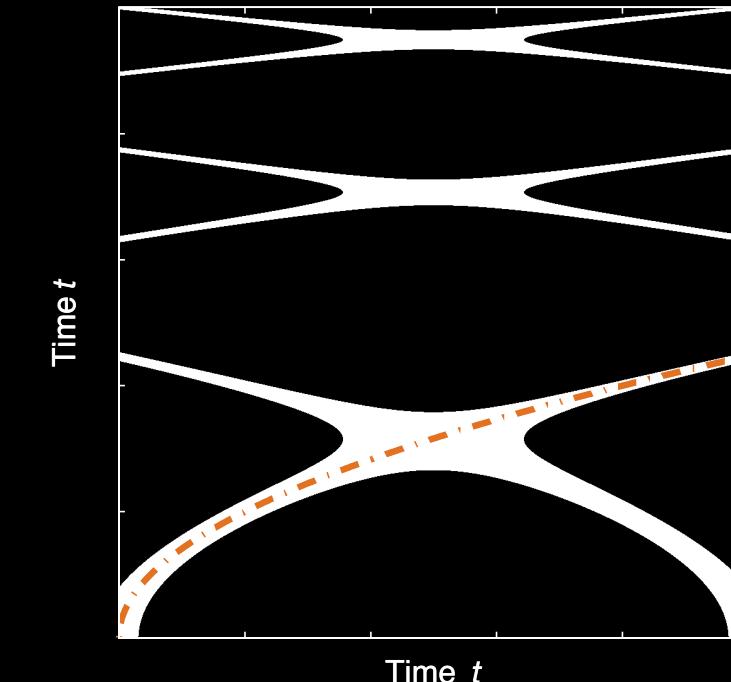
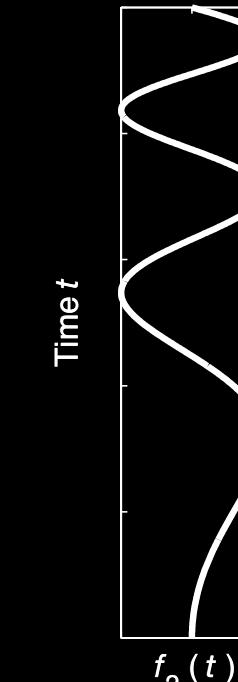
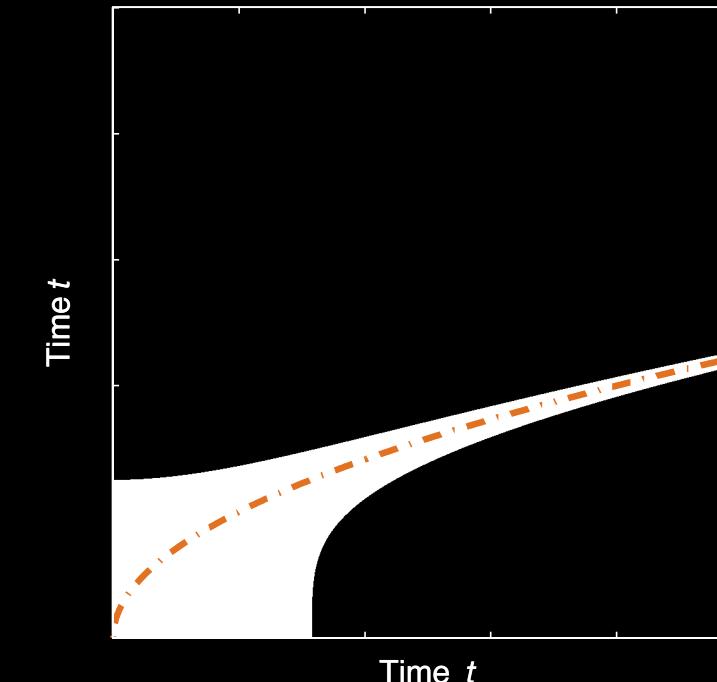
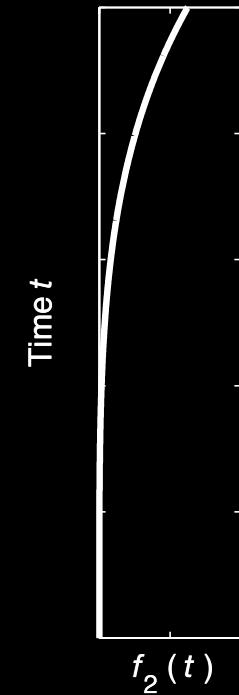
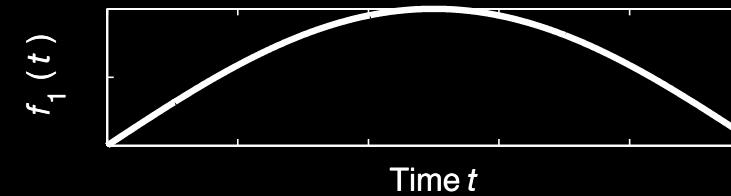
# NON-DIAOGONAL AND BOWED LINES

$$T_1 = t \quad T_2 = 5t^2$$

$$f(t) = t^2$$



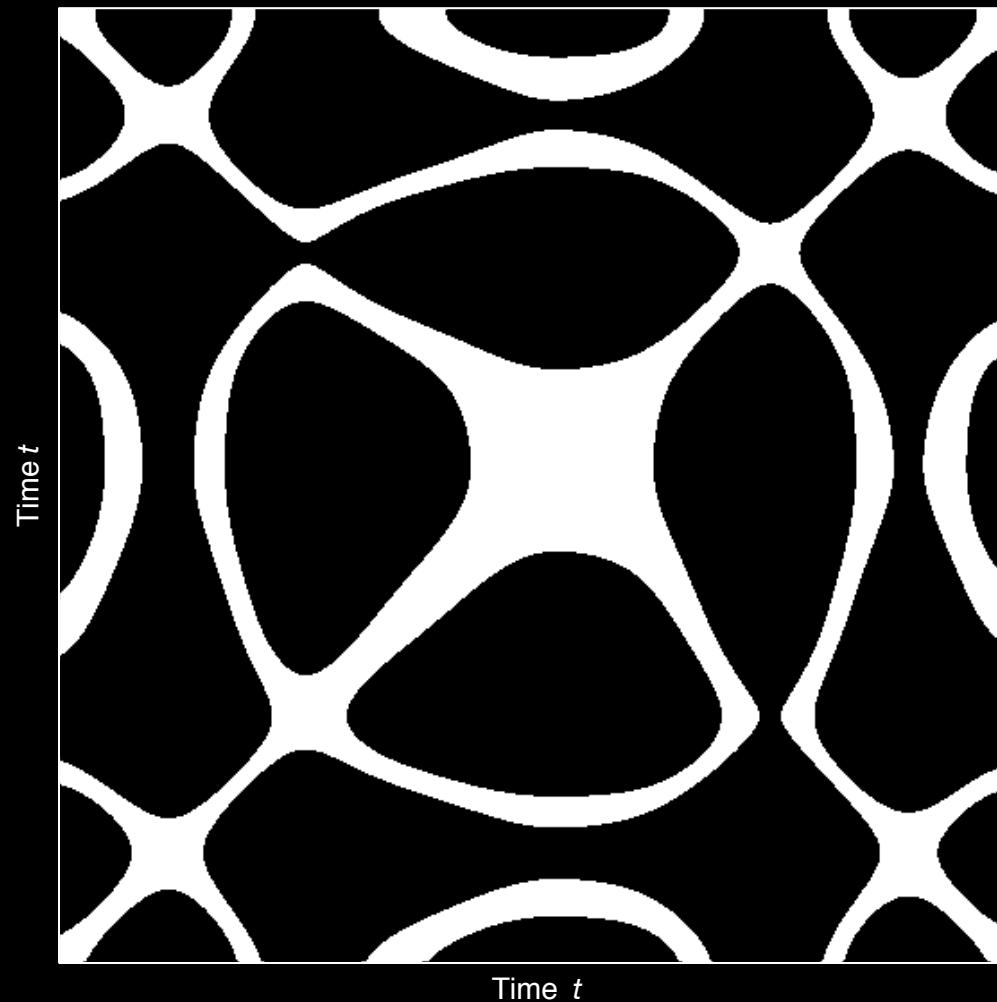
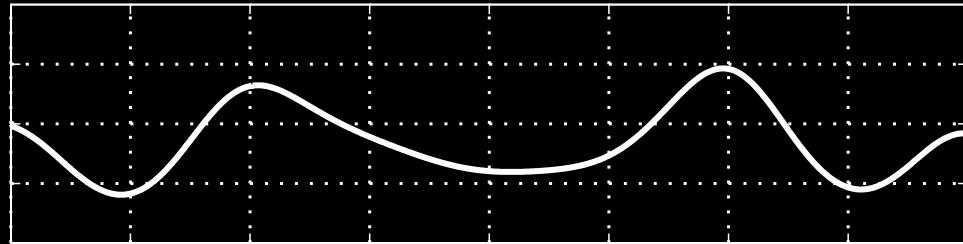
$$f(t) = \sin(\pi t)$$



$$\vartheta(t) = T_2^{-1} (T_1(t))$$

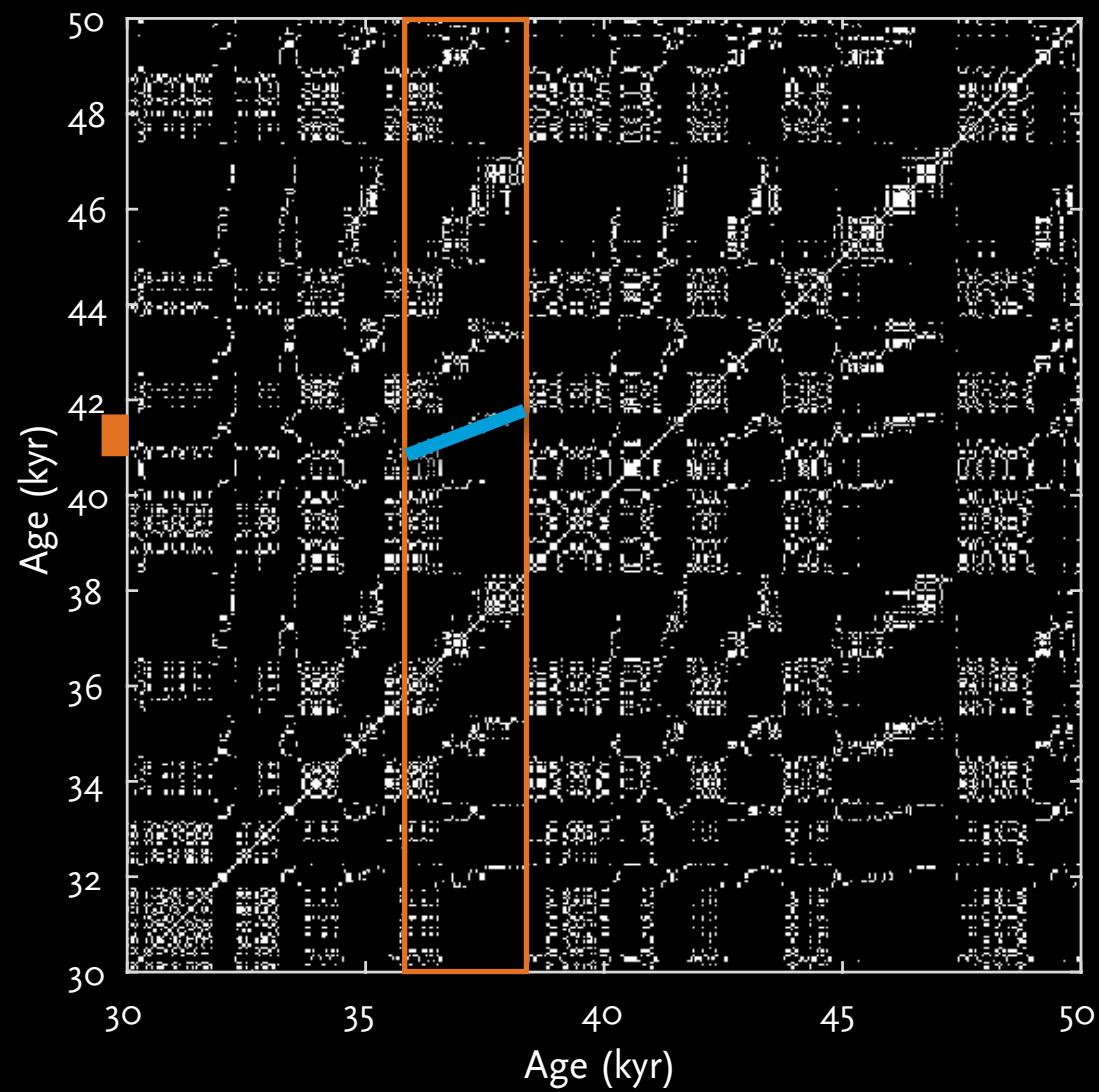
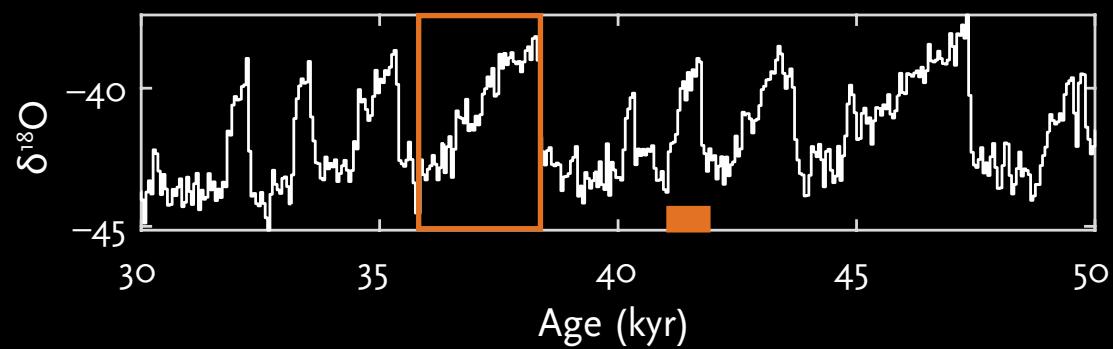
$$\vartheta(t) = \sqrt{0.2t}$$

# NON-DIAOGONAL AND BOWED LINES

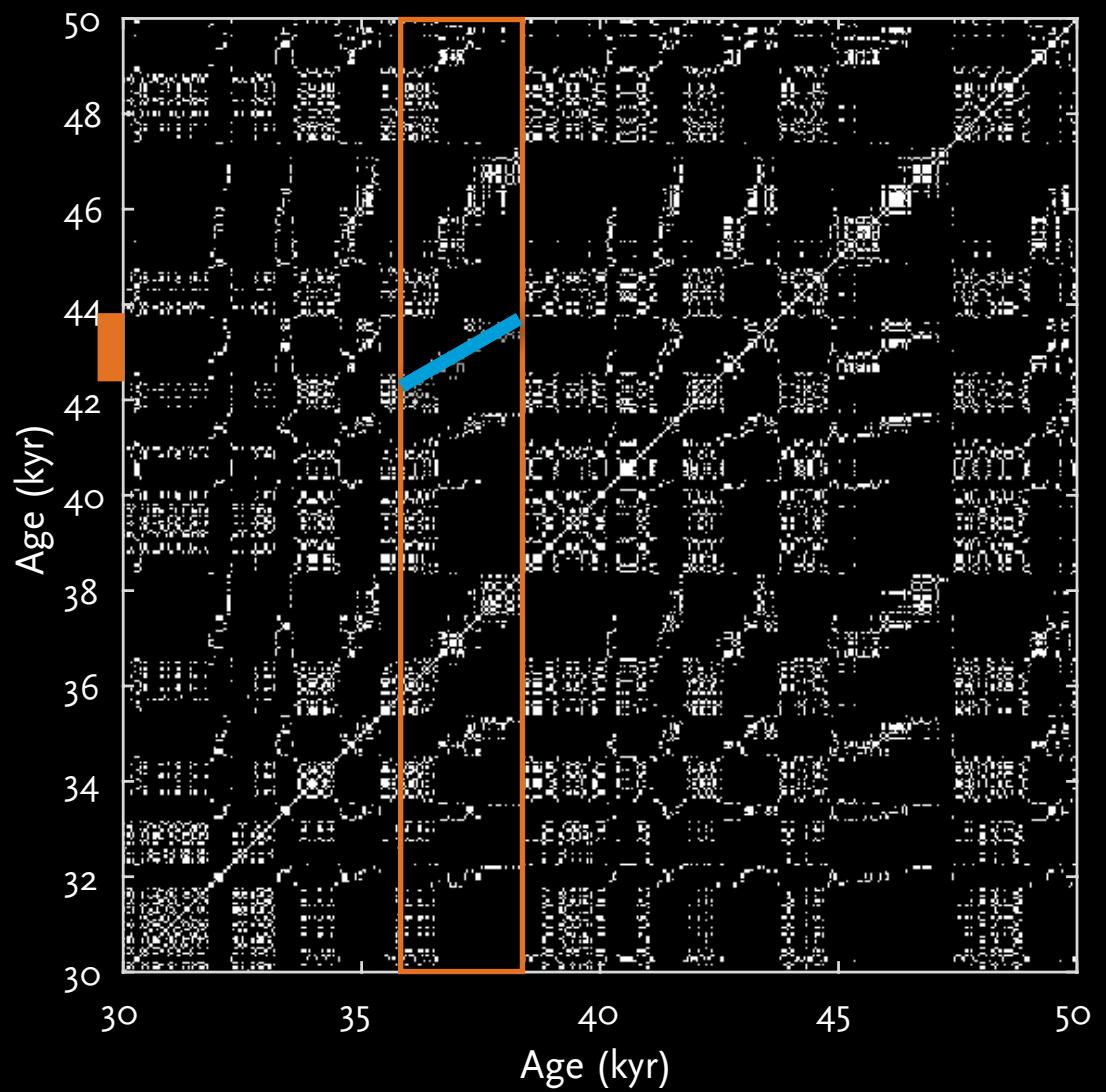
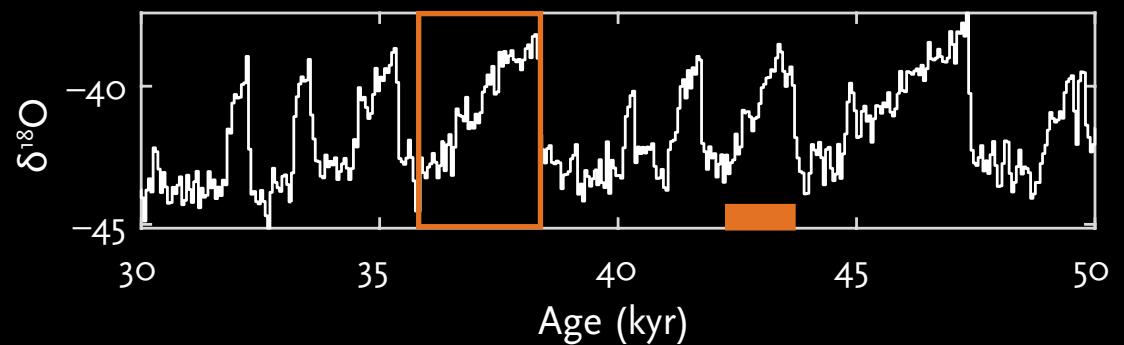


- Evolution of states is similar at different epochs but with different velocity
- Dynamics of the system could be changing

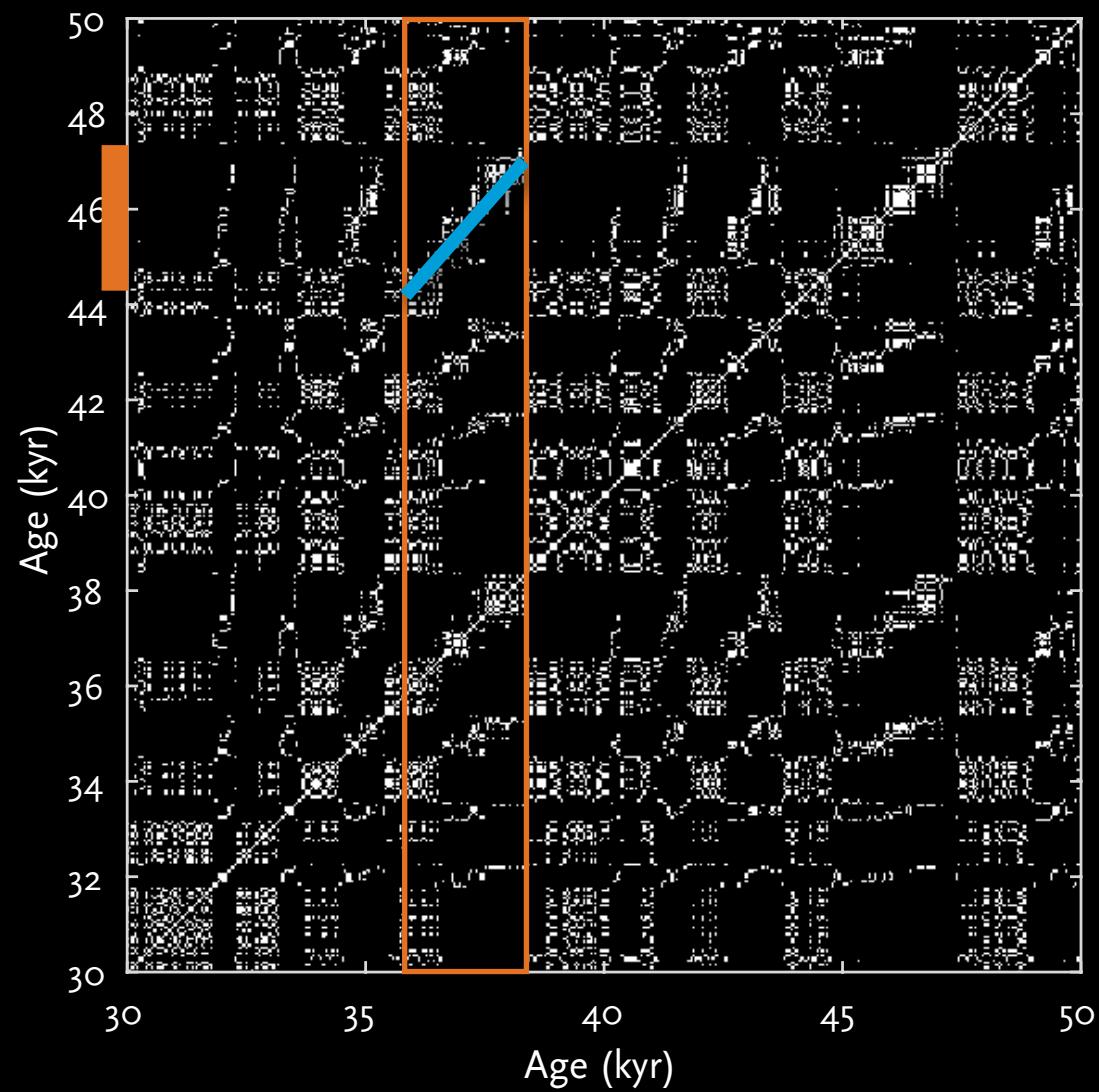
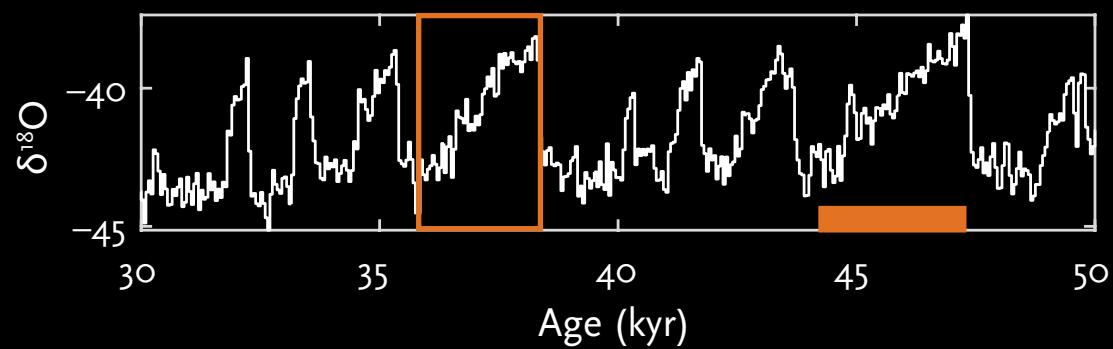
# NGRIP ICE CORE RECORD



# NGRIP ICE CORE RECORD



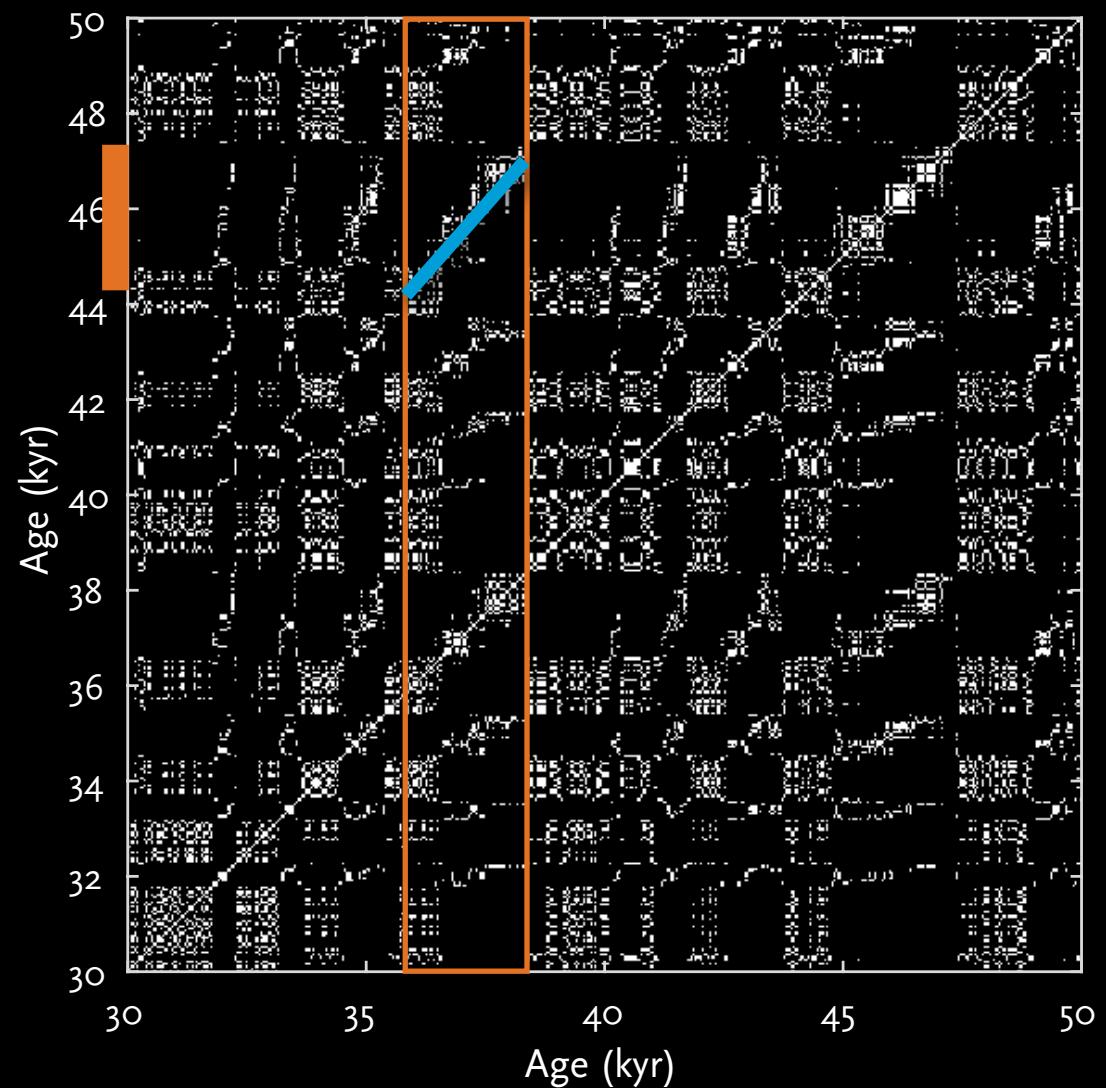
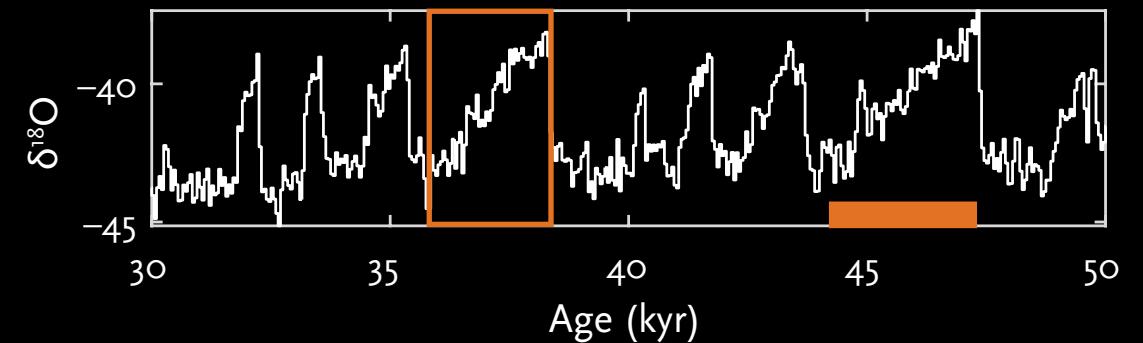
# NGRIP ICE CORE RECORD



# NON-DIAOGONAL AND BOWED LINES

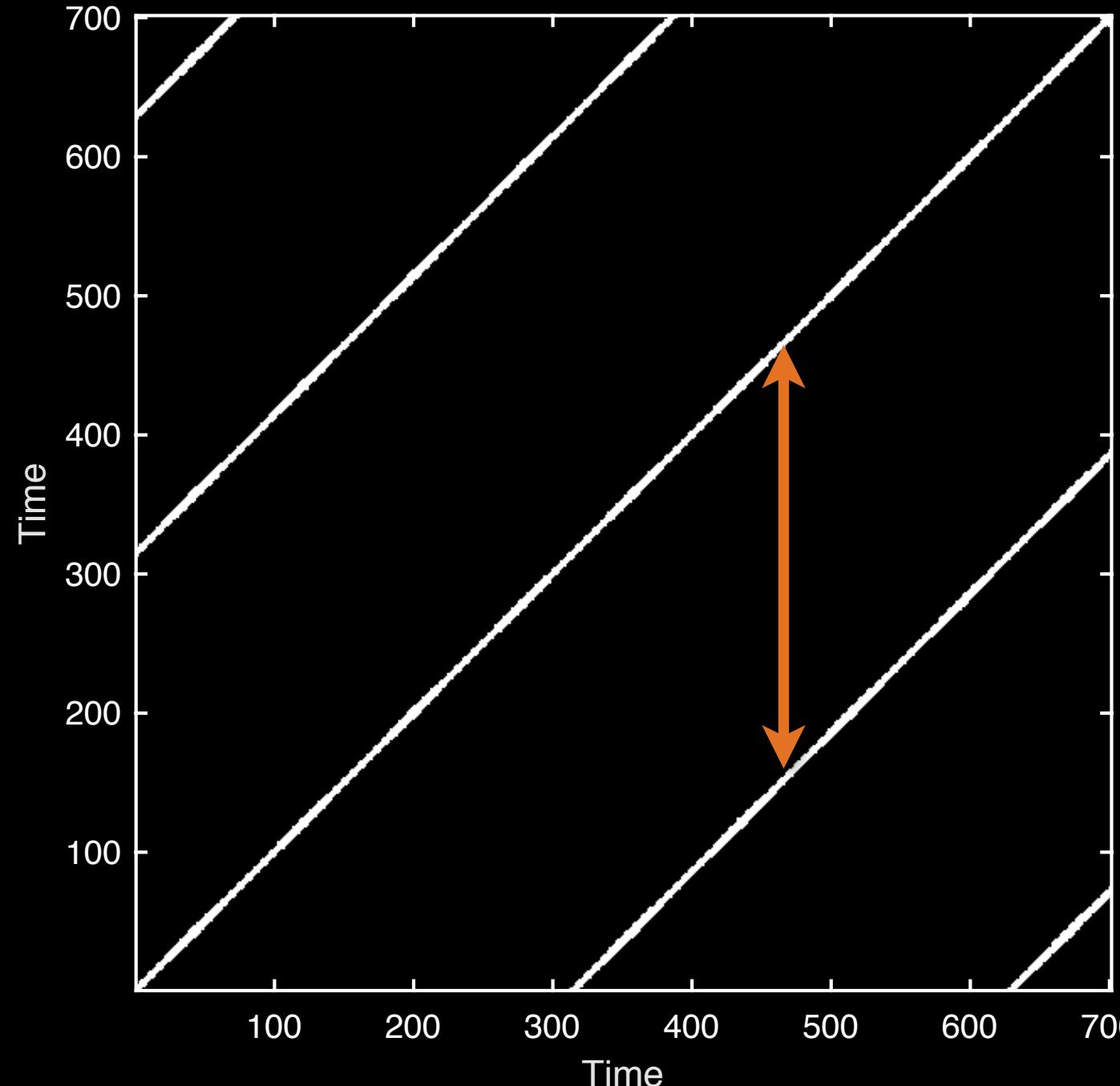
NGRIP

- Lines with slopes  $\approx 45^\circ$  are interesting!
- Change of temporal resolution  
(sampling issues!)
- Change of dynamics
- System with different time scales  
(e.g., high and low frequencies)
- Dynamics with self-similarity(?)



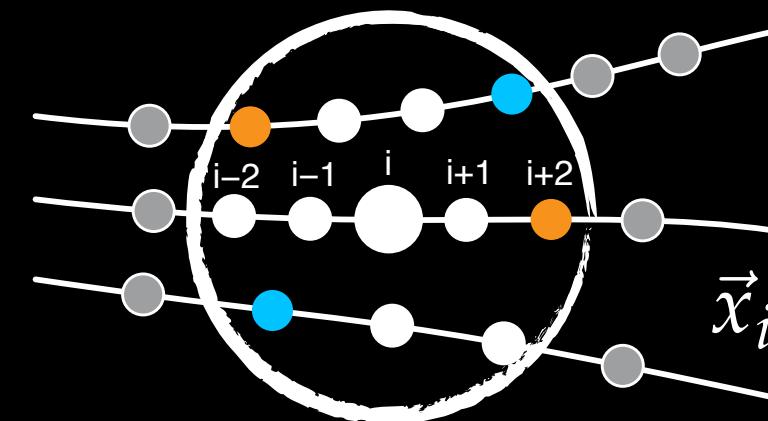
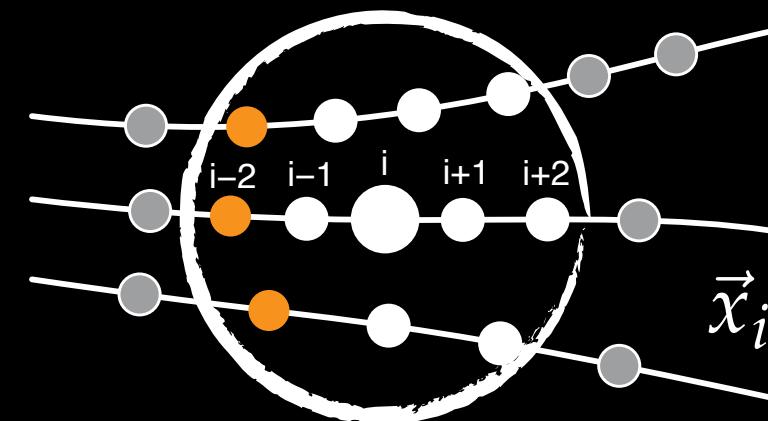
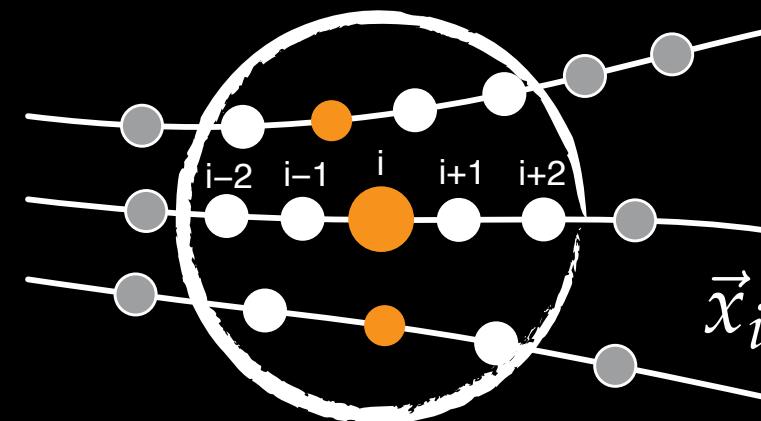
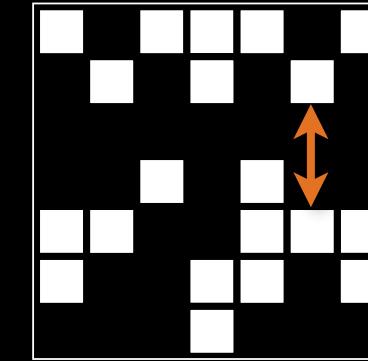
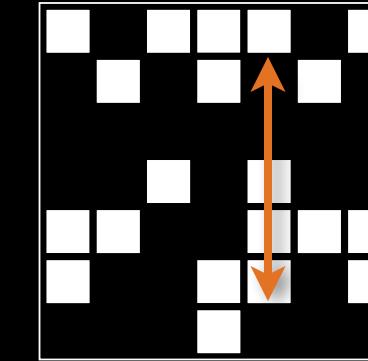
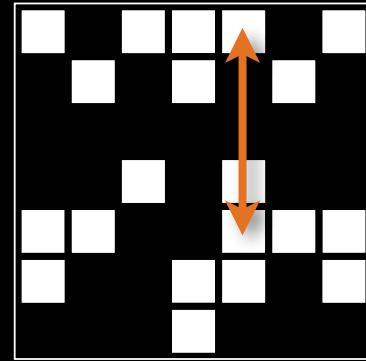
# RECURRENCE TIMES

Periodic dynamics



- Periodic length:  
vertical distance between  
recurrence points  
(vertical “empty” lines)

# RECURRENCE TIMES

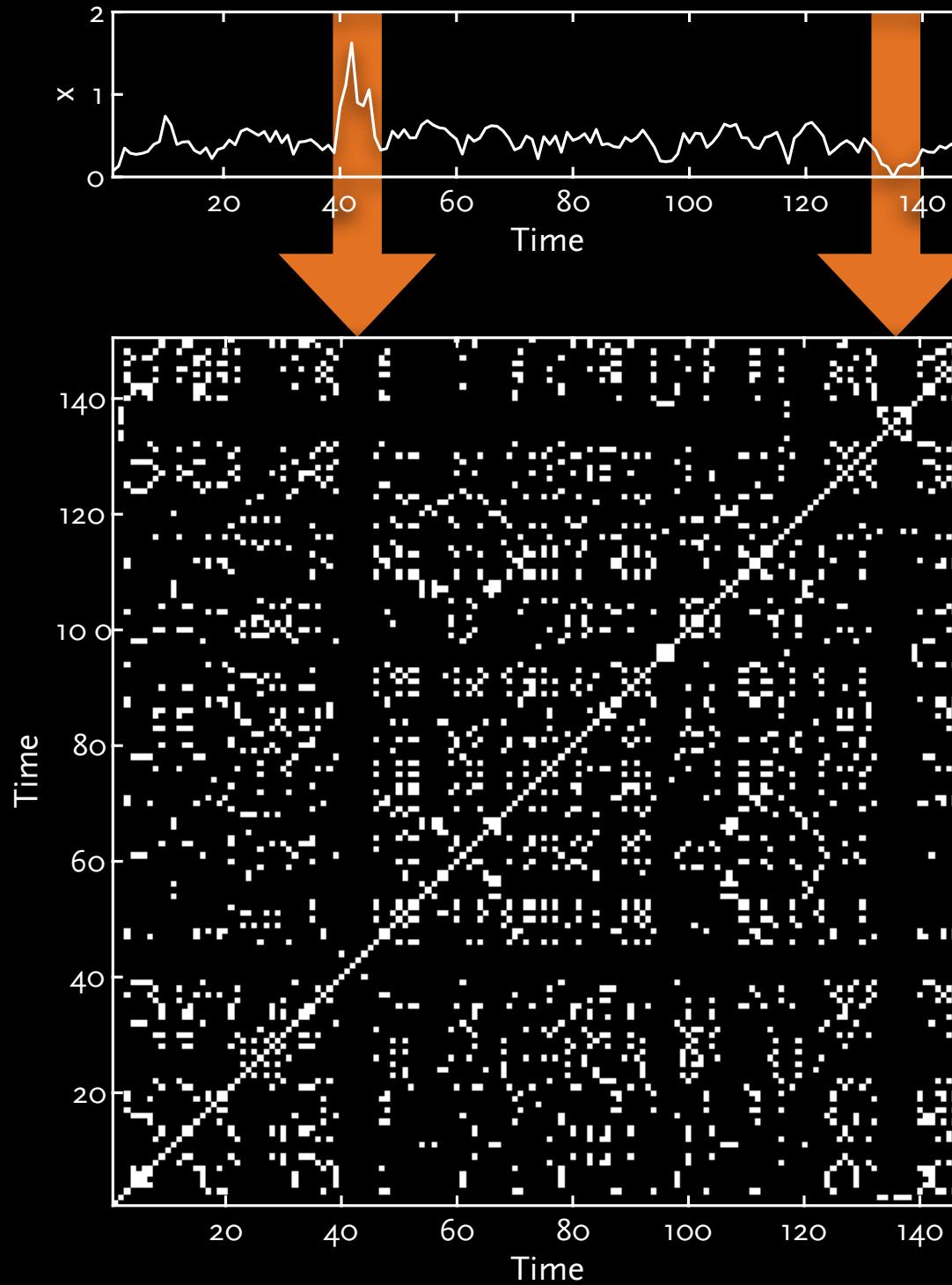


- Recurrence time distribution  $P(T)$
- Complexity of the time scale of recurrence (recurrence time entropy)

# VISUAL INTERPRETATION

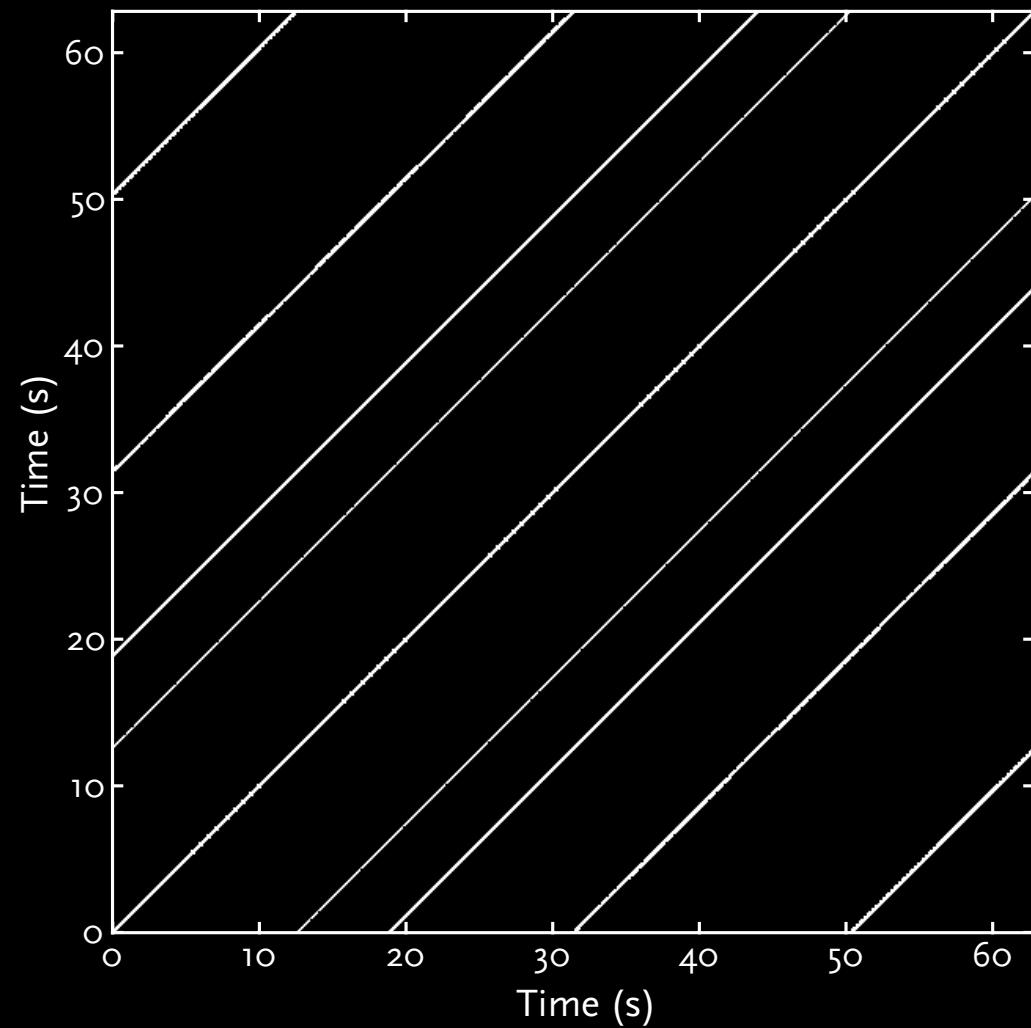
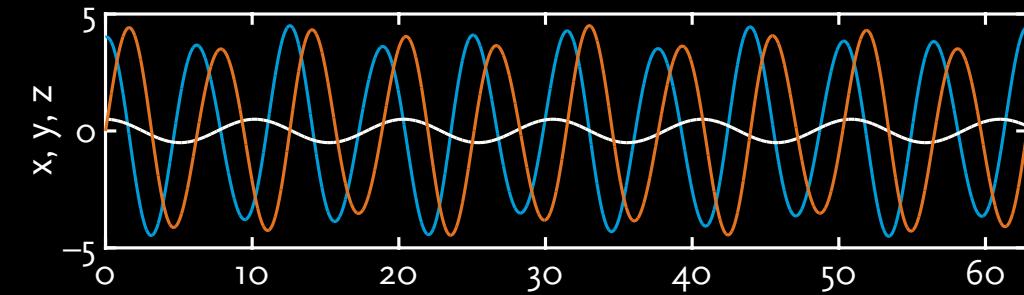
Typology (large scale appearance)	homogeneous	periodic	trend	disrupted, separated into blocks
Texture (small scale structures)	only single points	mainly diagonal lines	vertical lines, black blocks	curved pattern
Characteristic pattern	noninterrupted diagonal lines, periodic separation of diagonal lines	sparingly occupied columns/ bars	missing recurrences	local changes in recurrence point density

# DISRUPTIVE STRUCTURES IN RECURRENCE PLOTS



- Empty/ sparsely occupied columns
- Rare or extreme events
- Divide RP into block-like sub-regions

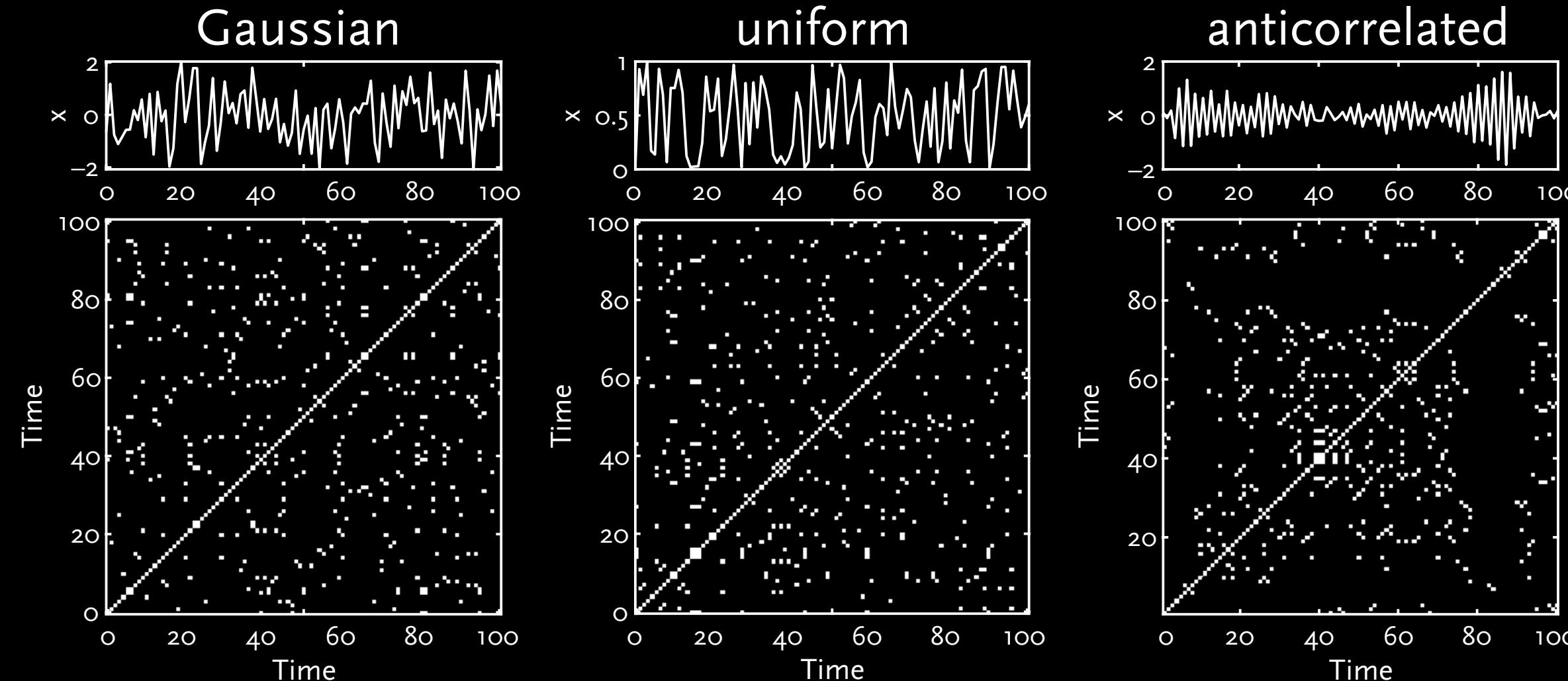
# PERIODIC AND QUASI-PERIODIC PATTERN



- Cyclicities in the process
- Time distance between periodic patterns (e.g. lines) corresponds to the periods
- Different distances between long diagonal lines reveal quasi-periodic processes

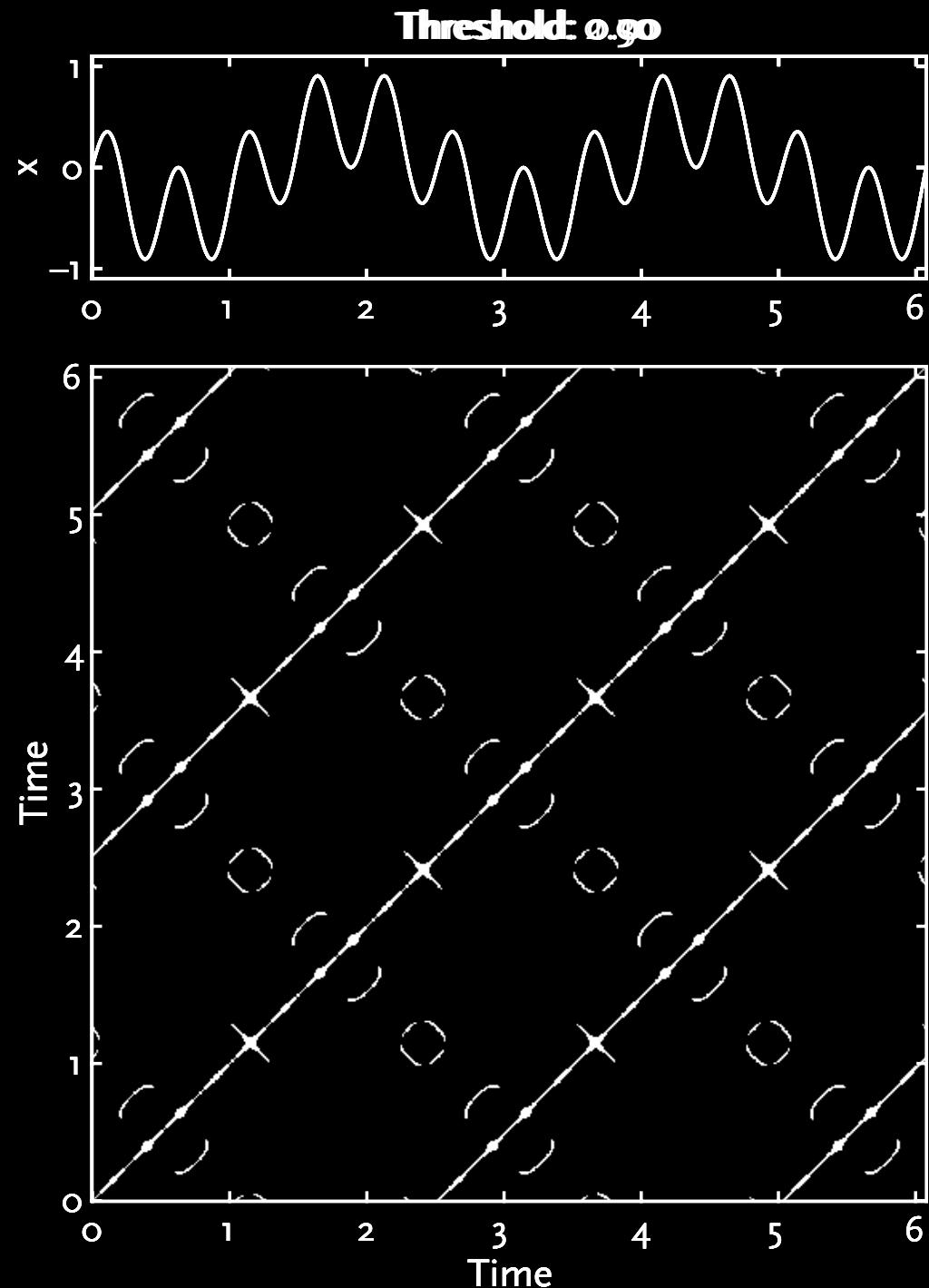
# SINGLE ISOLATED POINTS

- Strong fluctuation in the process
- If only single isolated points occur, the process may be an uncorrelated random or even anti-correlated process



# SENSITIVITY ON PARAMETERS

# RECURRENCE THRESHOLD



- Too small:
  - no points, no essential structures
- Too large:
  - interesting structures are merged, thick lines, completely full
- Optimal selection:
  - large variety of structures, thin lines, different time scales visible

# RECURRENCE THRESHOLD: QUANTILE APPROACH

- Preselect  $RR$  (e.g., to have  $RR = 10\%$ )  
 $RR(\varepsilon) \rightarrow \varepsilon$

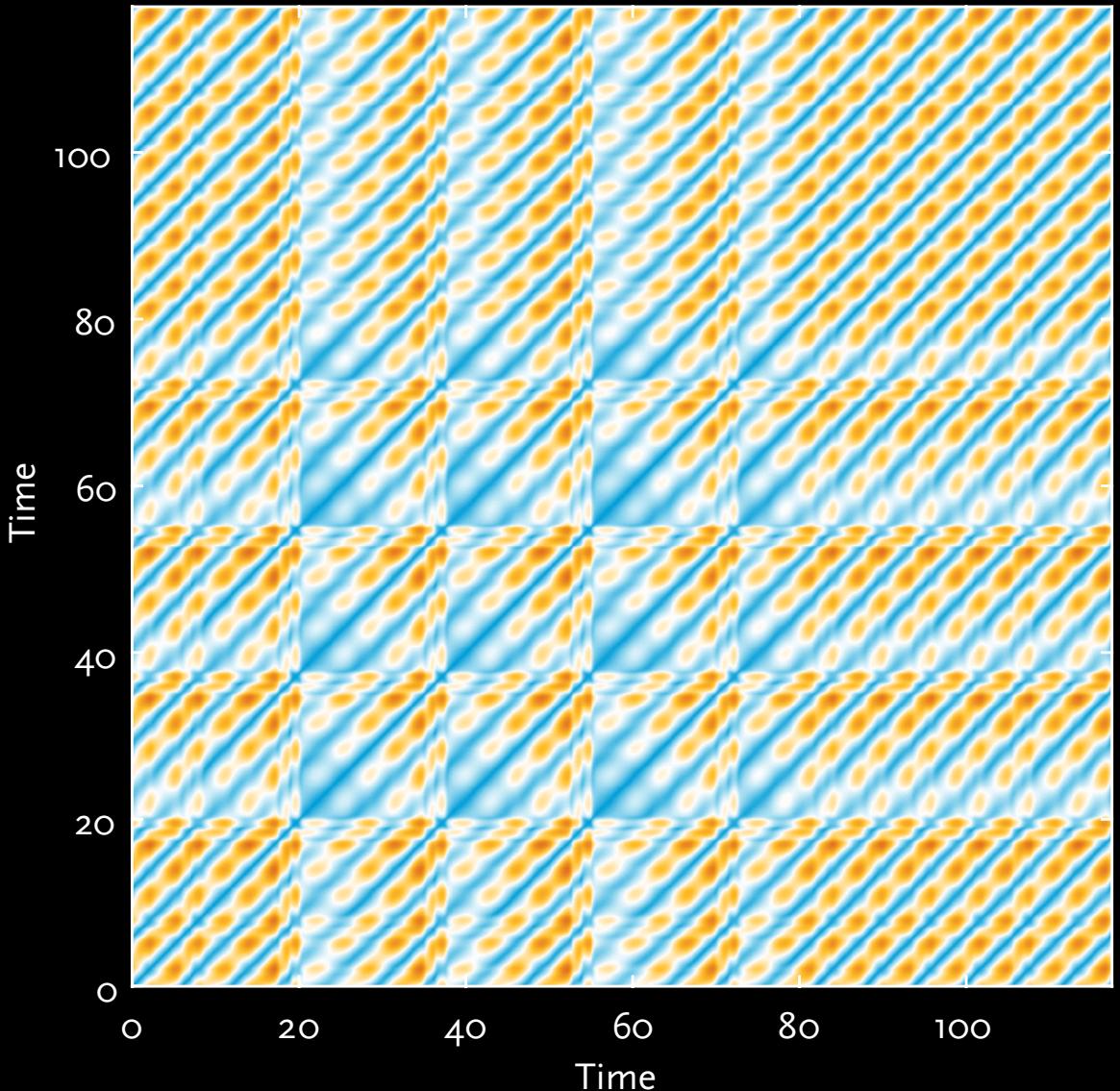
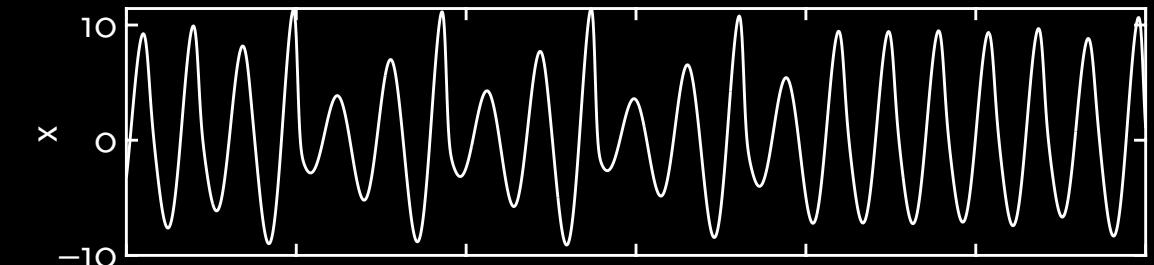
- Recurrence rate:

$$RR(\varepsilon) = \frac{1}{N(N-1)} \sum_{i,j,i \neq j}^N R_{i,j}(\varepsilon)$$

- Distance matrix:

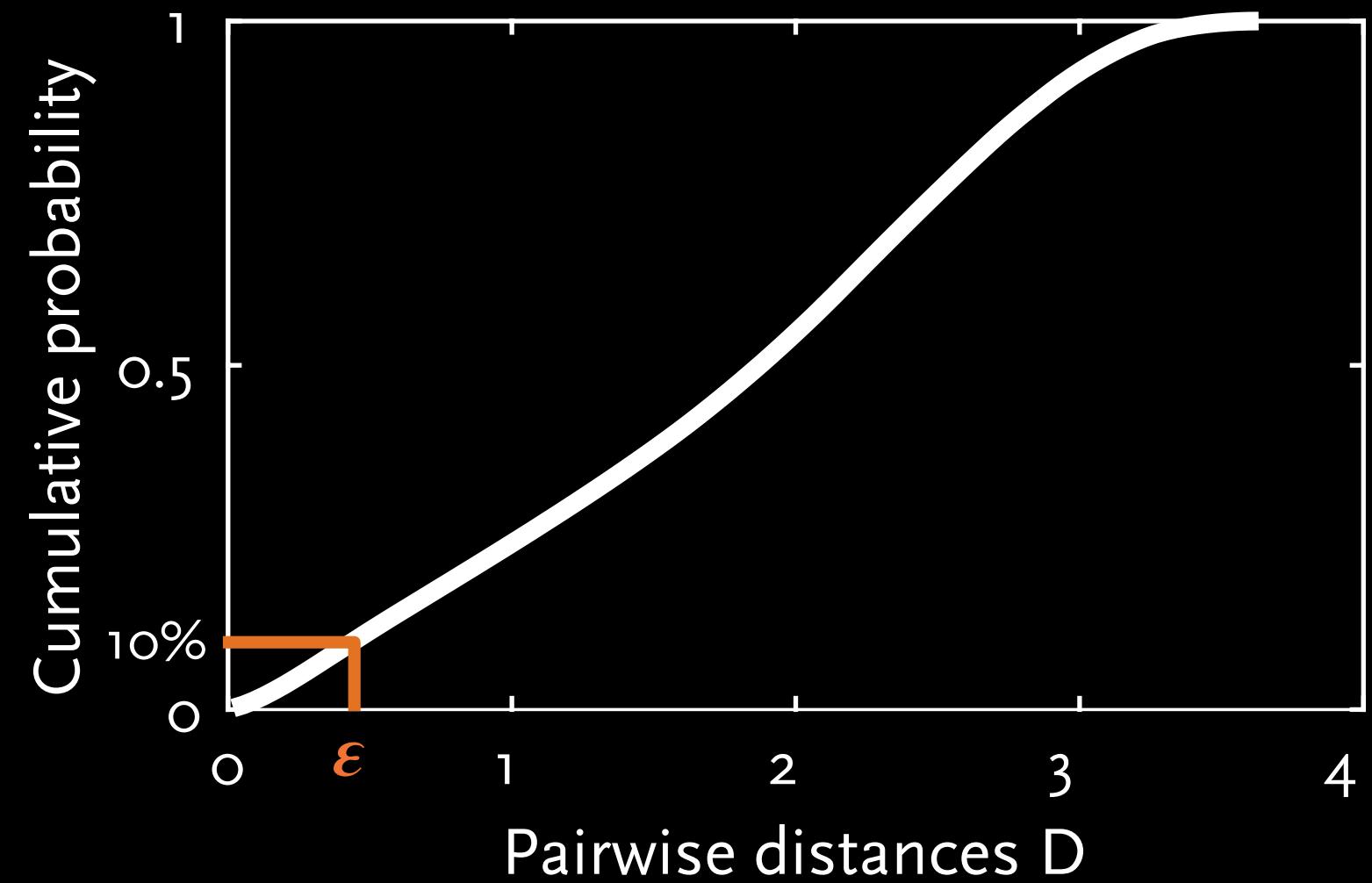
$$R_{i,j}(\varepsilon) = \Theta(\varepsilon - D_{i,j})$$

$$D_{i,j} = \|\vec{x}_i - \vec{x}_j\|$$



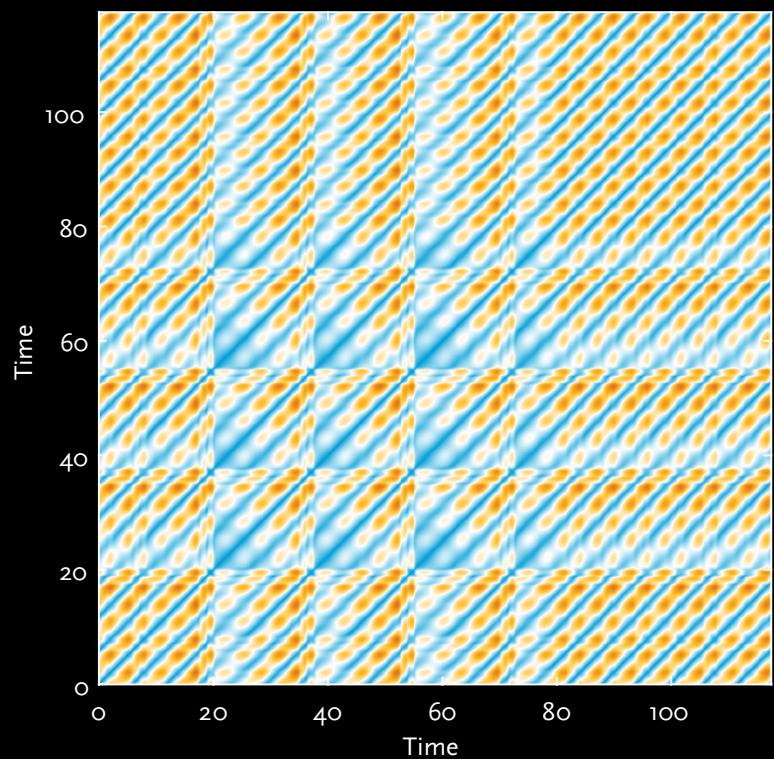
# RECURRENCE THRESHOLD: QUANTILE APPROACH

- Preselect  $RR$   
 $RR(\varepsilon) \rightarrow \varepsilon$
- Quantile of distance distribution  
 $p(x < D_{i,j})$
- e.g.,  $\varepsilon = D_{0.1}$   
i.e.,  $0.1 = p(\varepsilon < D_{i,j})$
- Quantile = recurrence rate

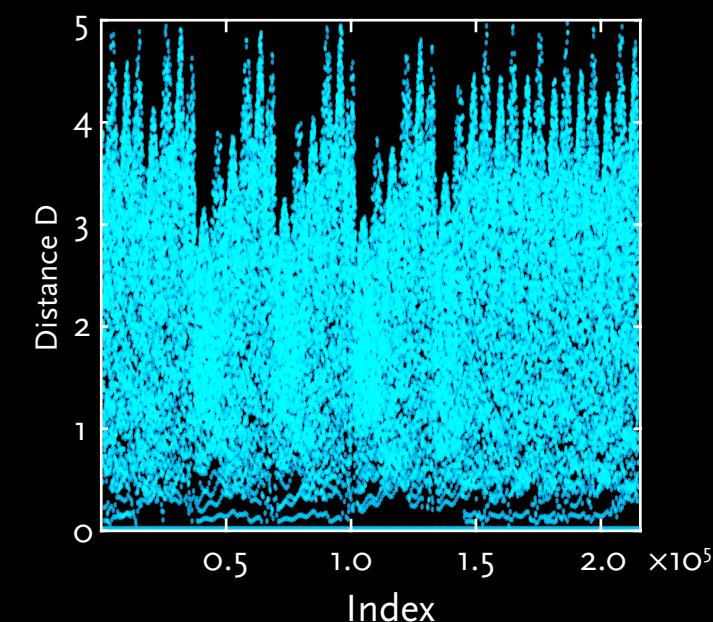


# RECURRENCE THRESHOLD: QUANTILE APPROACH

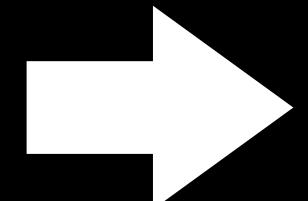
Distance matrix  $D_{i,j}$



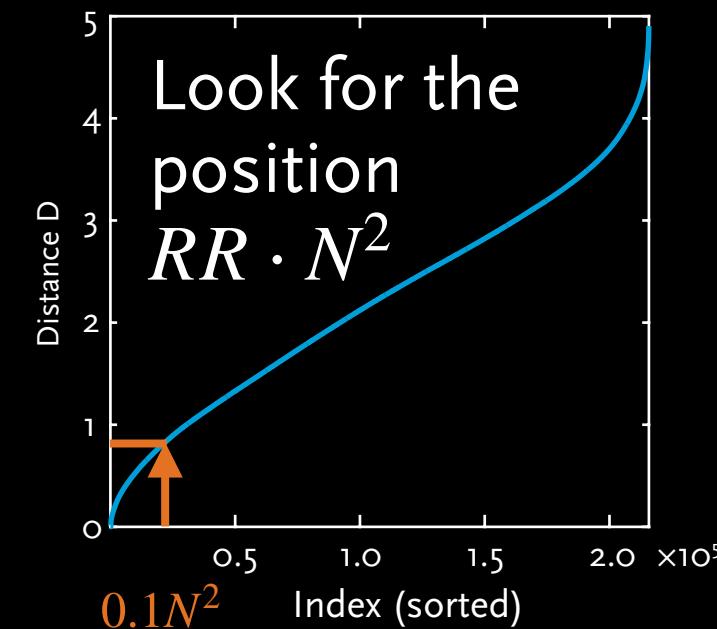
Distances  $D_{i,j}$



Sort  
distance  
values

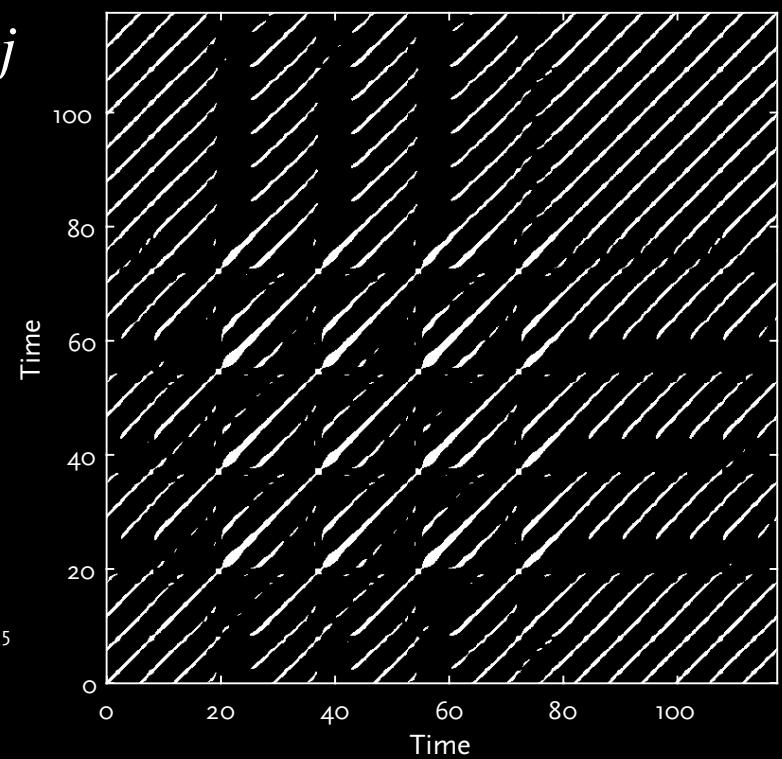


Sorted distances  $D_{i,j}$



e.g.,  $\varepsilon = D_{0.1}$ , i.e.,  $0.1N^2 = p(\varepsilon < D_{i,j})$

Recurrence  $D_{i,j} < \varepsilon$



- Robust recurrence characteristics in different embedding dimensions

# RECURRENCE THRESHOLD: CONNECTEDNESS

- Connectivity of recurrence matrix

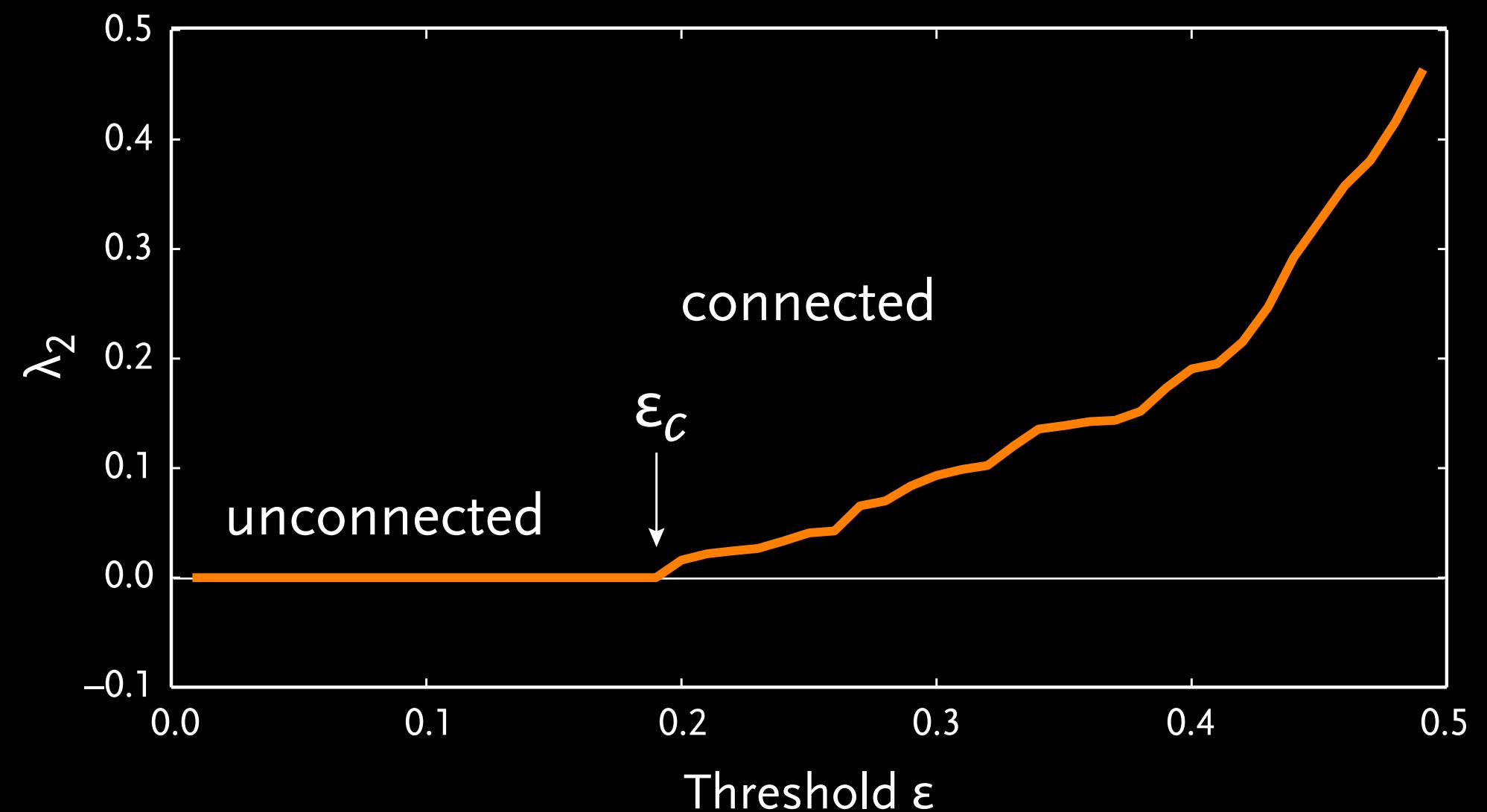
$$A_{i,j} = R_{i,j} - \delta_{i,j}$$

- Laplace matrix

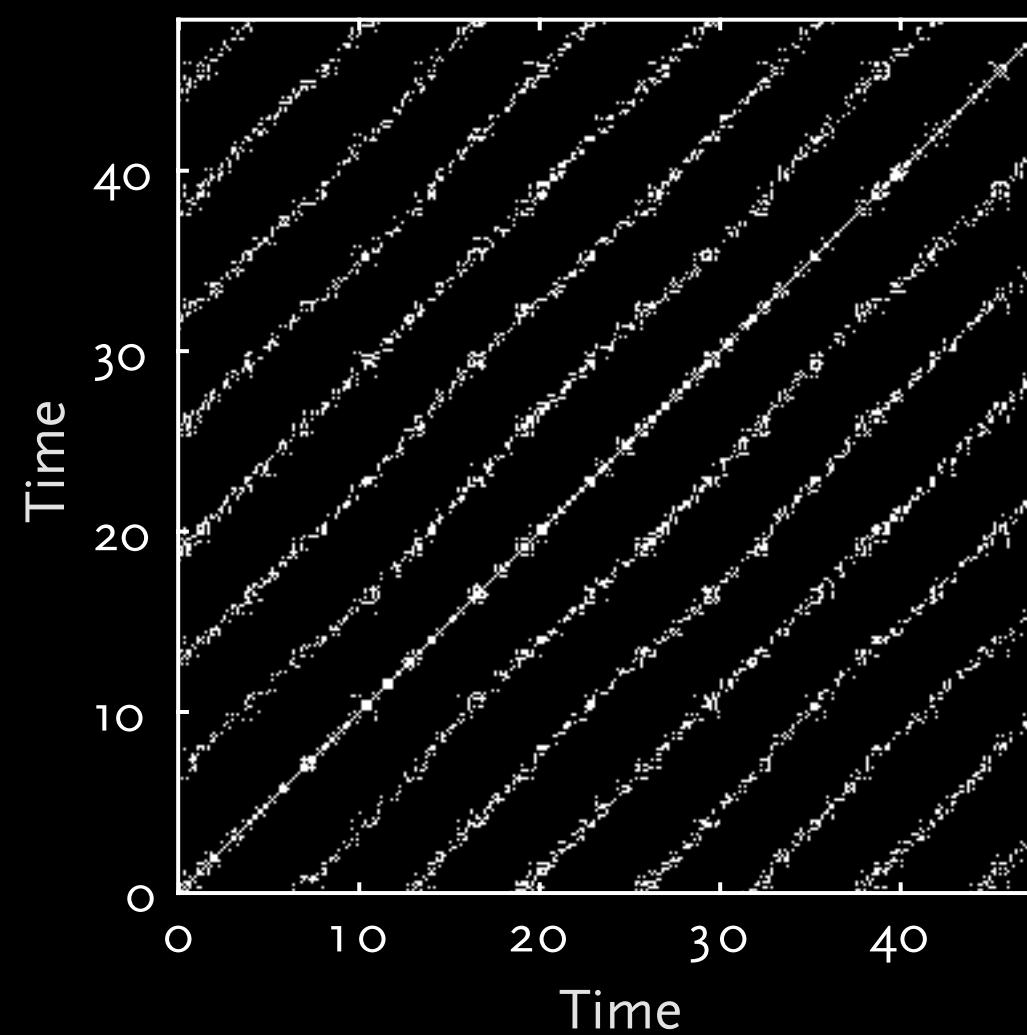
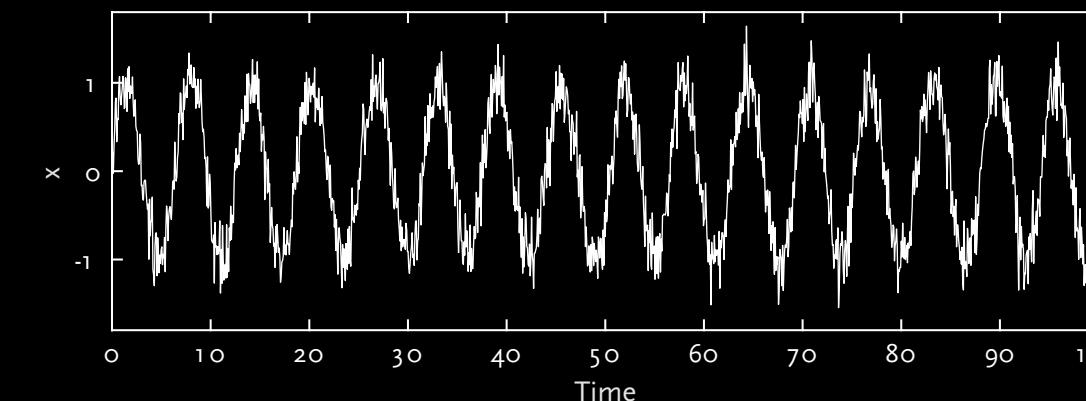
$$L_{i,j} = \delta_{i,j} \sum_j R_{i,j} - R_{i,j}$$

- $\epsilon \rightarrow 2^{\text{nd}}$  smallest

Eigenvalue  $\lambda_2 > 0$

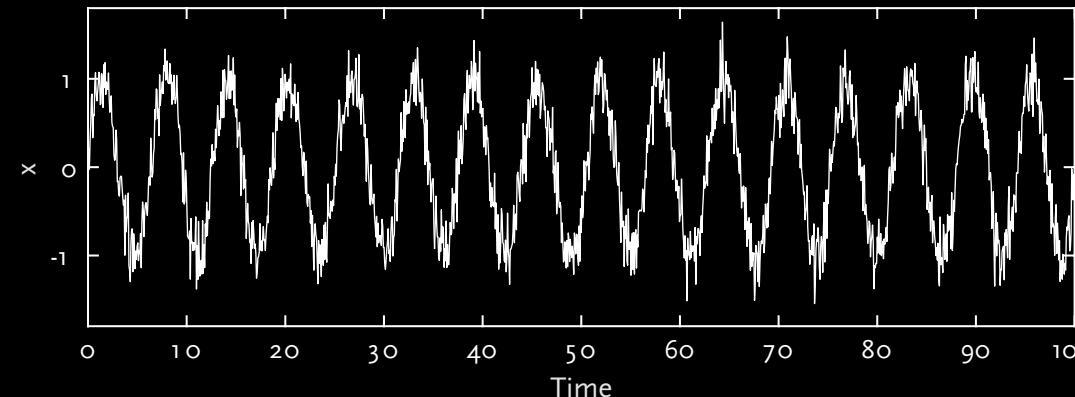


# RECURRENCE THRESHOLD: OPTIMIZING DIAGONALS



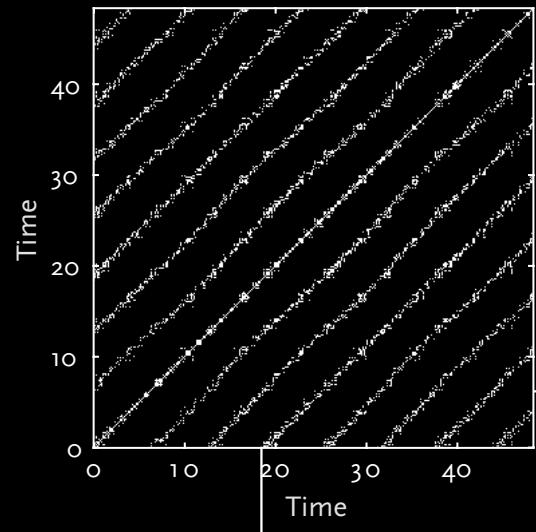
- Minimizing the fragmentation and thickness of the diagonal lines (for cyclical signals)

# RECURRENCE THRESHOLD: OPTIMIZING DIAGONALS



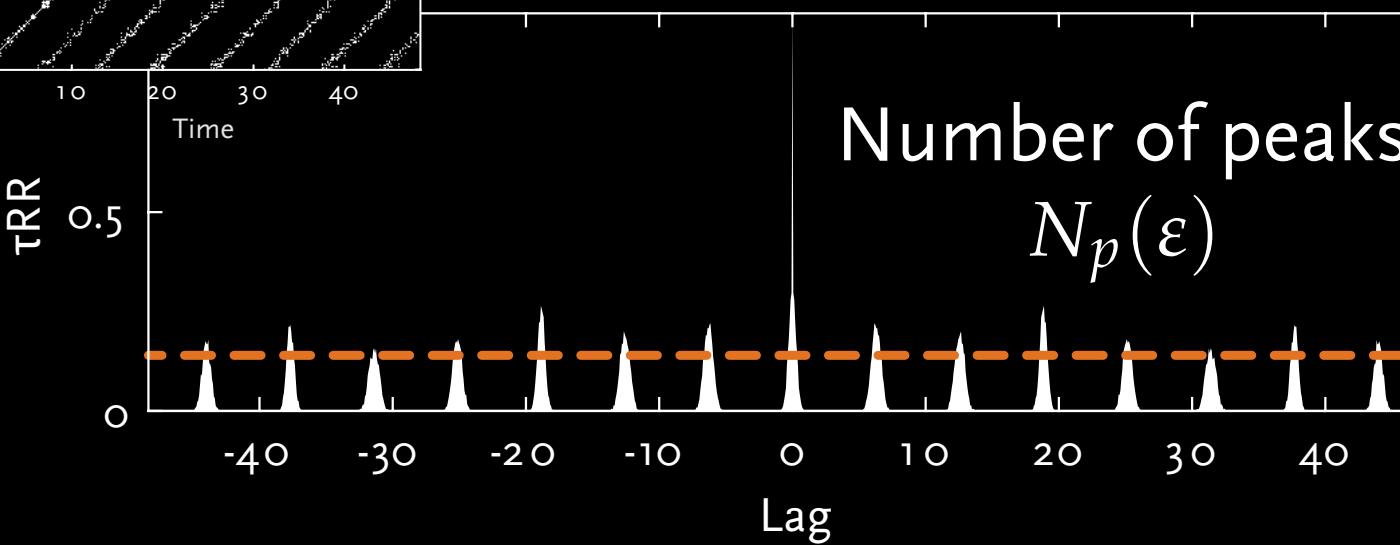
Number of neighbours

$$N_n(\varepsilon) = \frac{1}{N} \sum_{i,j} R_{i,j}(\varepsilon)$$



Number of peaks

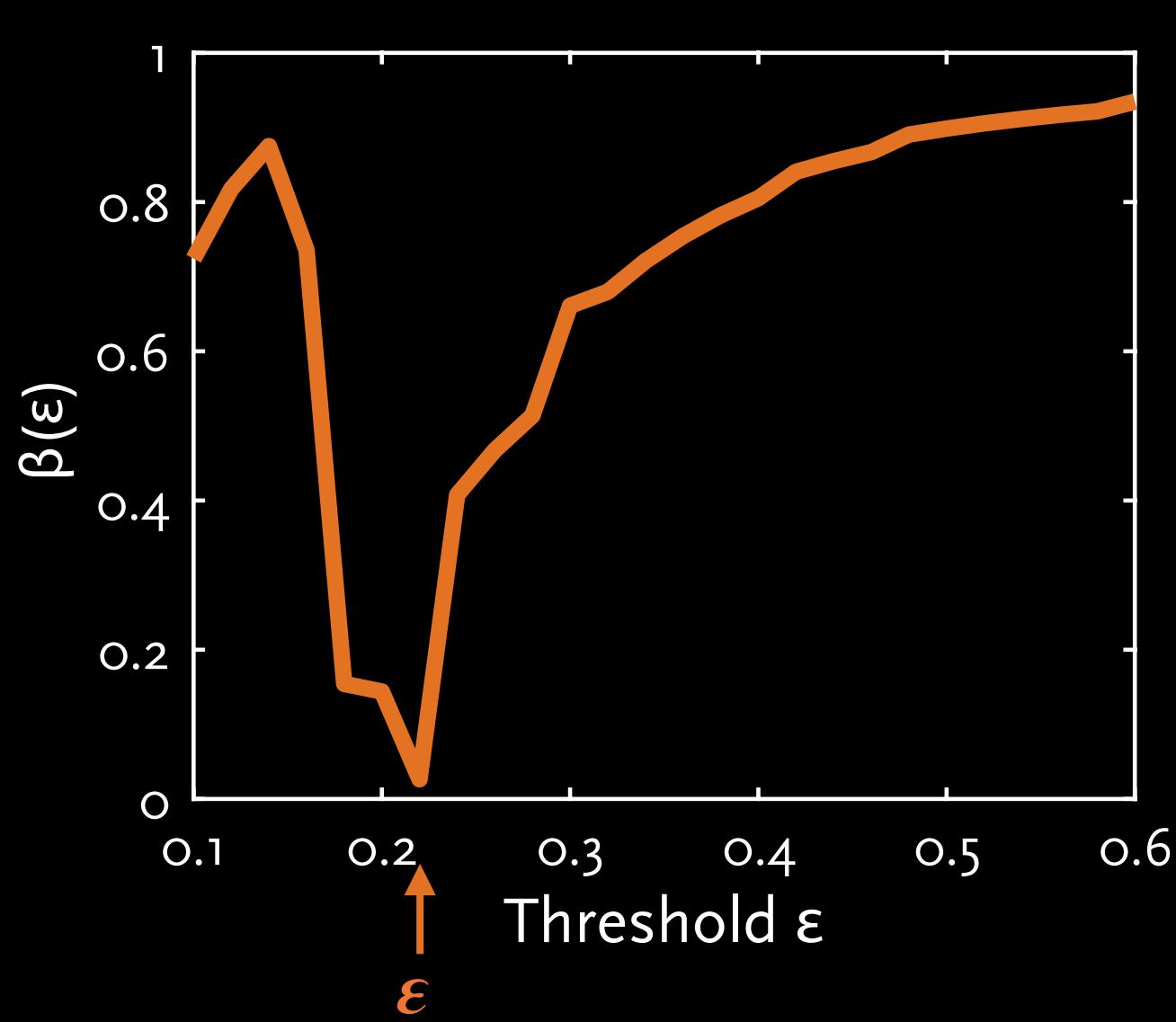
$$N_p(\varepsilon)$$



- Minimizing the fragmentation and thickness of the diagonal lines (for cyclical signals)
  - $N_p$  maximal
  - $N_n$  close to  $N_p$

$$\beta(\varepsilon) = \frac{|N_n(\varepsilon) - N_p(\varepsilon)|}{N_n(\varepsilon)}$$

# RECURRENCE THRESHOLD: OPTIMIZING DIAGONALS



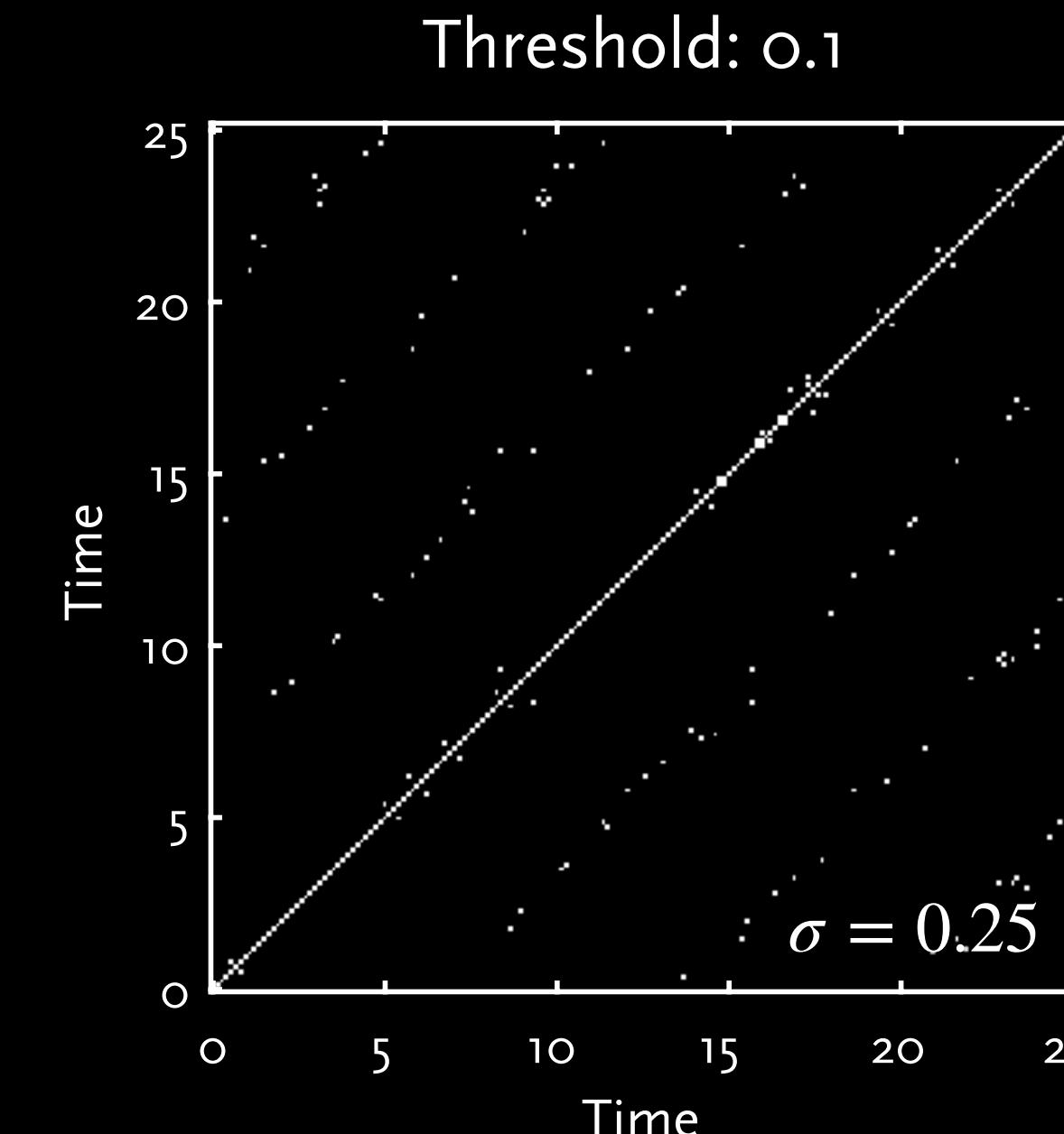
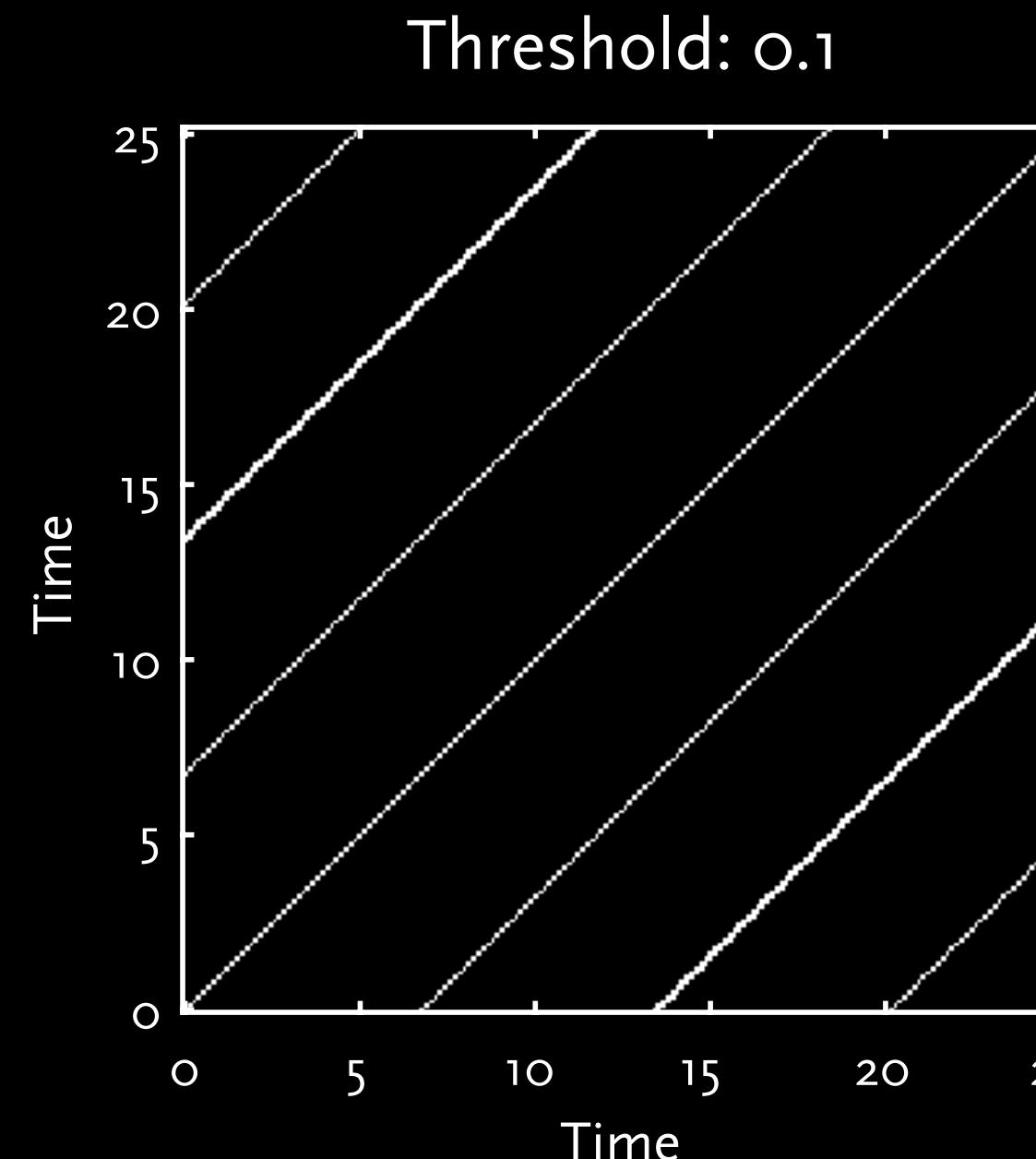
- Minimizing the fragmentation and thickness of the diagonal lines (for cyclical signals)
  - $N_p$  maximal
  - $N_n$  close to  $N_p$

$$\beta(\varepsilon) = \frac{|N_n(\varepsilon) - N_p(\varepsilon)|}{N_n(\varepsilon)}$$

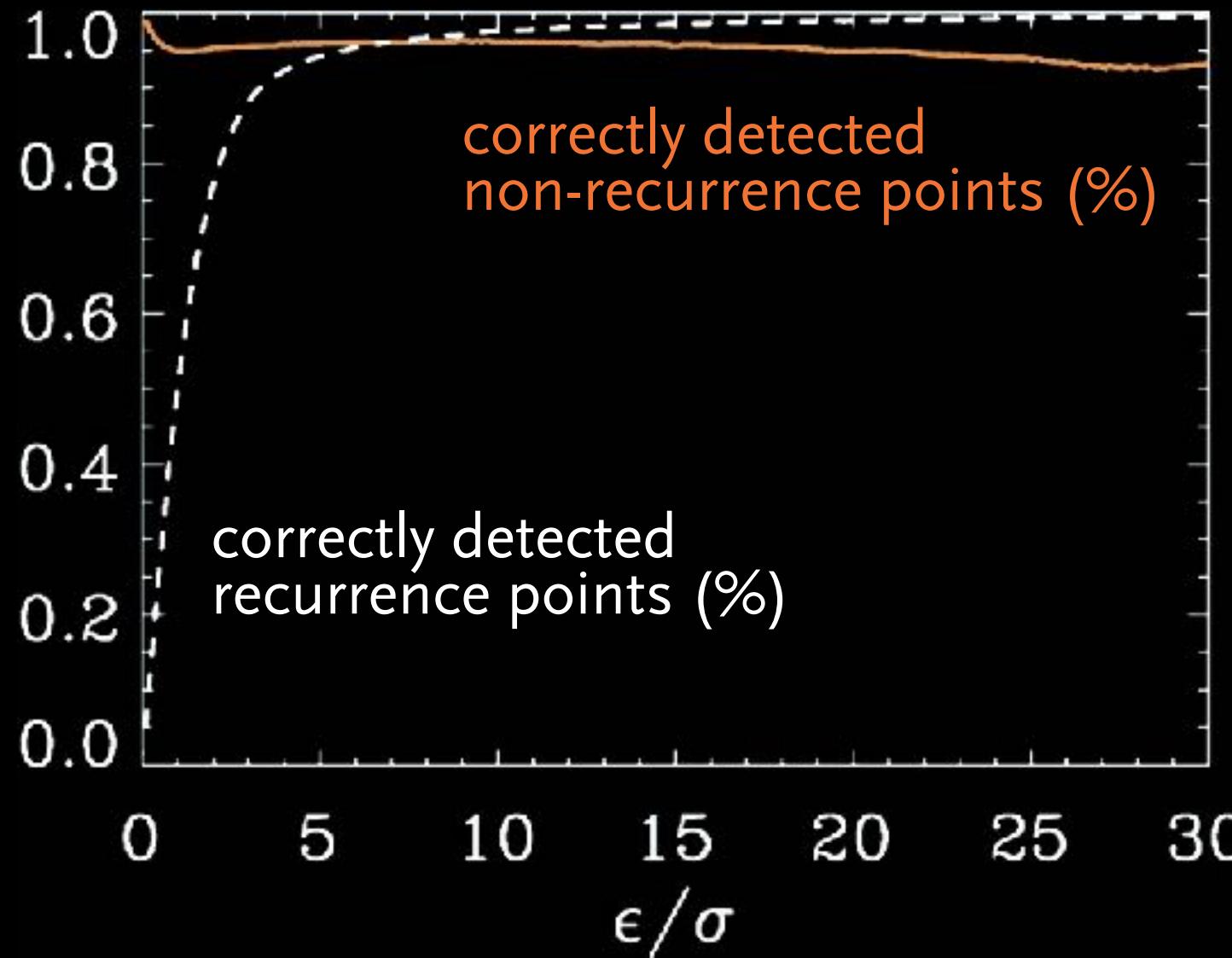
$$\Rightarrow \varepsilon = \arg \min_{\varepsilon} \beta(\varepsilon)$$

# RECURRENCE THRESHOLD: OBSERVATIONAL NOISE

$$x(t) = \hat{x}(t) + \sigma\xi$$



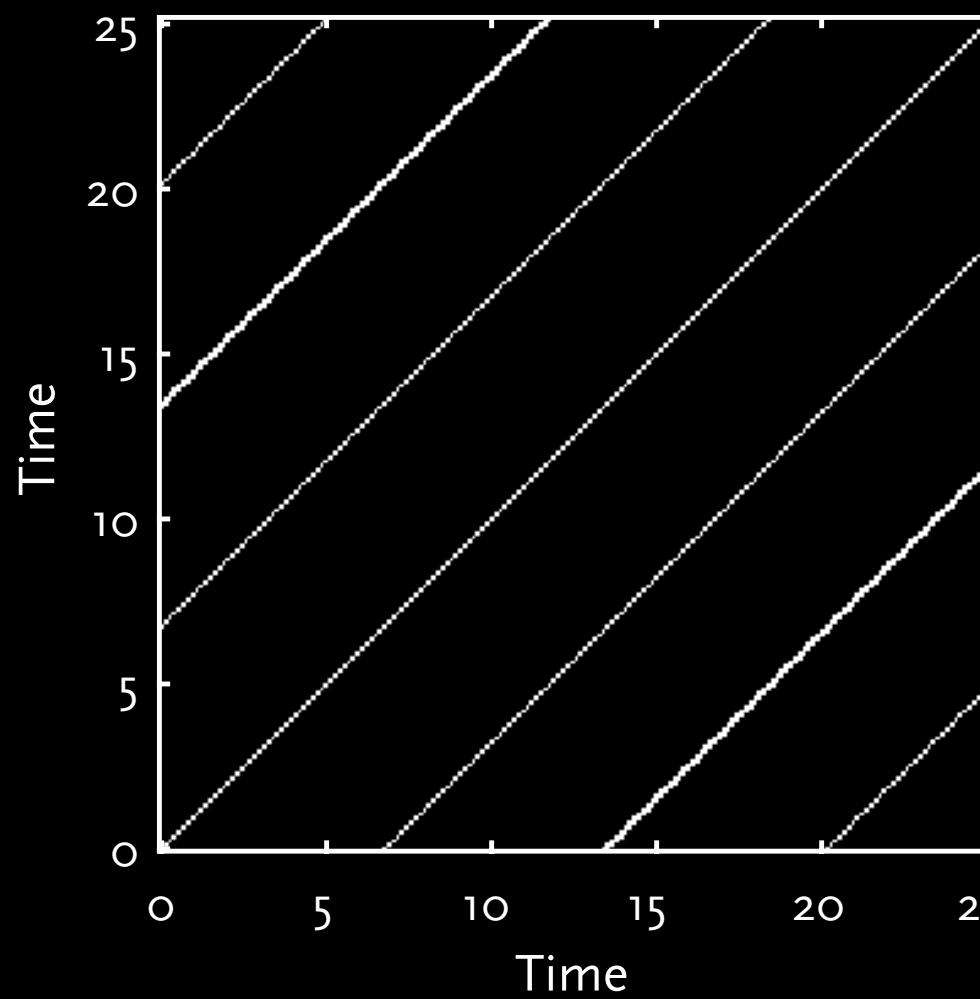
# RECURRENCE THRESHOLD: OBSERVATIONAL NOISE



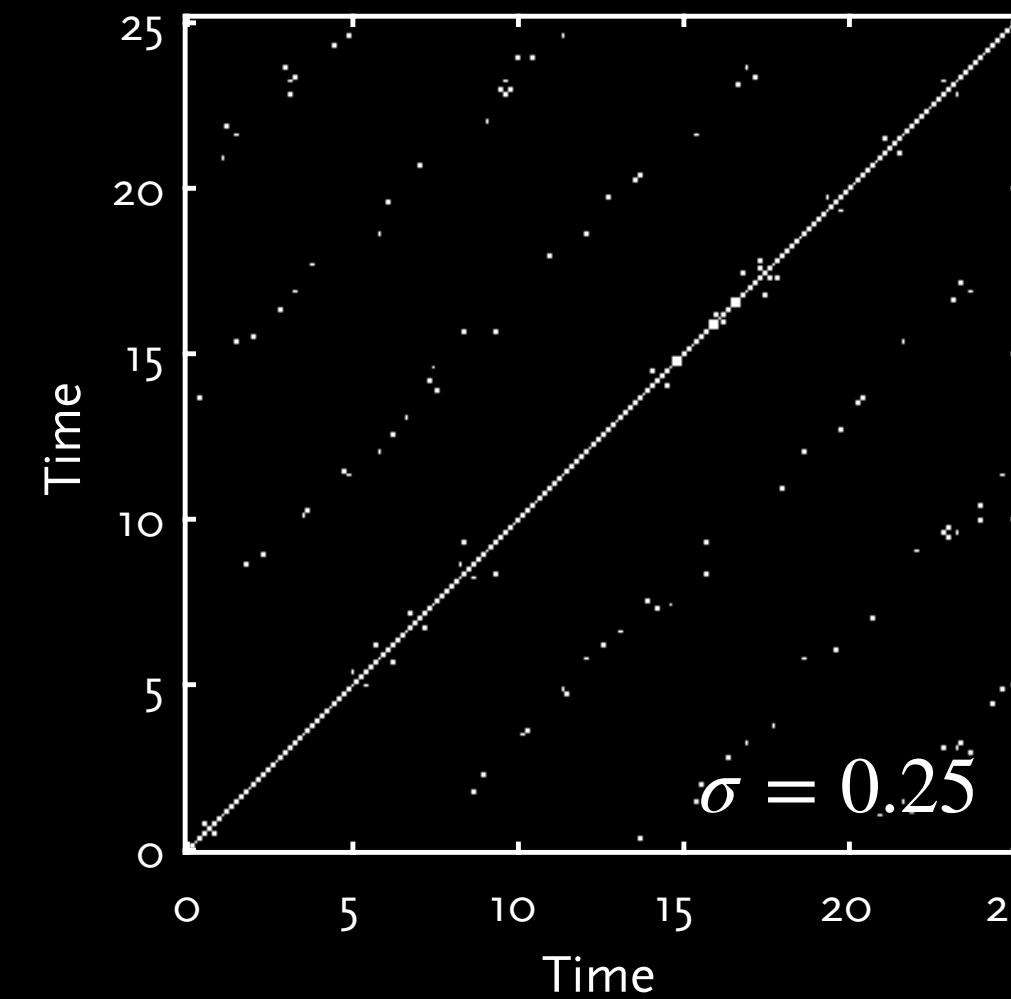
- Observational noise  $\sigma$
- threshold  $\epsilon > 5\sigma$

# RECURRENCE THRESHOLD: OBSERVATIONAL NOISE

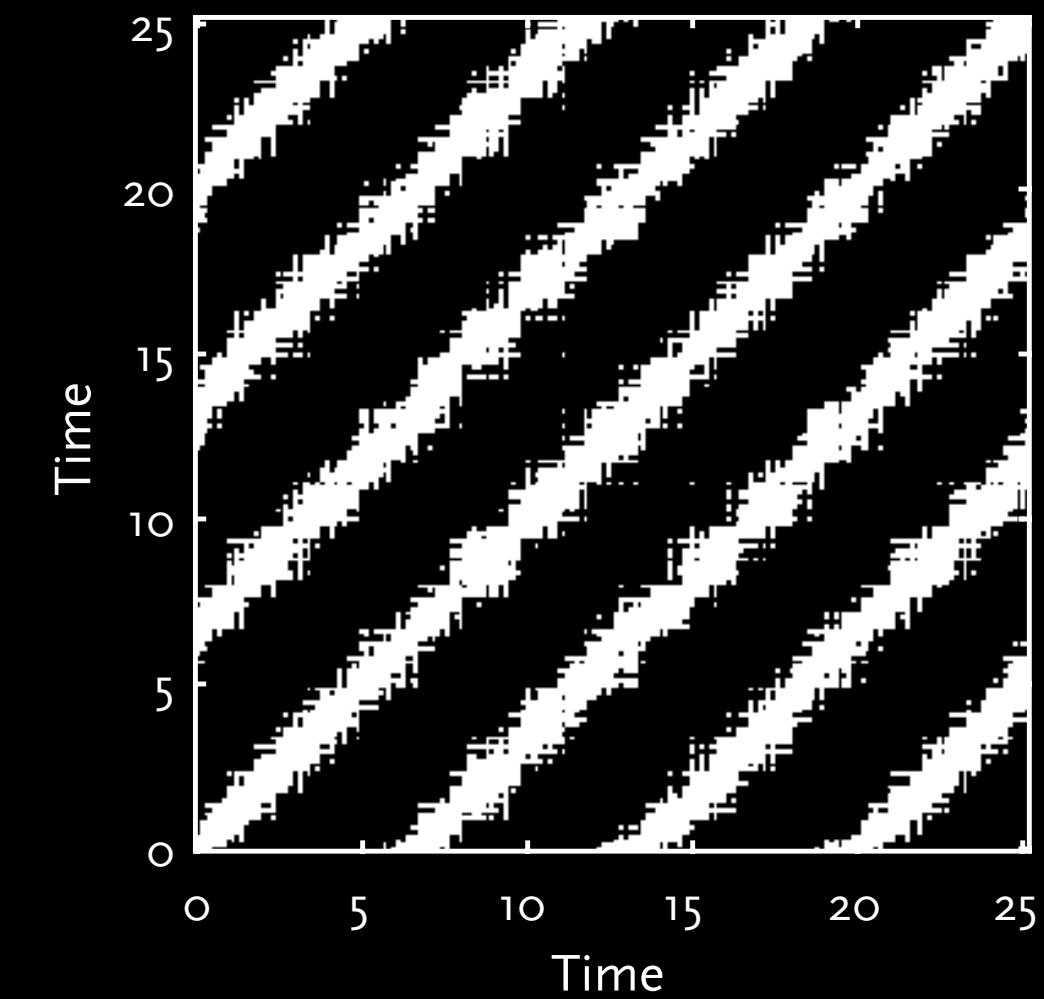
Threshold: 0.1



Threshold: 0.1



Threshold: 1.25



# RECURRENCE THRESHOLD: TOPOLOGICAL SIMILARITY

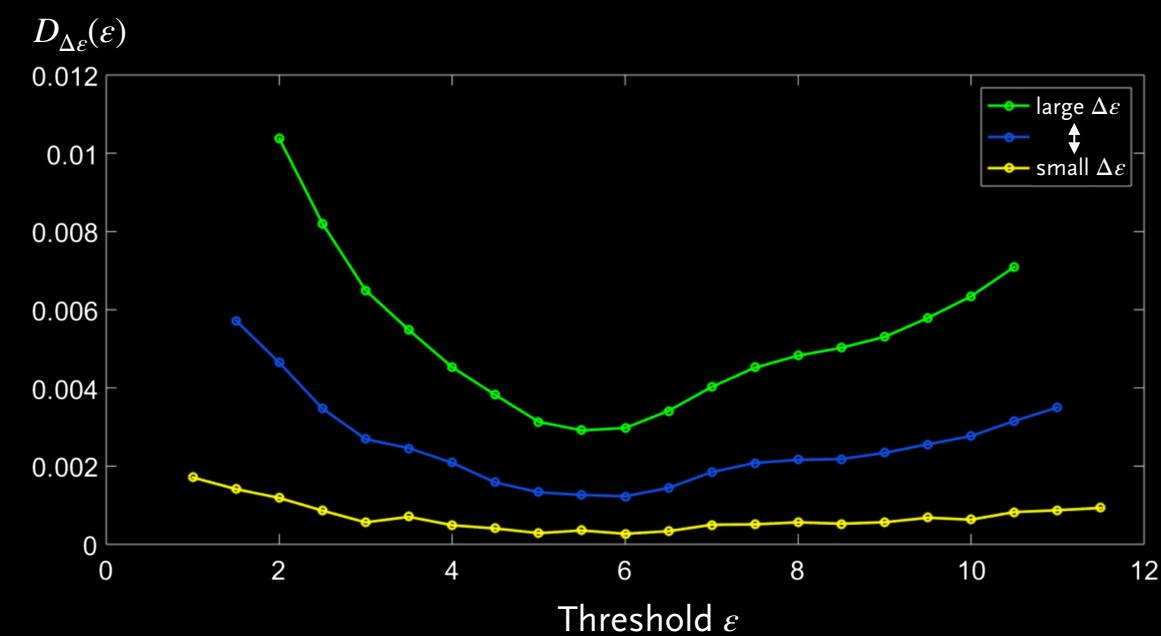
Vega et al., Phys A 445, 2016

- Similarity in number and size of network modules
- Number of modules  
 $C(\mathbf{R}(\varepsilon - \Delta\varepsilon)) = C(\mathbf{R}(\varepsilon)) = C(\mathbf{R}(\varepsilon + \Delta\varepsilon)) > 1$
- Size of module  $k$   
 $|C_k(\mathbf{R}(\varepsilon + \Delta\varepsilon))| - |C_k(\mathbf{R}(\varepsilon))| < \xi$

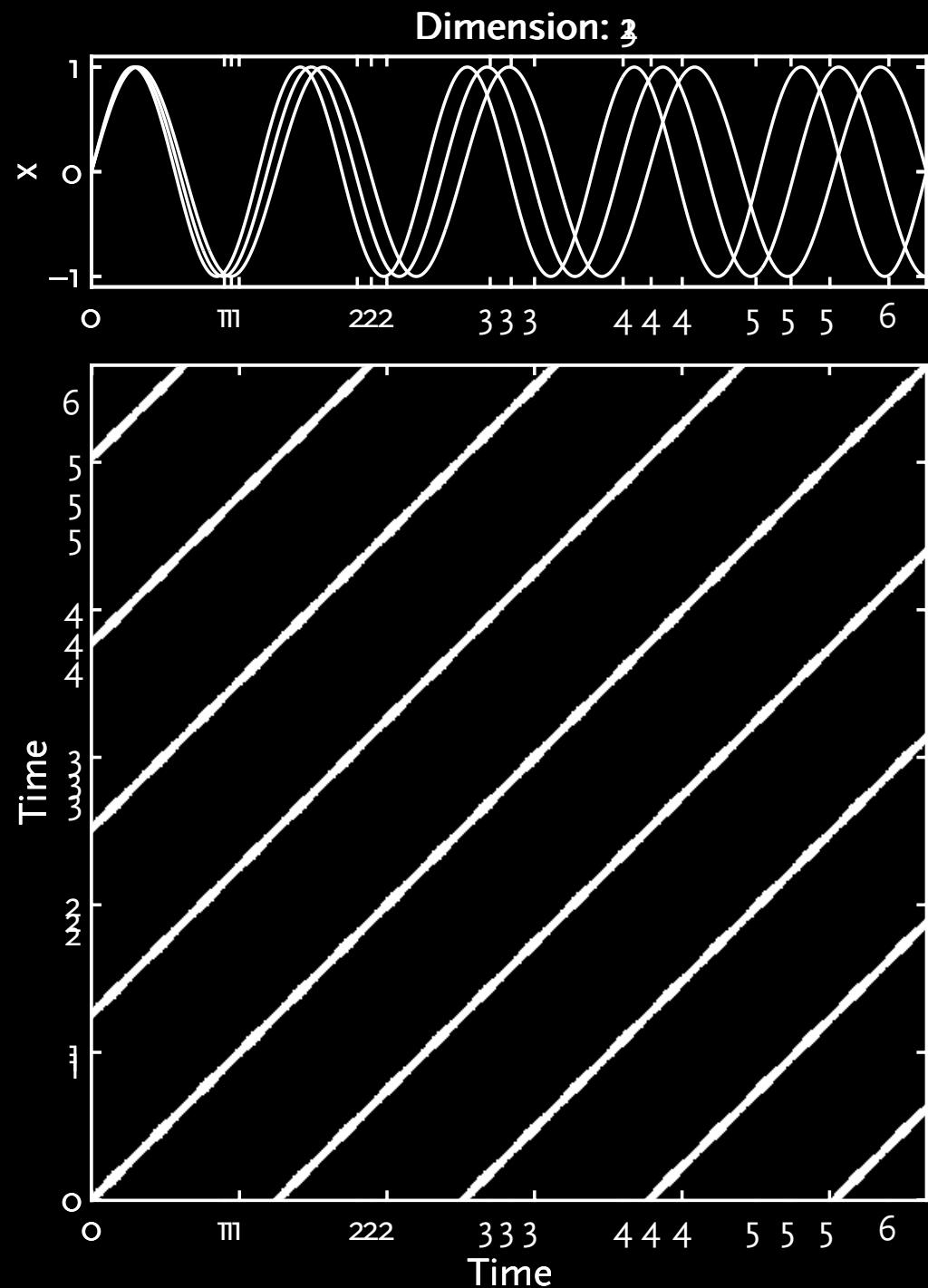
Andreadis et al., Chaos 30, 2020

- Minimise deviation of overall recurrences
- Compare  $\mathbf{R}(\varepsilon)$  with  $\mathbf{R}(\varepsilon + \Delta\varepsilon)$ :

$$D_{\Delta\varepsilon}(\varepsilon) = |\text{dist}(\mathbf{R}(\varepsilon + \Delta\varepsilon), \mathbf{R}(\varepsilon - \Delta\varepsilon))|$$



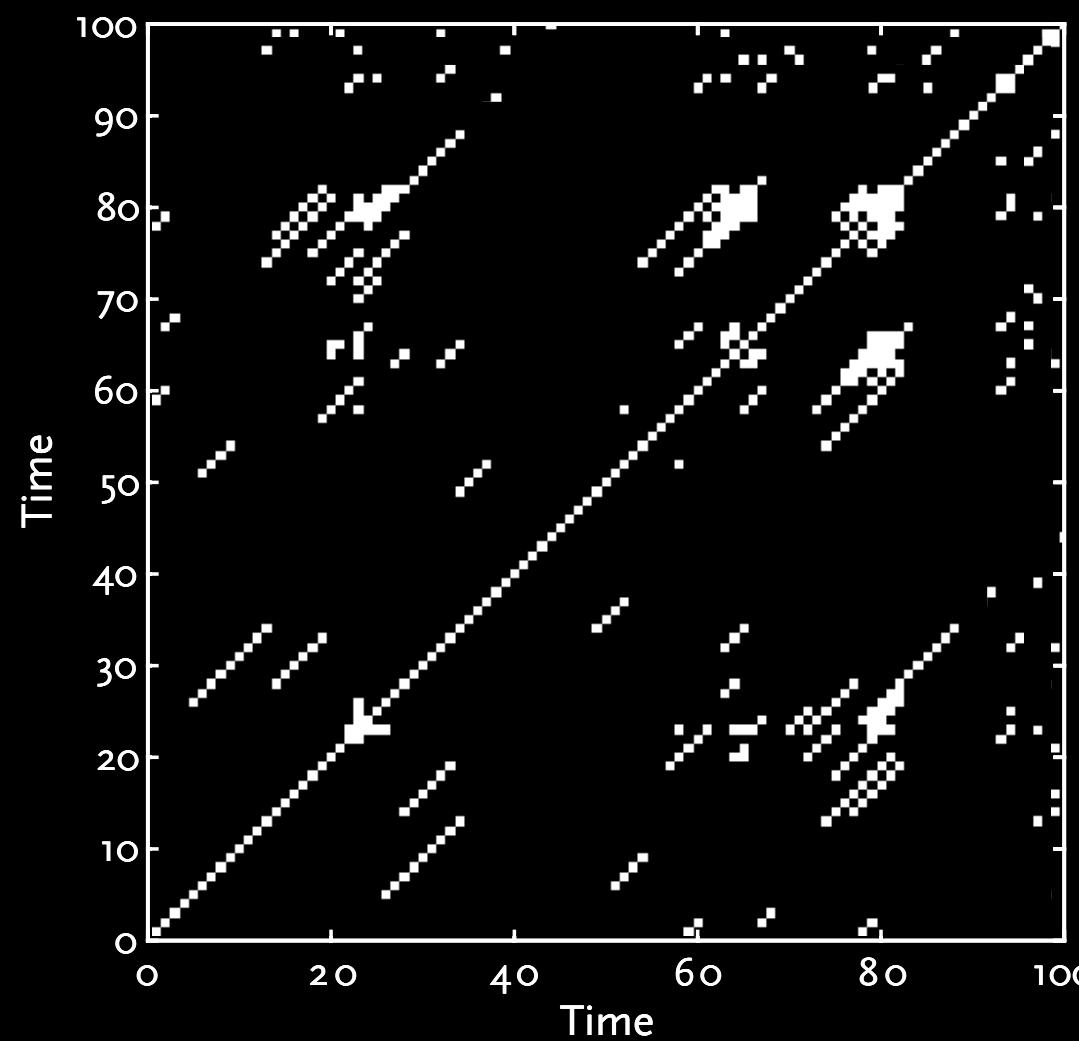
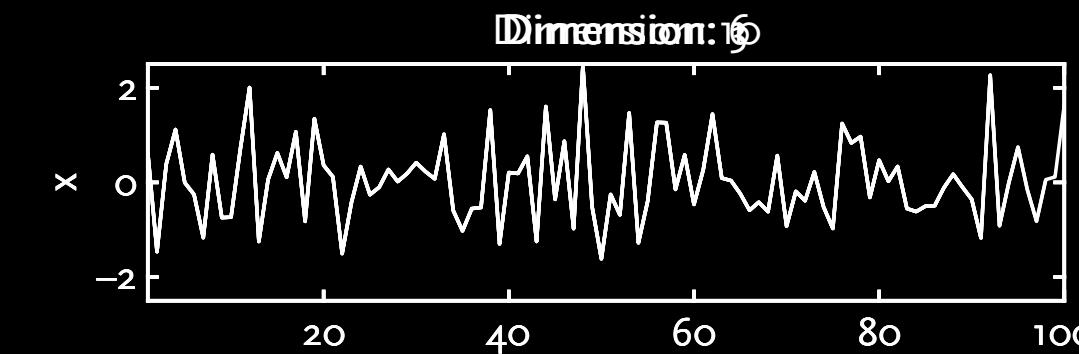
# INFLUENCE OF EMBEDDING DIMENSION



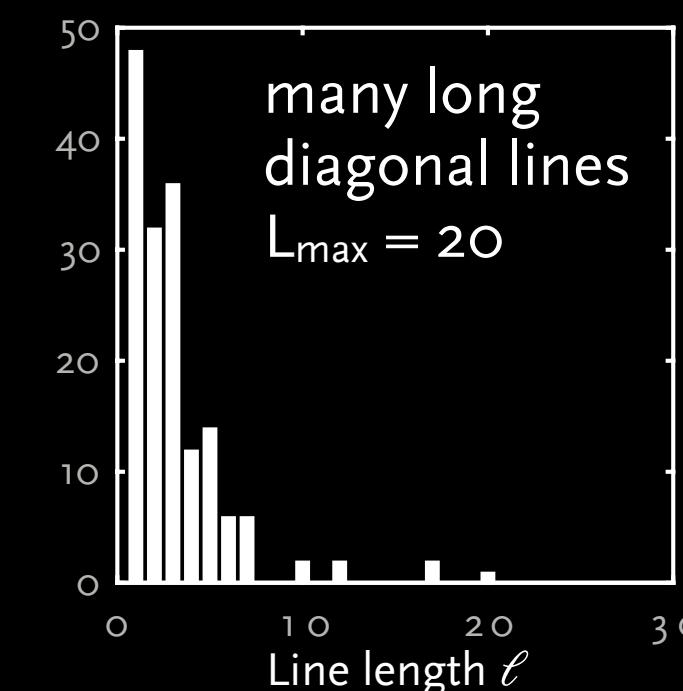
- Too low embedding dimension:
  - attractor (phase space trajectory) is not enough unfolded
  - trajectory segments cross each other or even evolve in opposite direction
- Too high embedding dimension:
  - over-embedding usually not a problem (within limits)



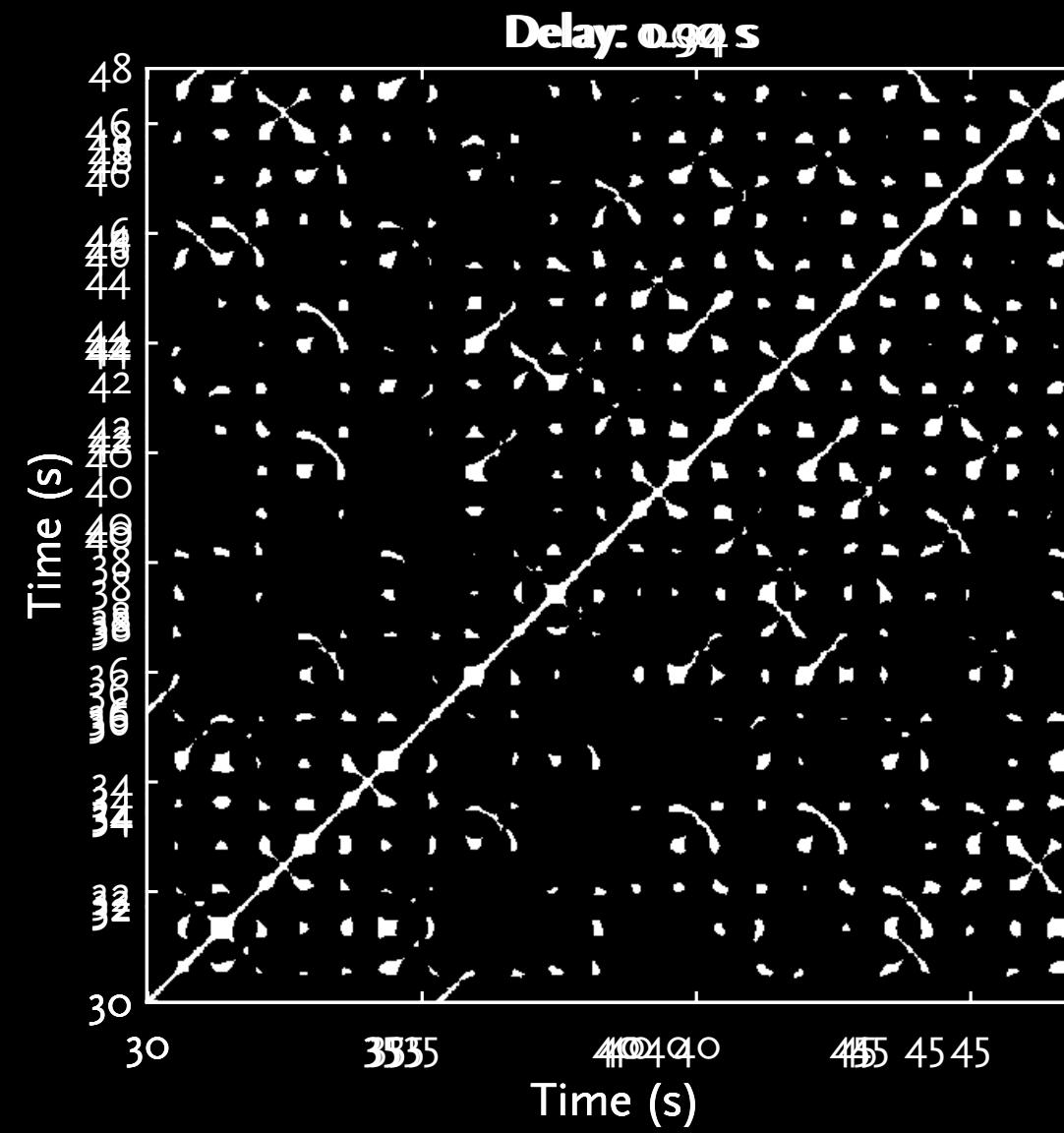
# INFLUENCE OF EMBEDDING DIMENSION



- Embedding can cause spurious correlations
- Artificially increase of diagonal line lengths (when using adaptive threshold)



# INFLUENCE OF EMBEDDING DELAY



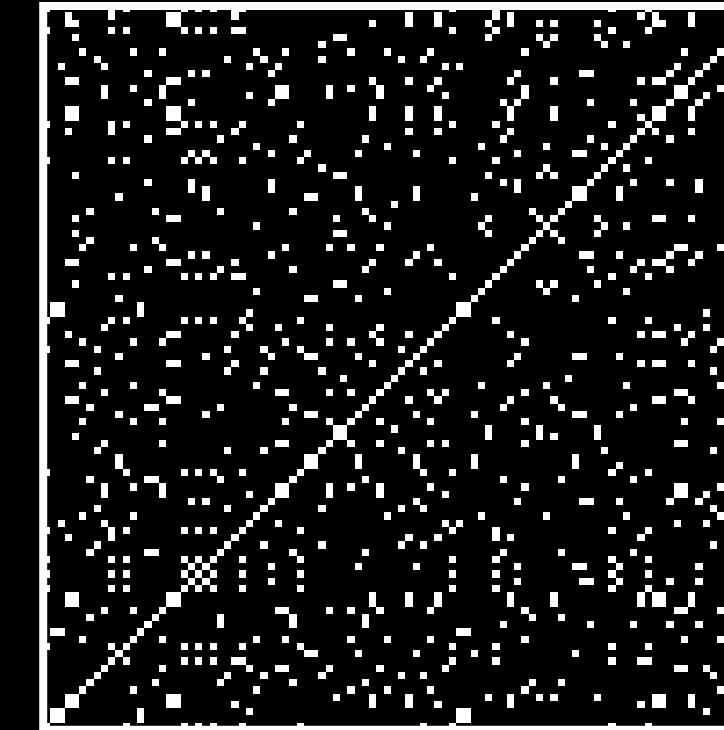
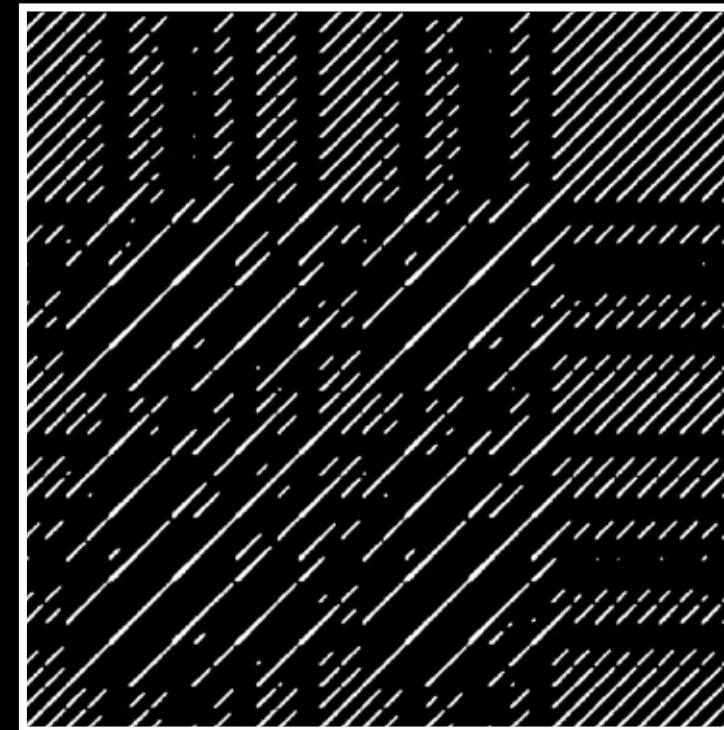
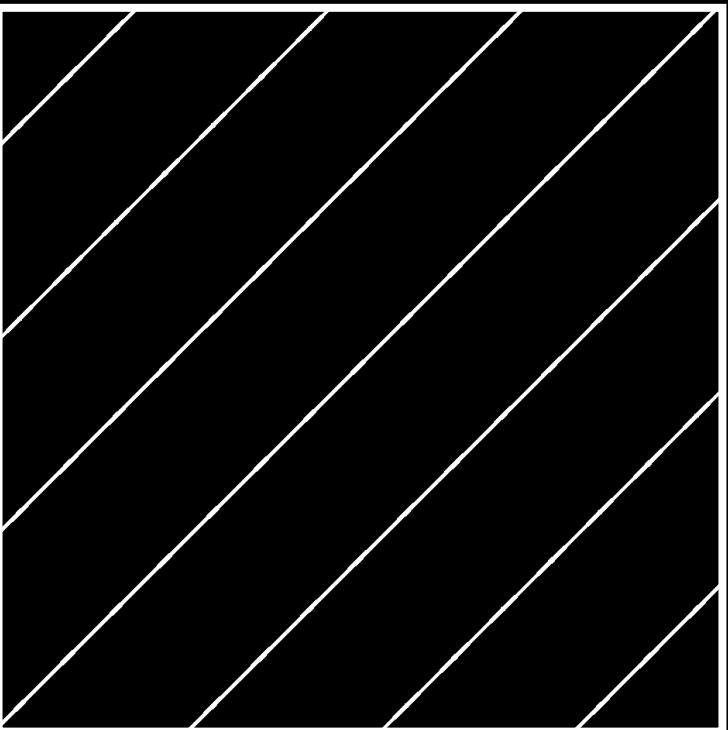
Lorenz system

- Embedding delay necessary to unfold attractor
- Can destroy line structures
- Optimal delay: mainly non-interrupted line structures

# RECURRENCE QUANTIFICATION ANALYSIS

# RECURRENCE QUANTIFICATION ANALYSIS

- 1992, 1994, Webber & Zbilut:  
Quantify the visually apparent differences in the RP



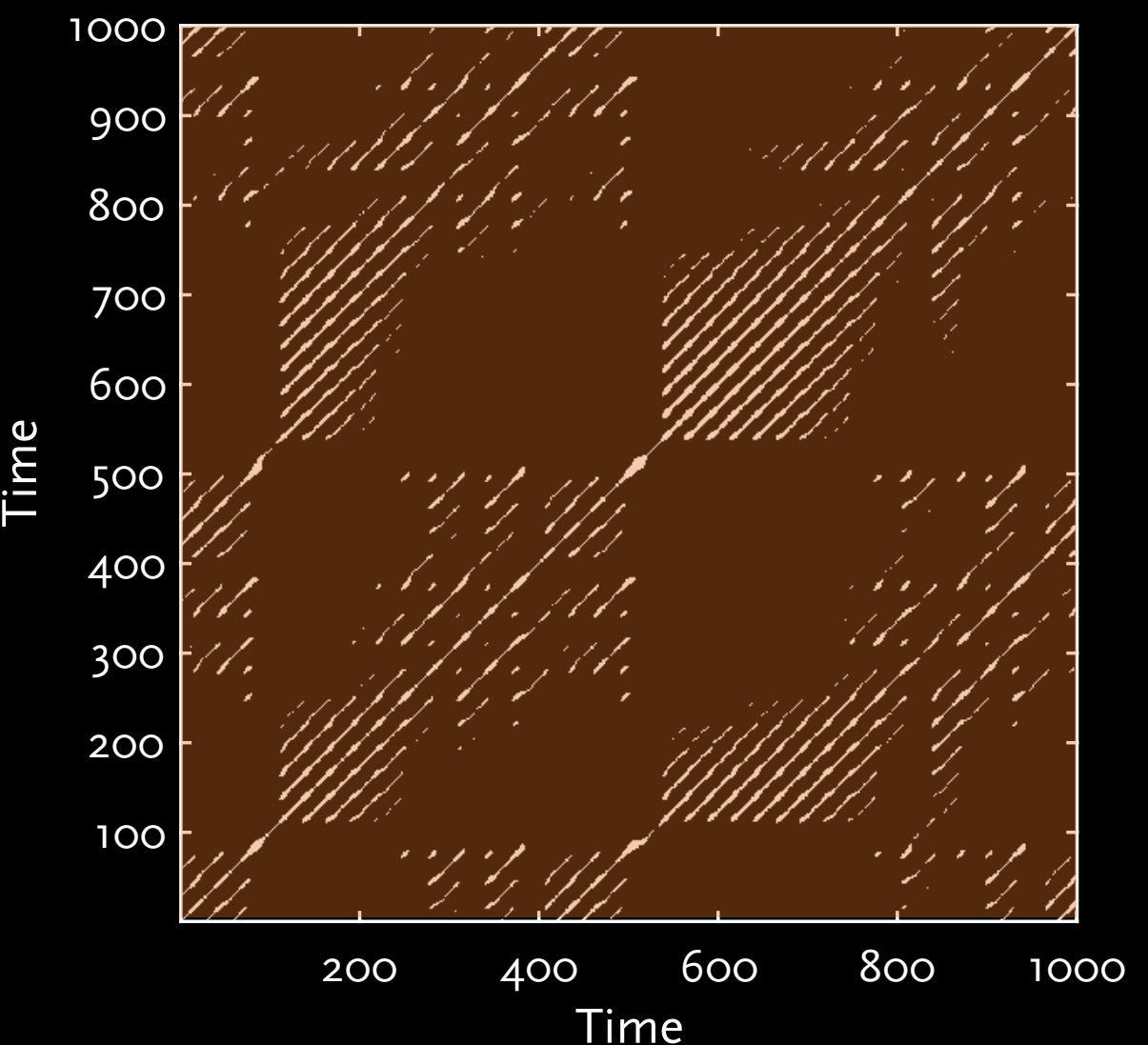
# RECURRENCE RATE

- Density of recurrence points in whole RP

$$RR(\varepsilon, m) = \frac{1}{N_m(N_m - 1)} \sum_{i=1}^{N_m} \sum_{\substack{j=1 \\ j \neq i}}^{N_m} R_{i,j}(\varepsilon, m)$$

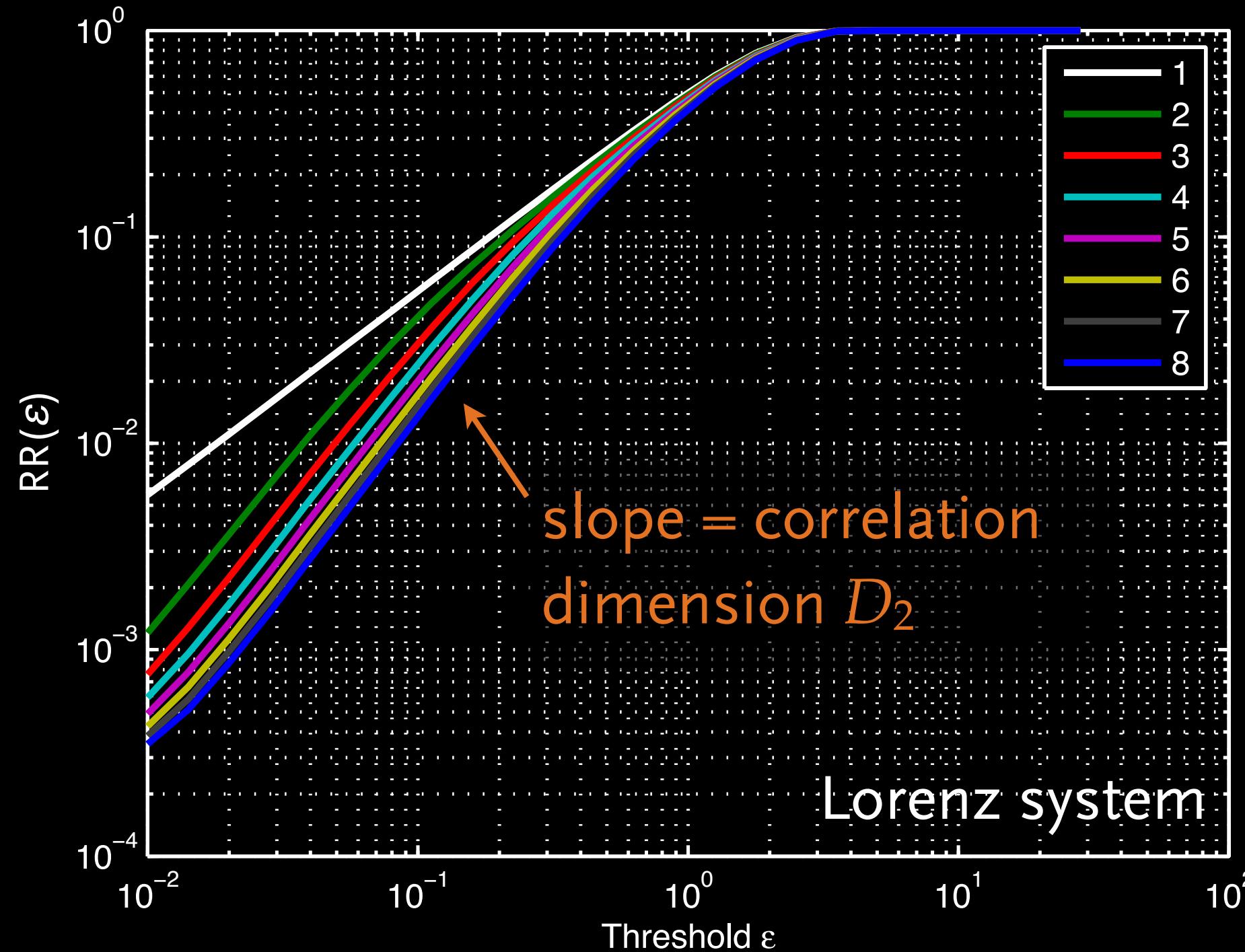
Probability that the system will return  
to any (randomly selected) former state

- Correlation sum





# RECURRENCE RATE



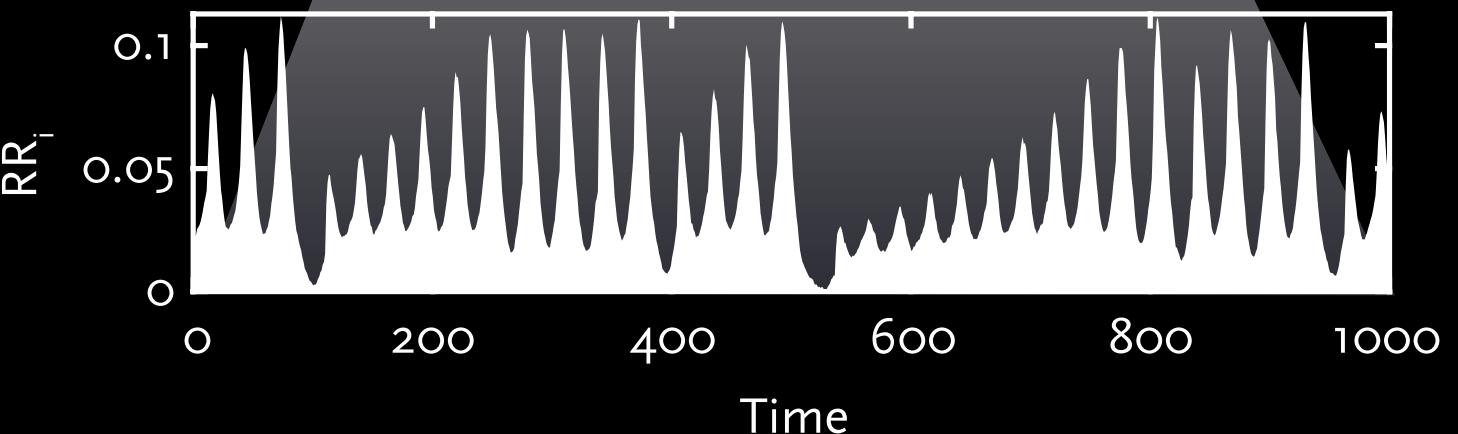
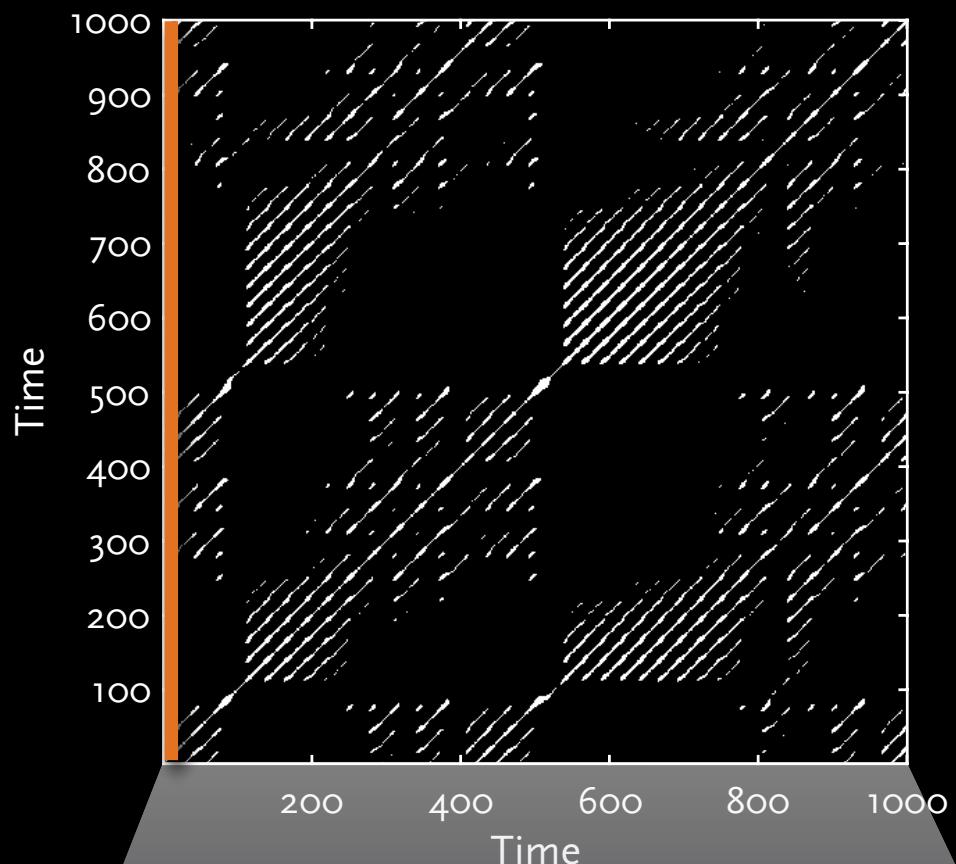
- Useful when testing scaling behaviour
- RR alone closely related to variance of the signal

# COLUMN-WISE RECURRENCE RATE

- Density of recurrence points per column

$$RR_i(\varepsilon, m) = \frac{1}{N_m - 1} \sum_{\substack{j=1 \\ j \neq i}}^{N_m} R_{i,j}(\varepsilon, m)$$

Probability of state at time point  $i$



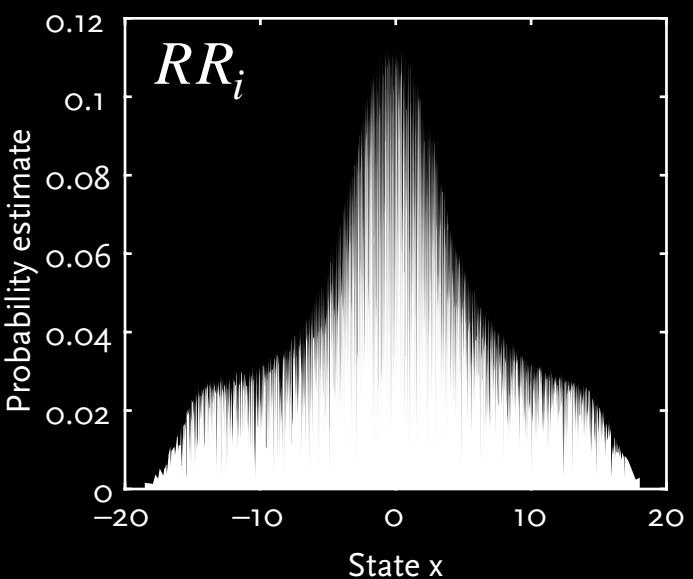
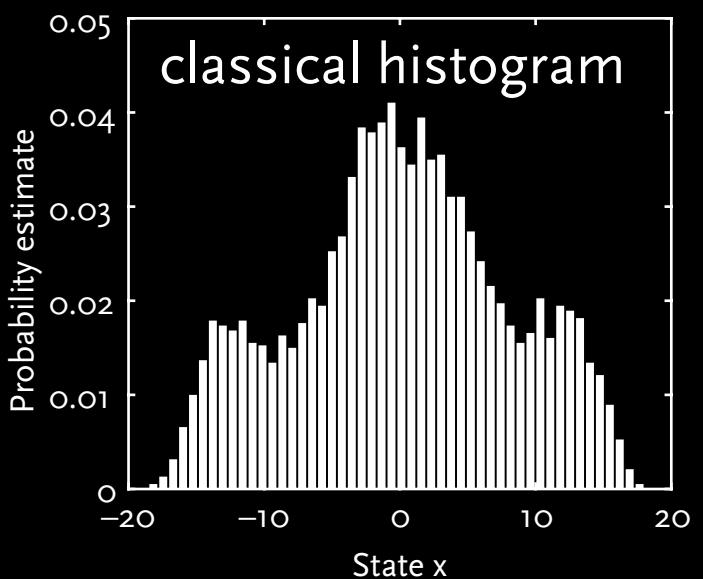
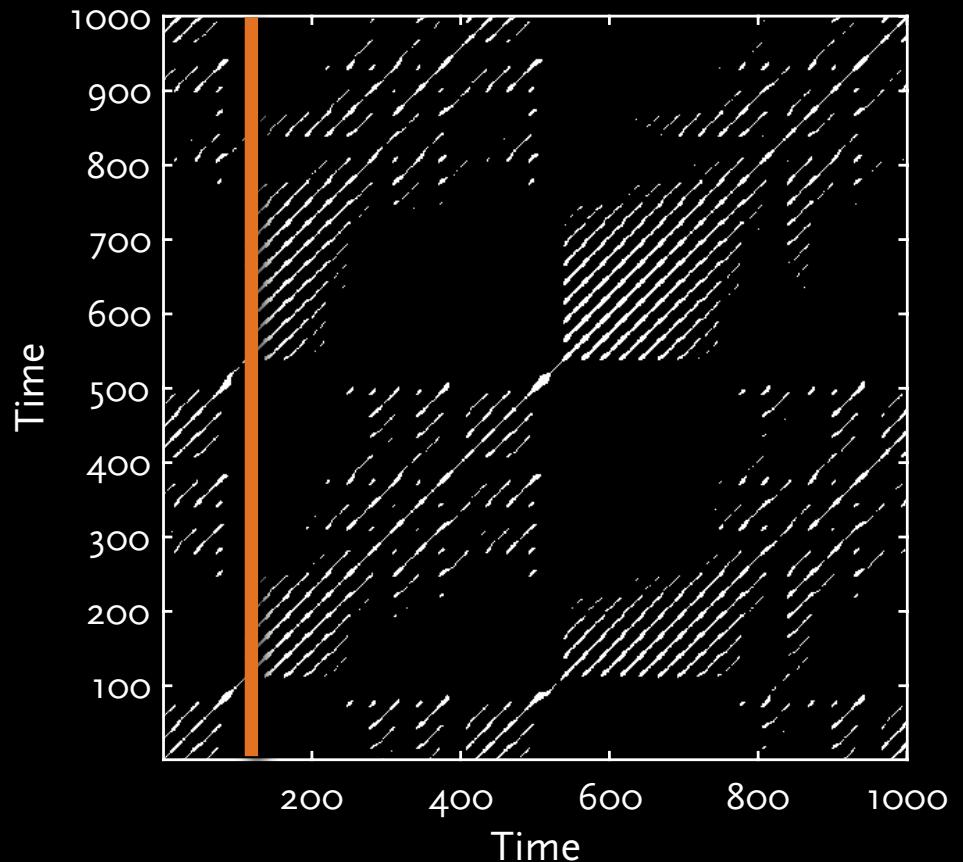
# COLUMN-WISE RECURRENCE RATE

- Density of recurrence points per column

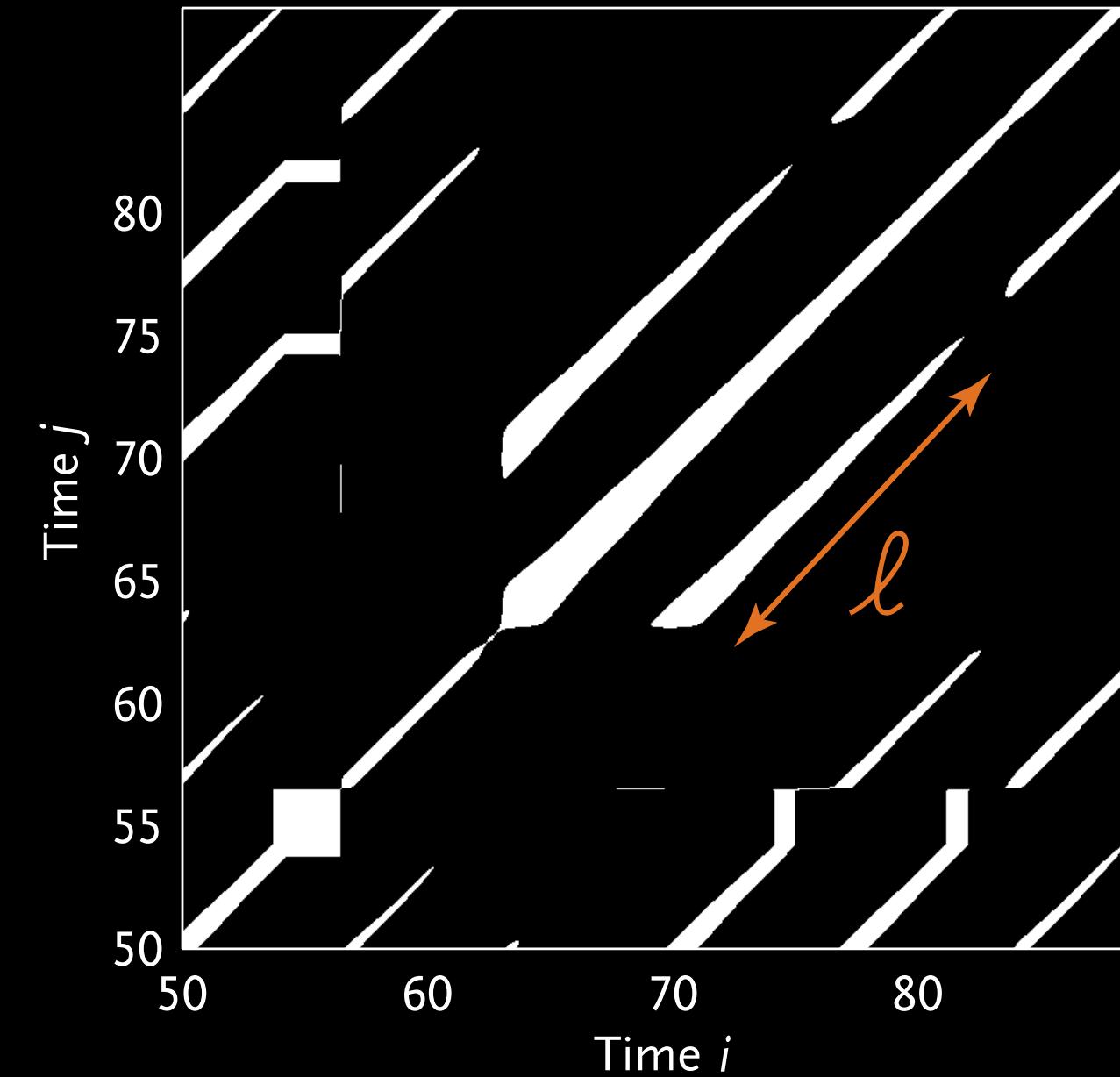
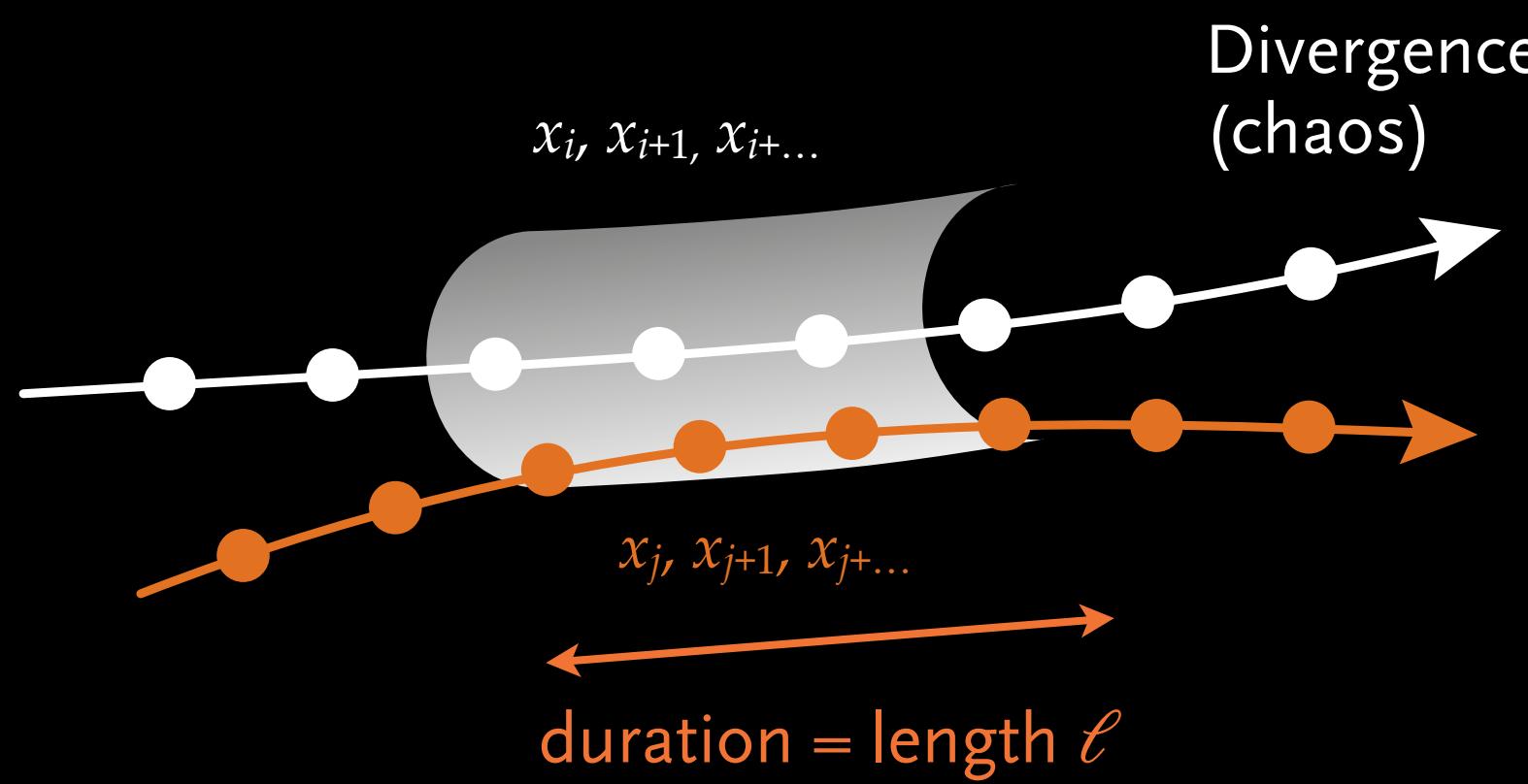
$$RR_i(\varepsilon, m) = \frac{1}{N_m - 1} \sum_{\substack{j=1 \\ j \neq i}}^{N_m} R_{i,j}(\varepsilon, m)$$

Probability of state at time point  $i$

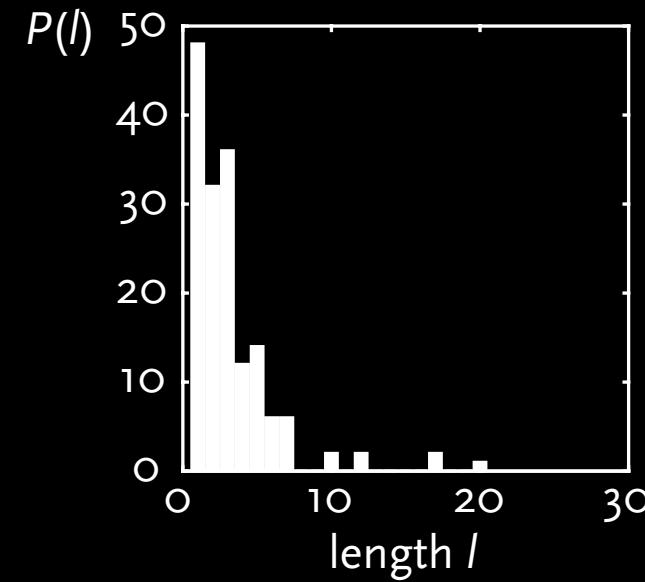
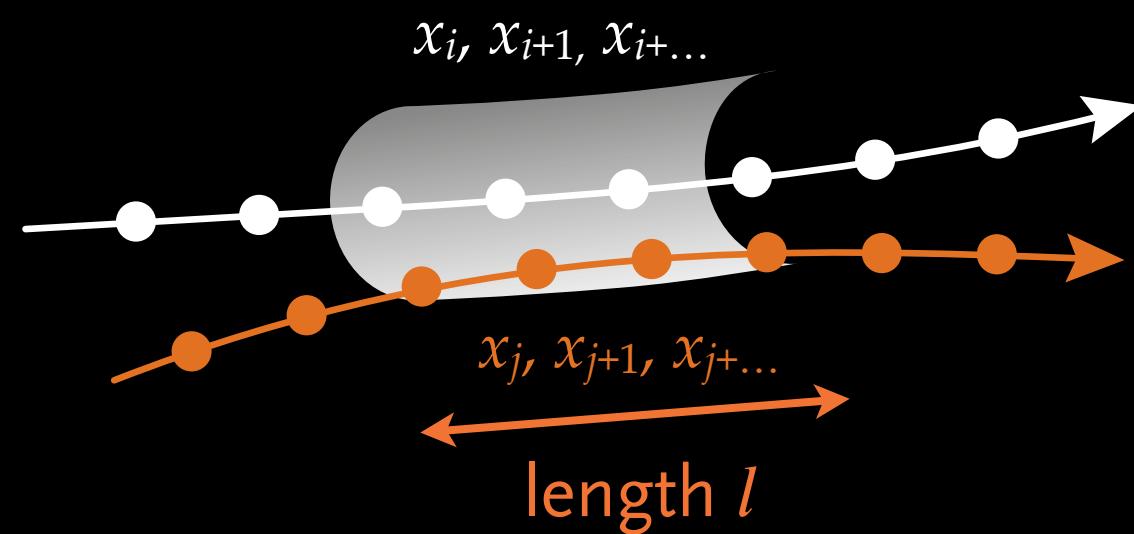
- Alternative probability estimator for many probability based methods, such as mutual information



# DIAGONAL LINES



# DIAGONAL LINE ENTROPY



- Shannon entropy of diagonal line length distribution

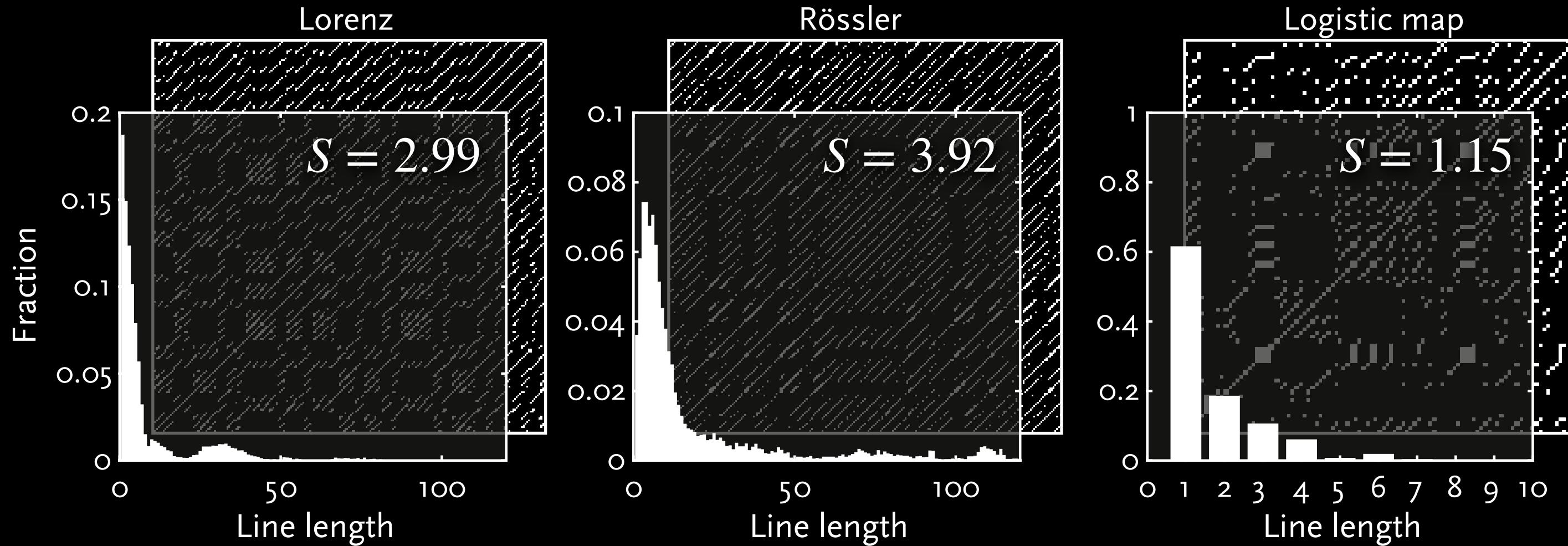
$$S = - \sum_{l=1}^N p(l) \ln p(l)$$

“... the more complex the deterministic structure of the recurrence plot, the larger the [entropy].”

# DIAGONAL LINE ENTROPY

- Entropy of line length distribution  $p(l)$

$$S = - \sum_{l=1}^N p(l) \ln p(l)$$



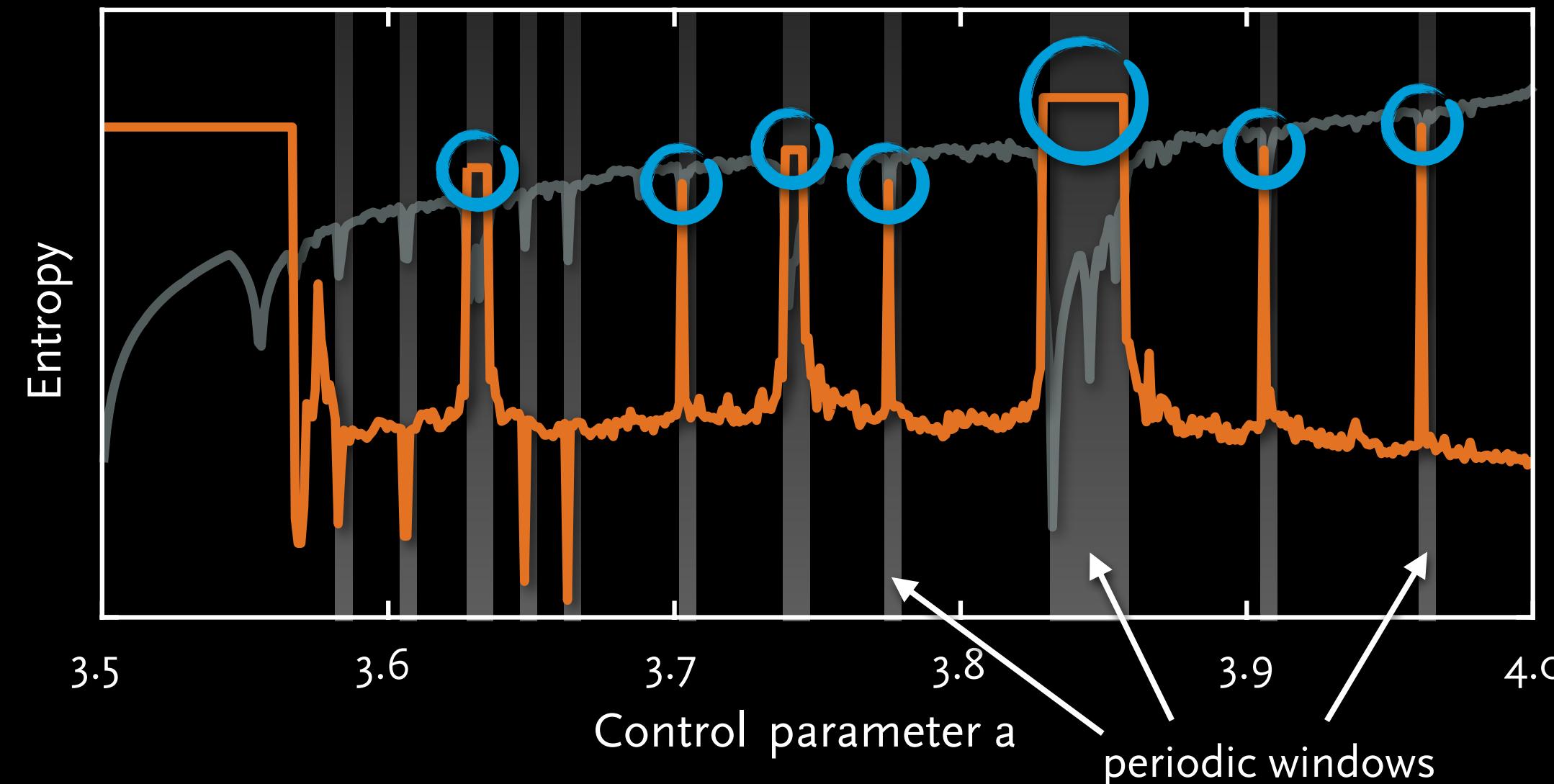


# DIAGONAL LINE ENTROPY

Larger entropy = more complex dynamics

Counterintuitive results?

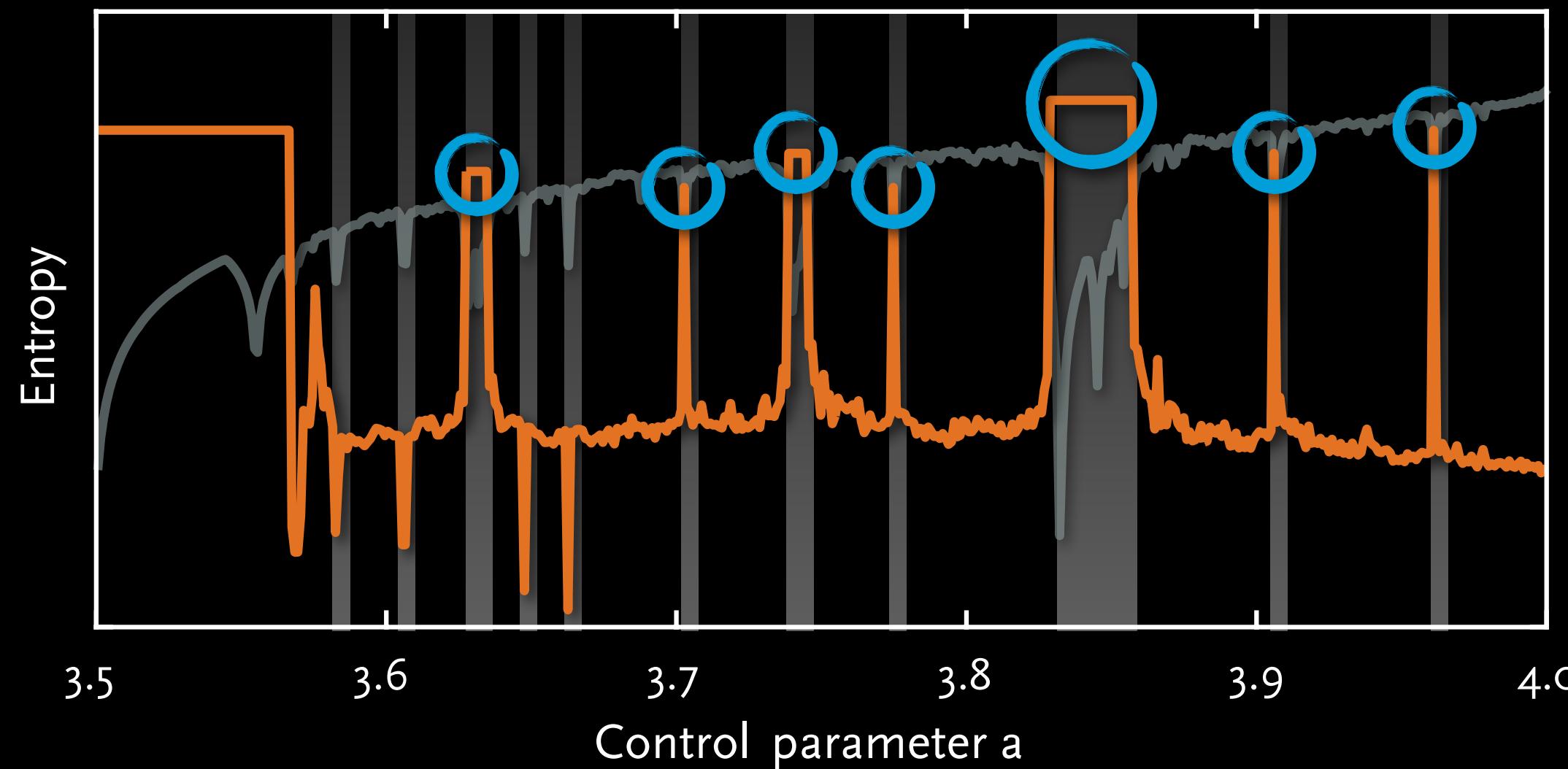
Large values  
for periodic  
dynamics



Q u i z Z

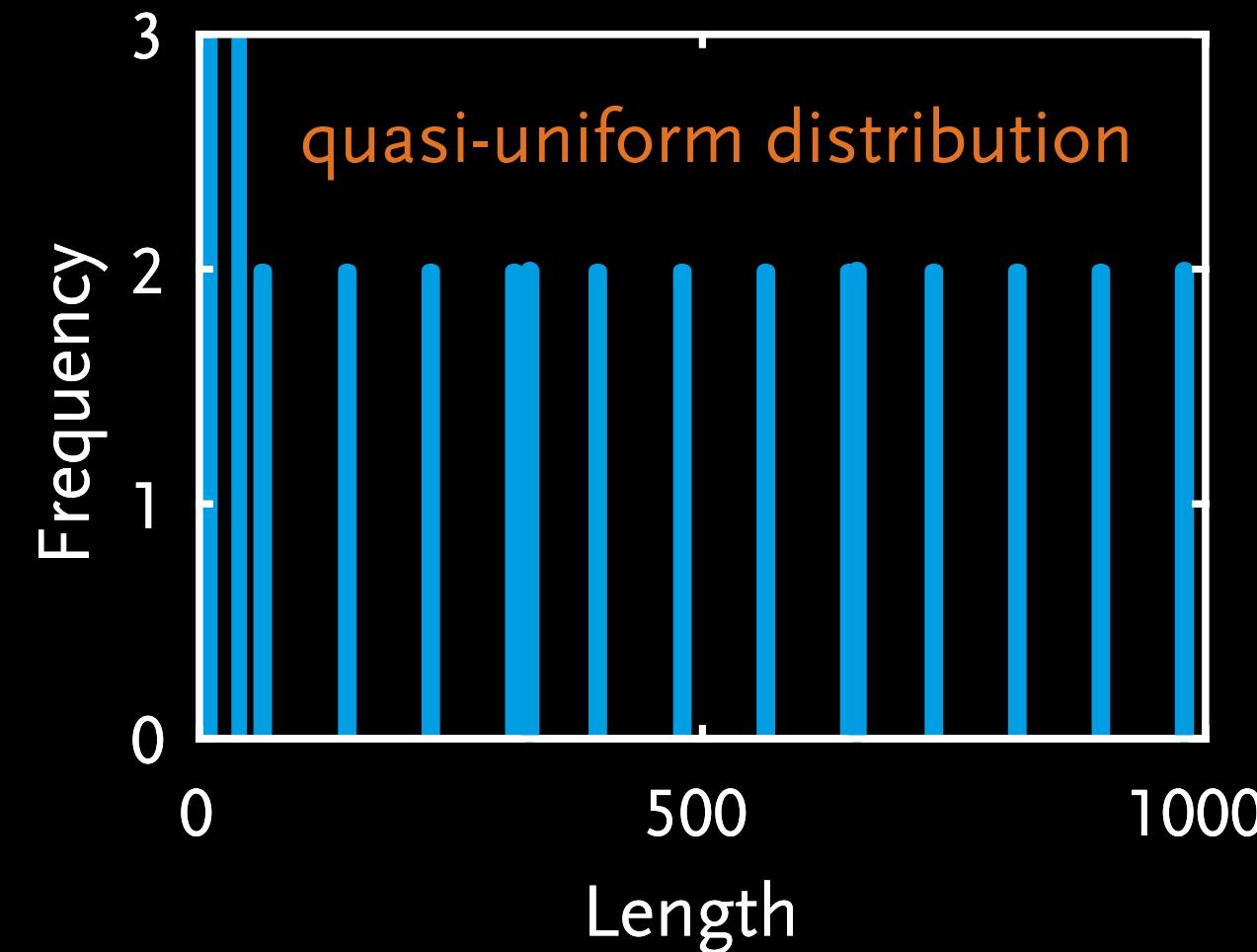
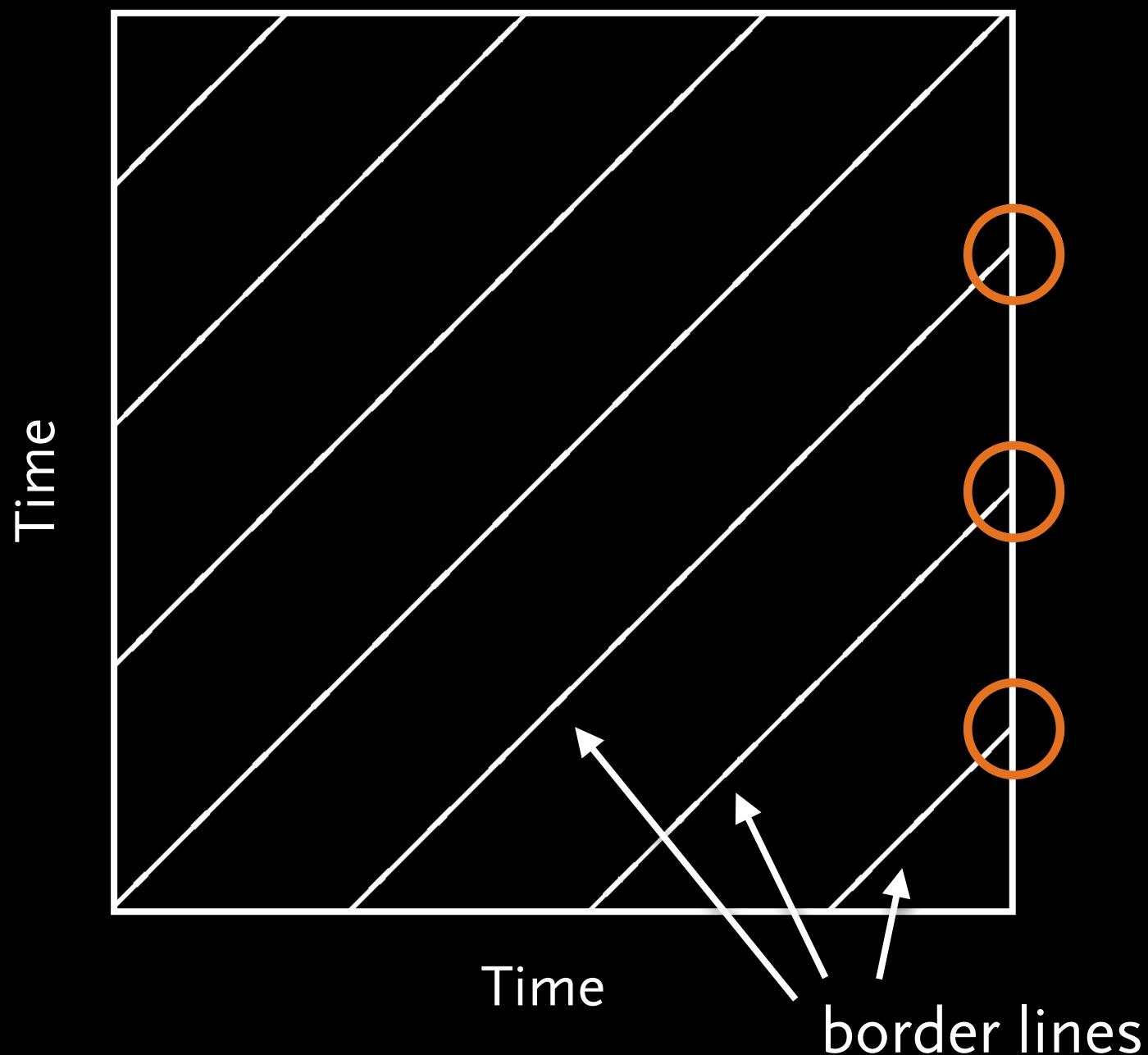
# Quizz

- Why large entropy values for periodic regimes?



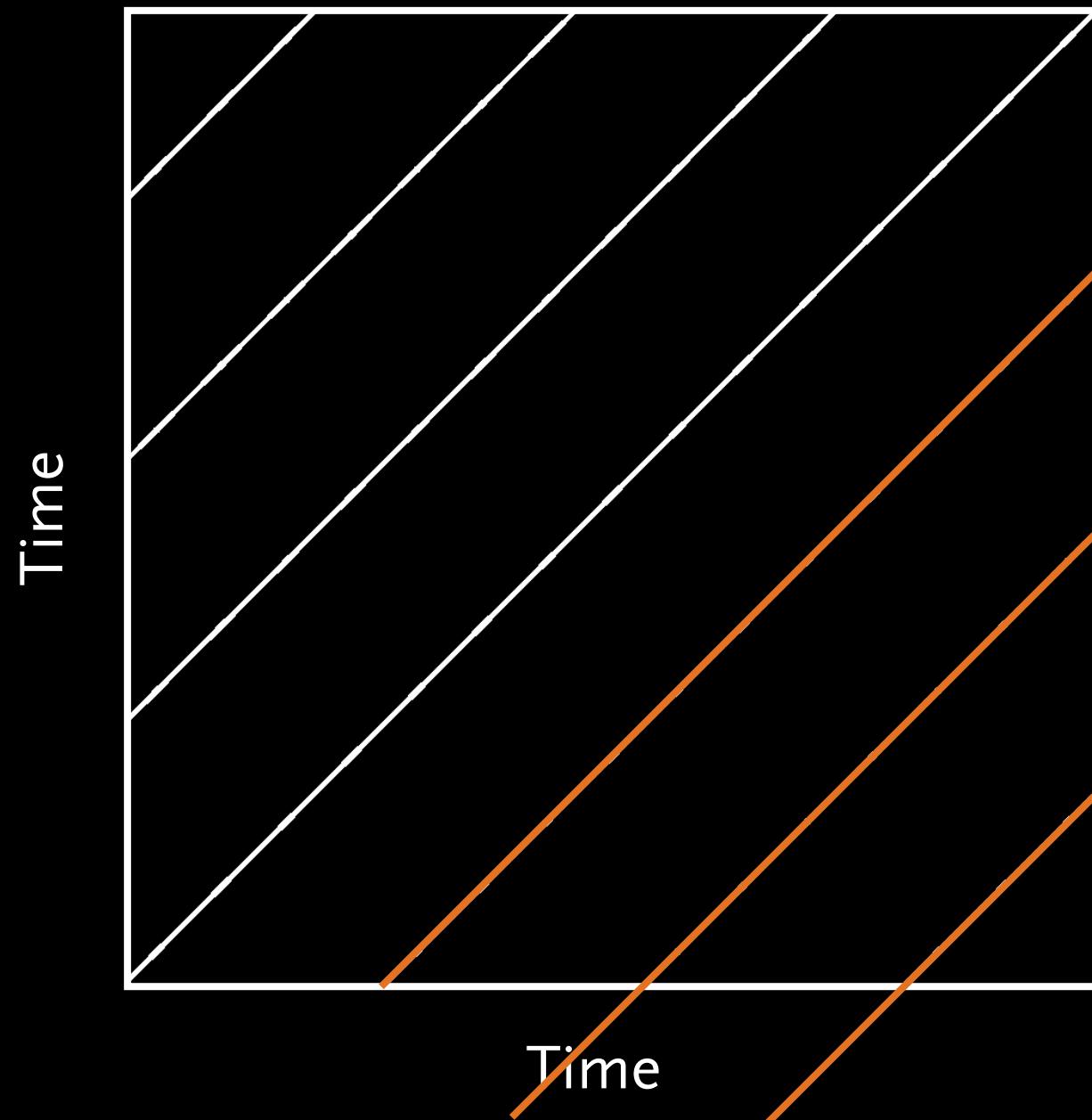


# DIAGONAL LINES FOR PERIODIC DYNAMICS

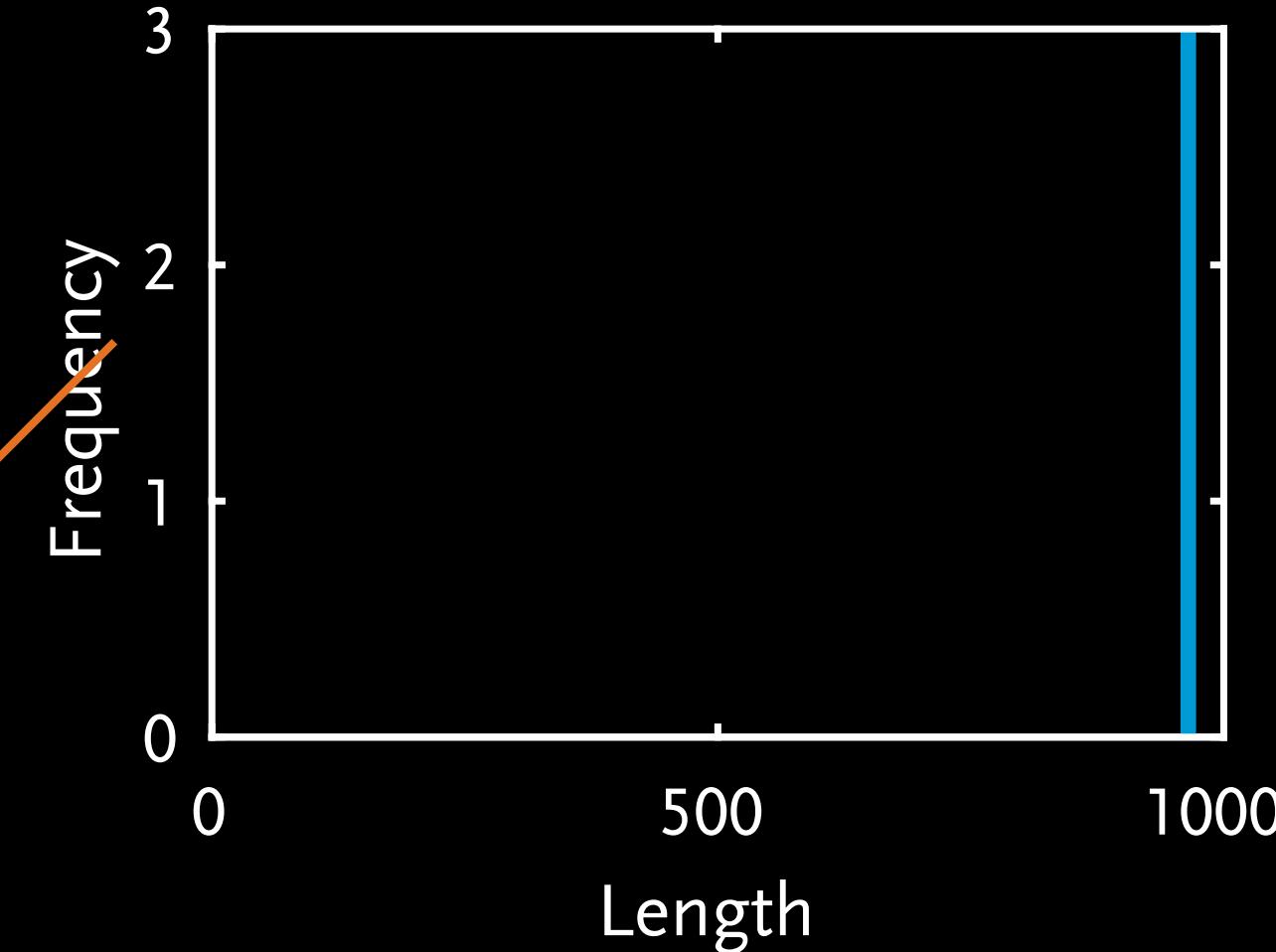




# CORRECTION OF DIAGONAL LINE LENGTHS



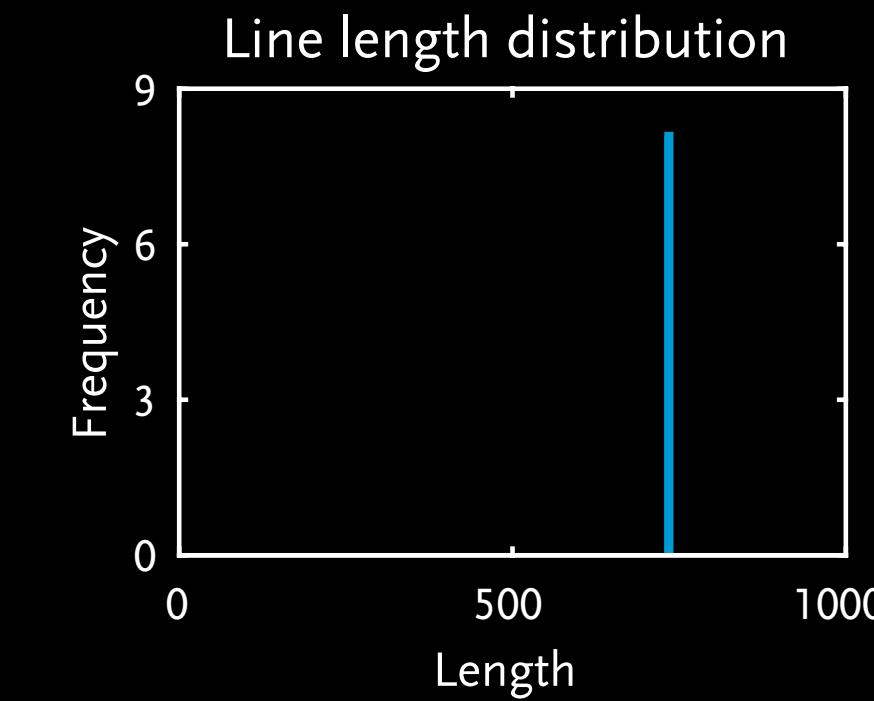
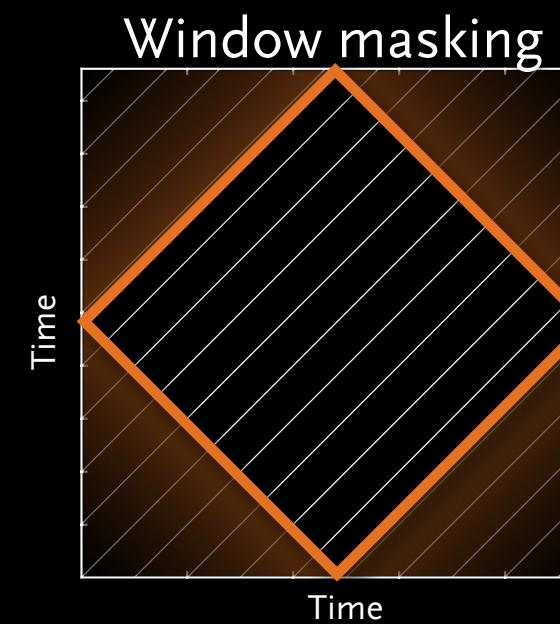
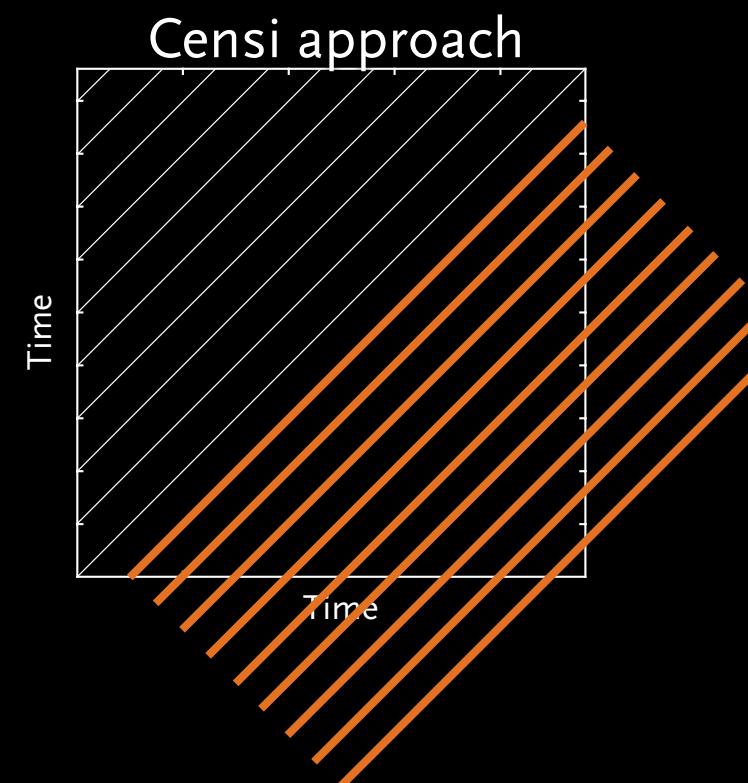
- Replace lengths of border lines by **longest** length





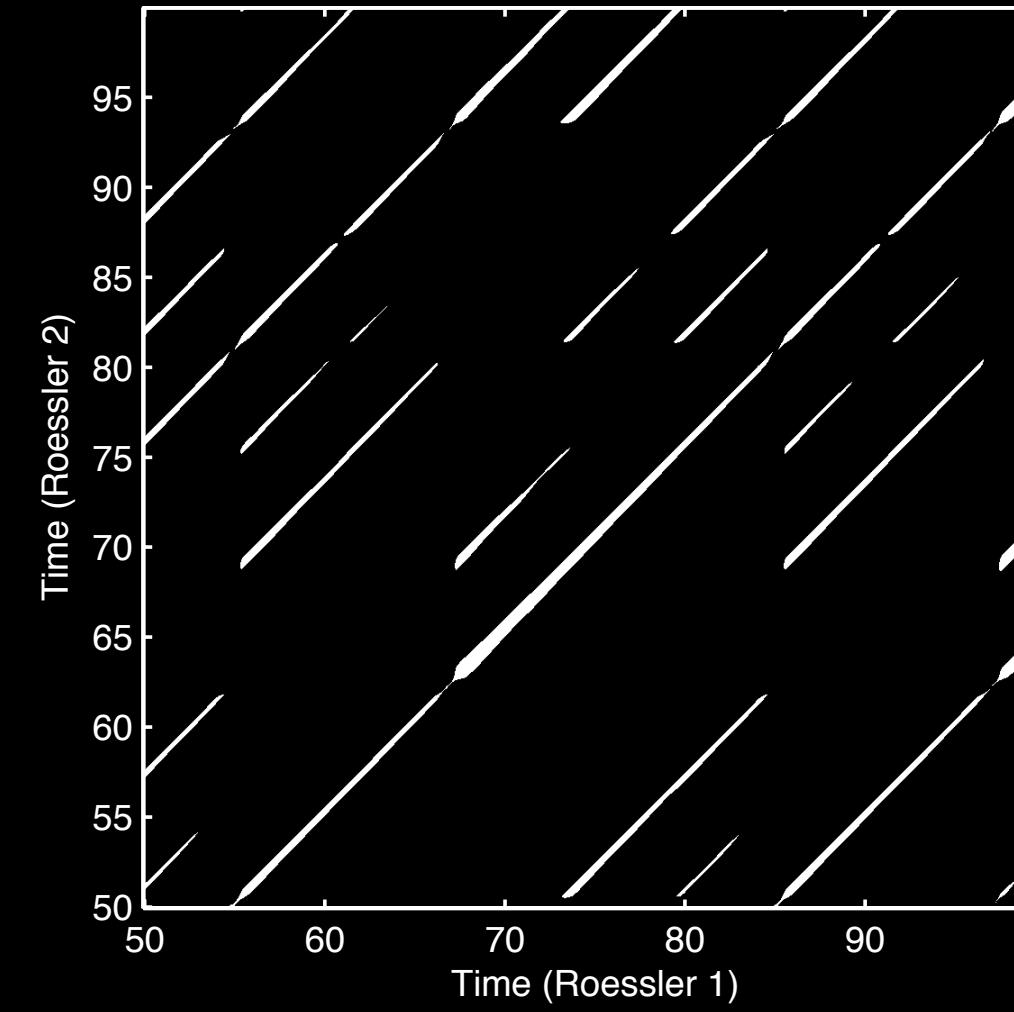
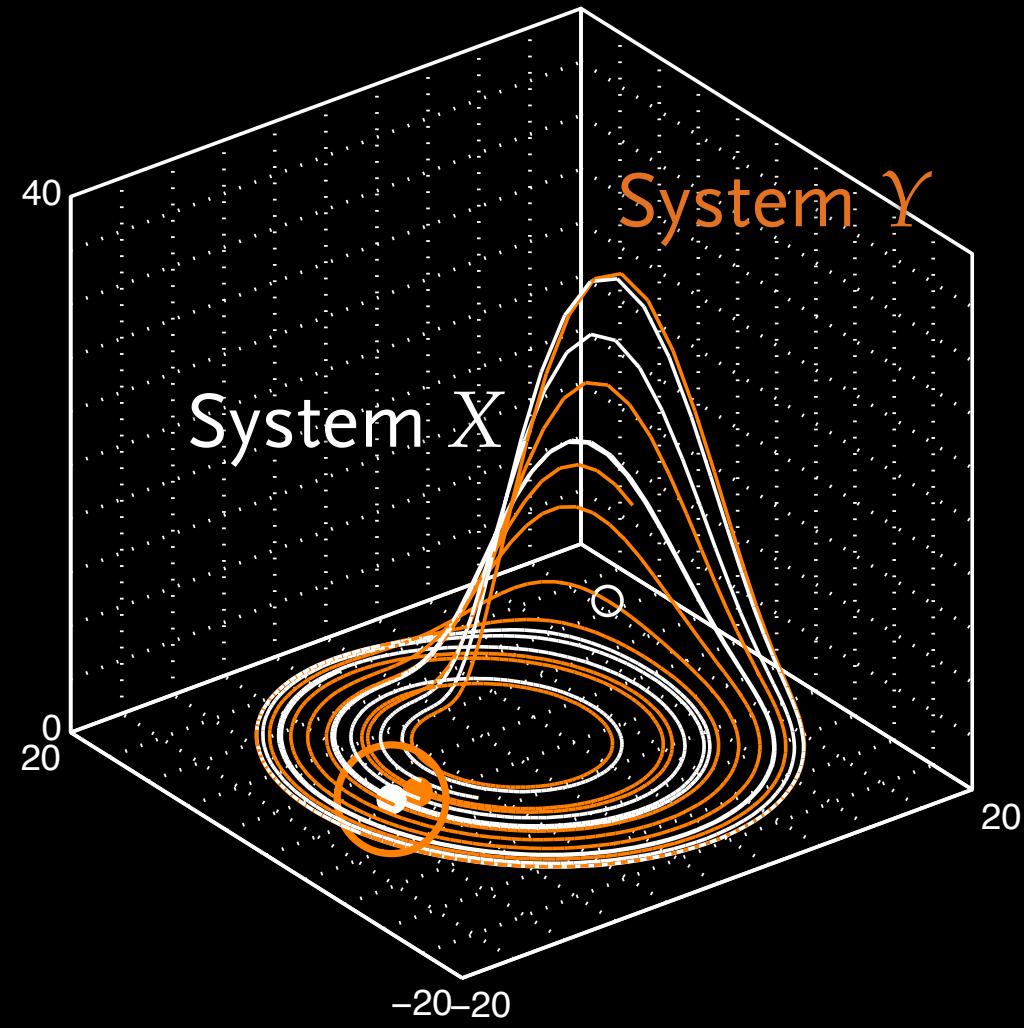
# CORRECTION OF DIAGONAL LINE LENGTHS

1. Discard border diagonals (dibo)
2. Keep only longest border diagonal (kelo)
3. Replace lengths of border dia-gonals by longest length (Censi)
4. Window masking



# COUPLING ANALYSIS

# CROSS RECURRENCE PLOT

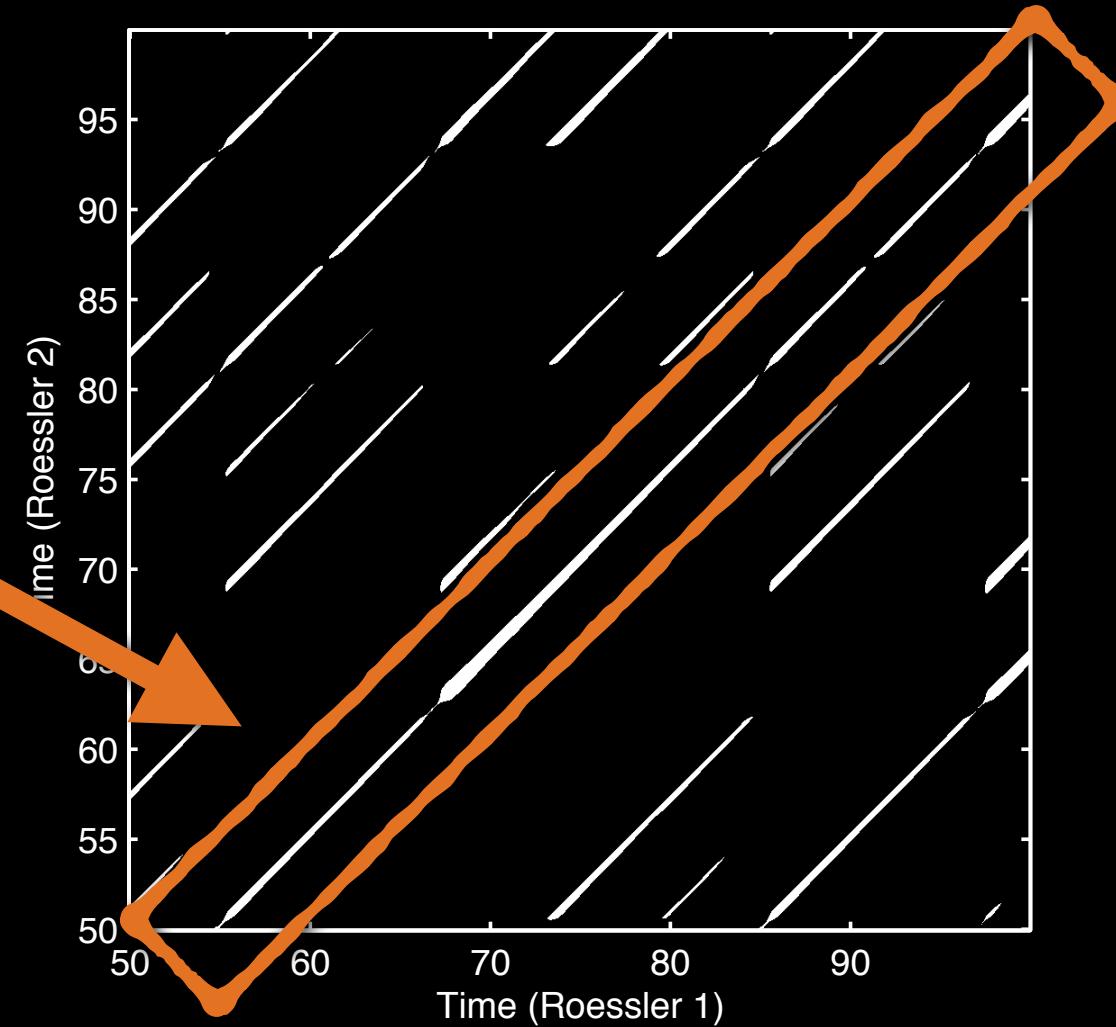
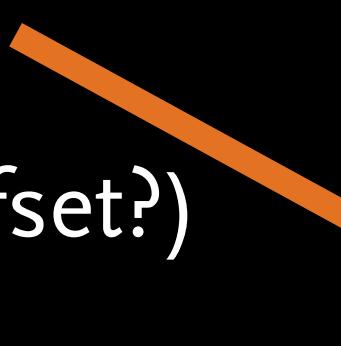


$$\text{CR}_{i,j}^{\vec{x},\vec{y}} = \Theta(\varepsilon - ||\vec{x}_i - \vec{y}_j||), \quad i = 1, \dots, N, j = 1, \dots, M$$

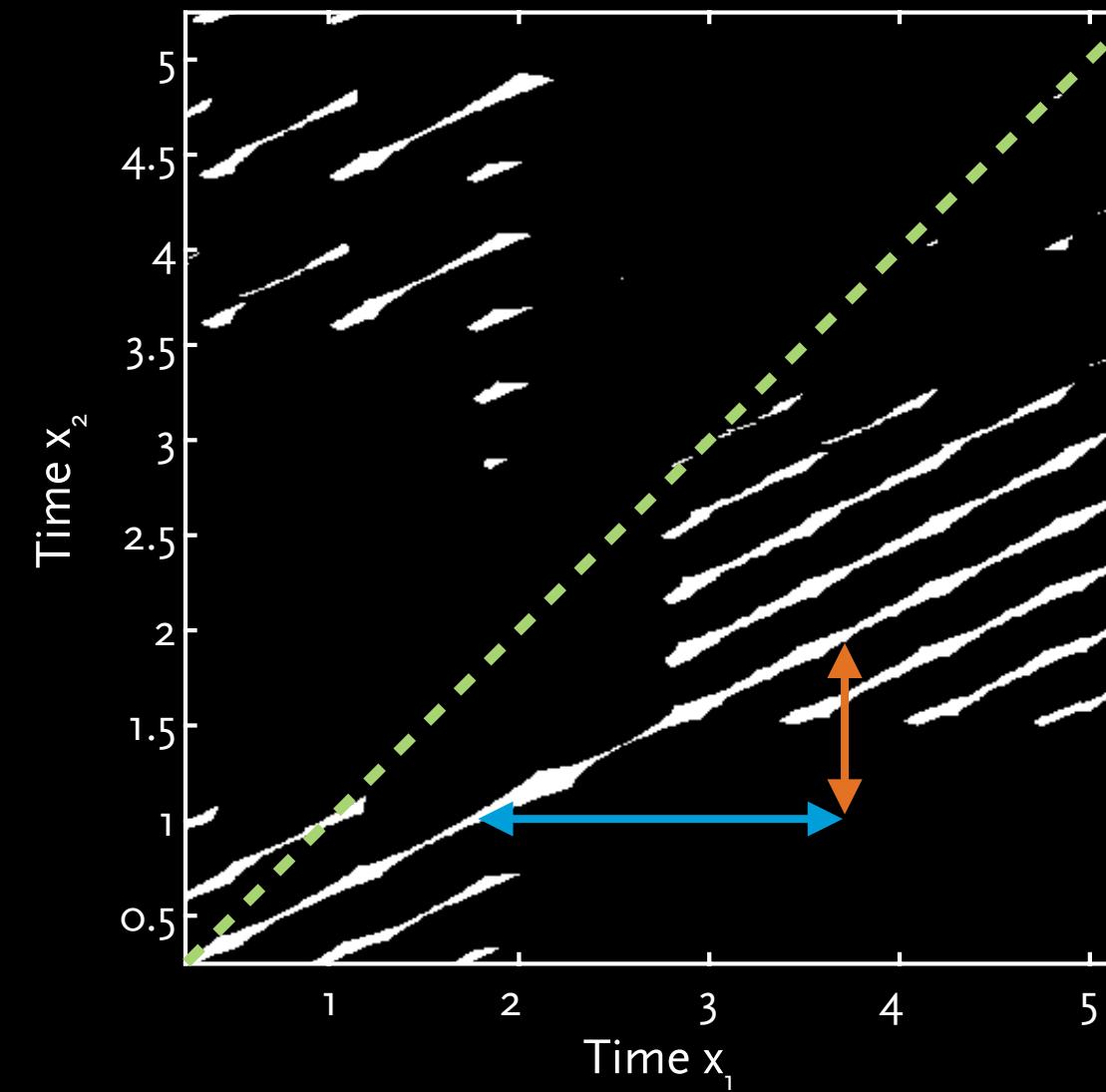
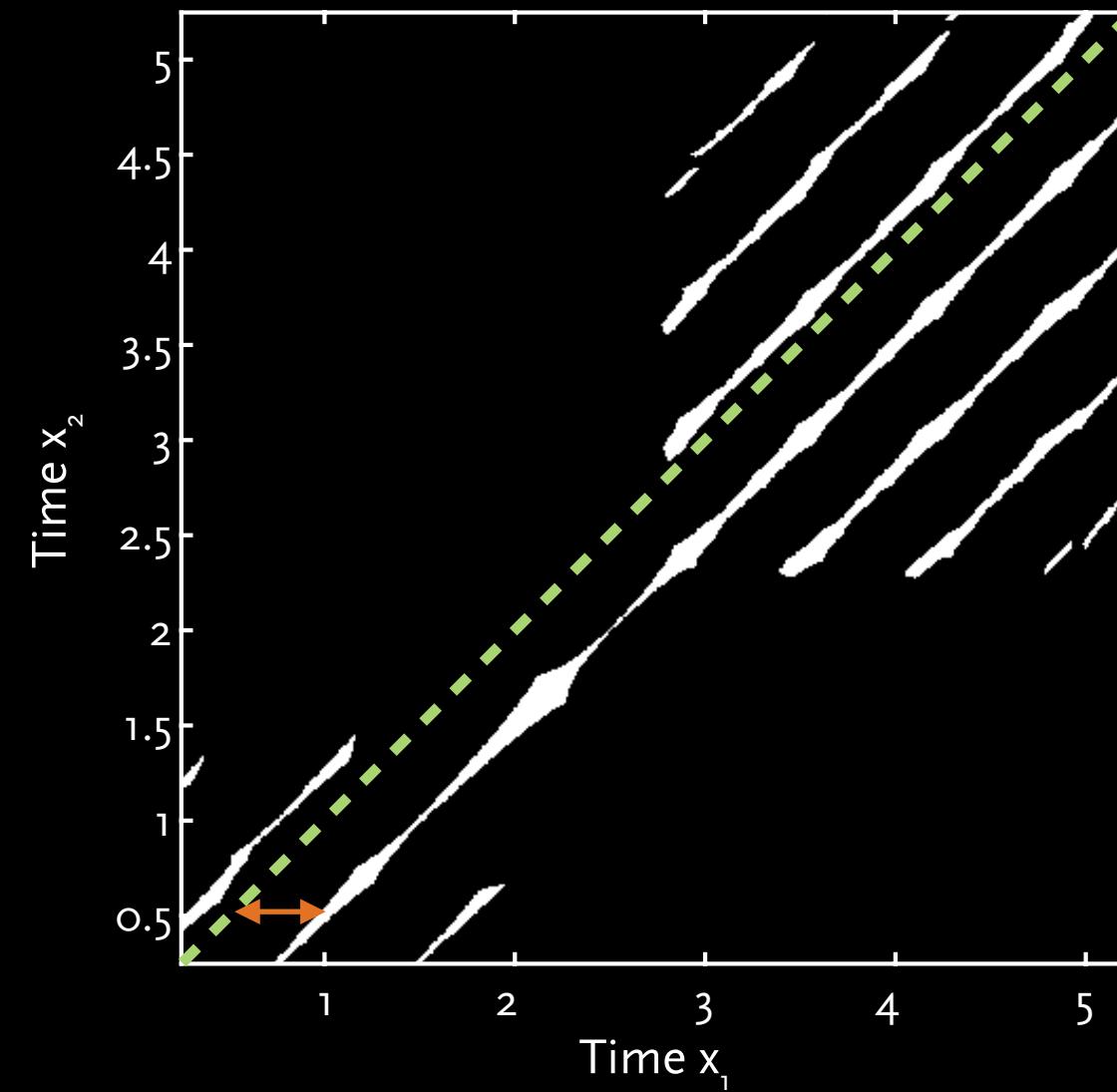
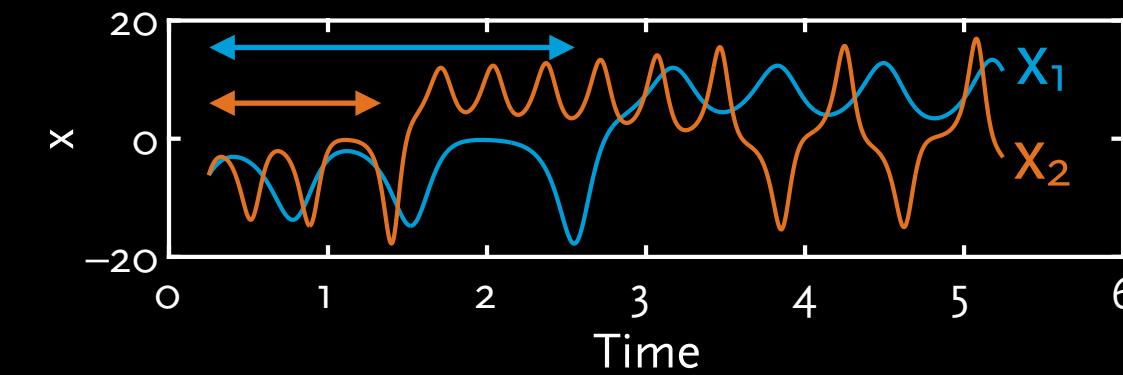
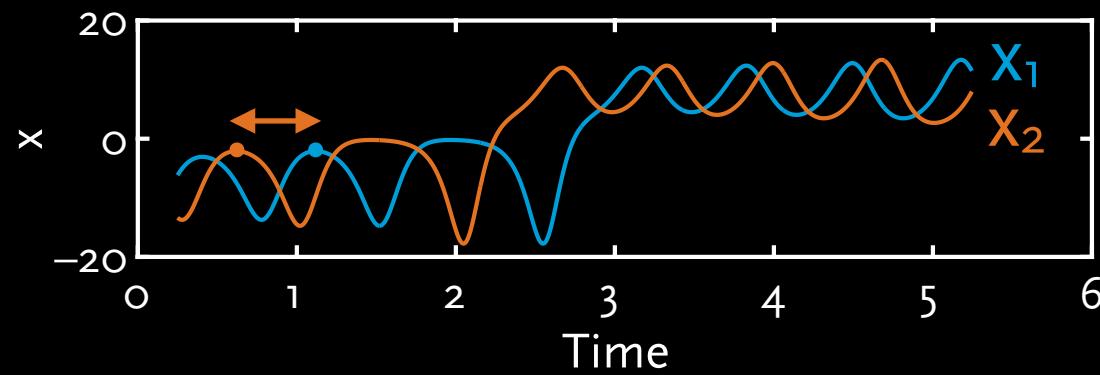
Q u i z Z

# Quiz

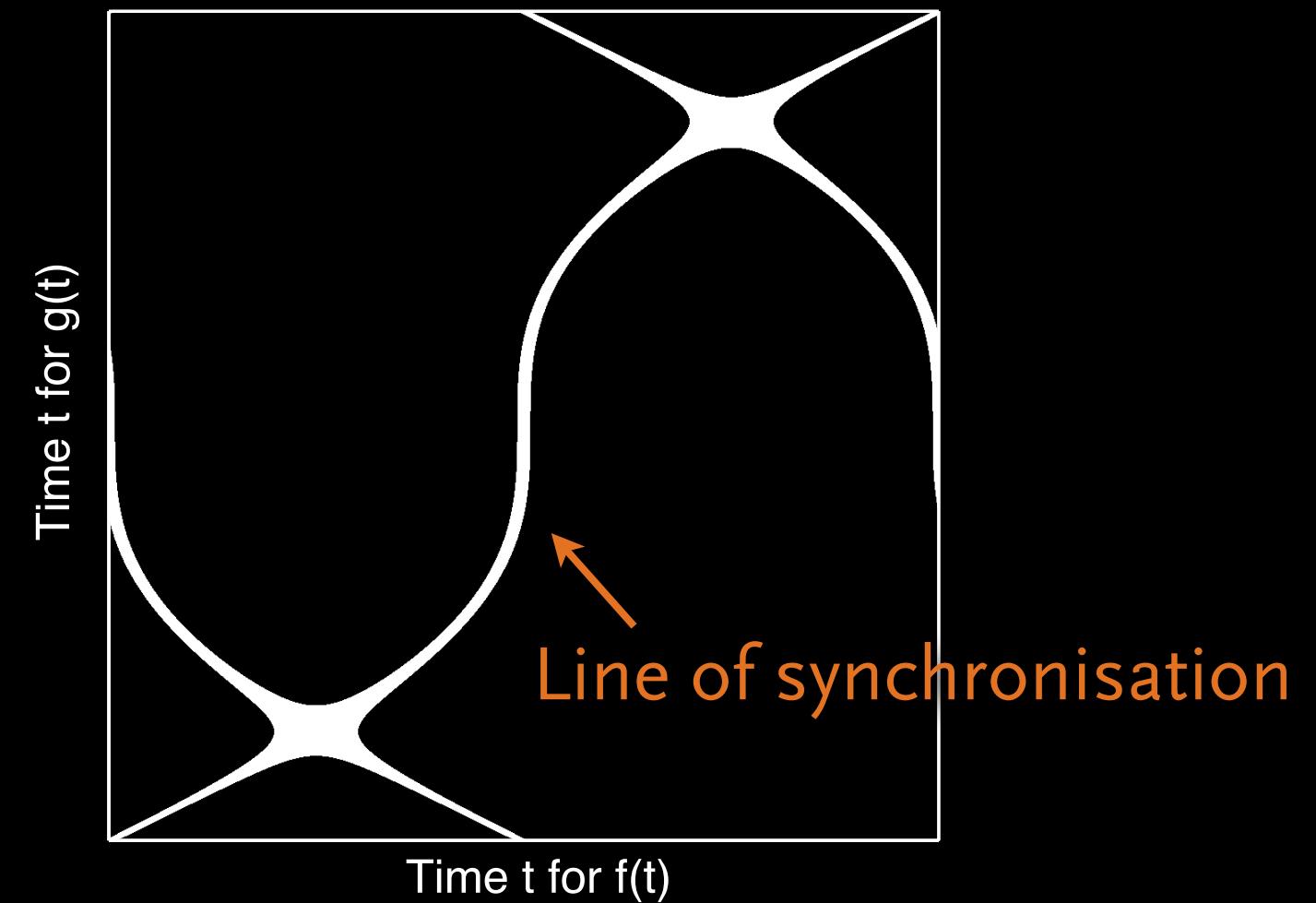
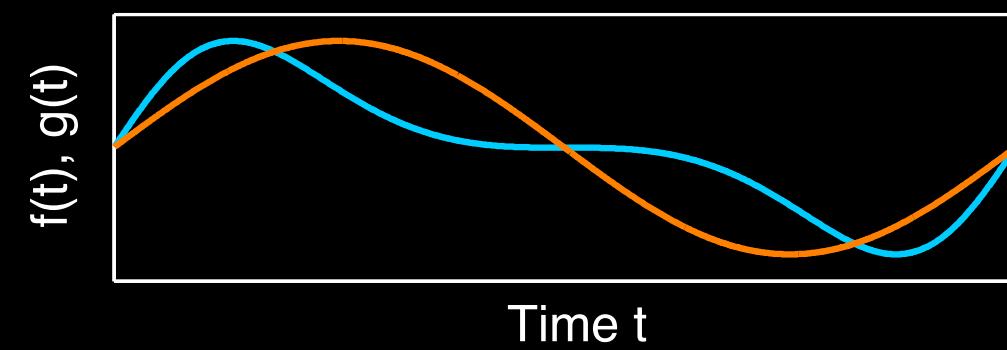
- What does this line mean/ how to interpret its shape?
  - main diagonal? (offset?)
  - straight line?
  - $45^\circ$ ?



# CROSS RECURRENCE PLOT



# CROSS RECURRENCE PLOT

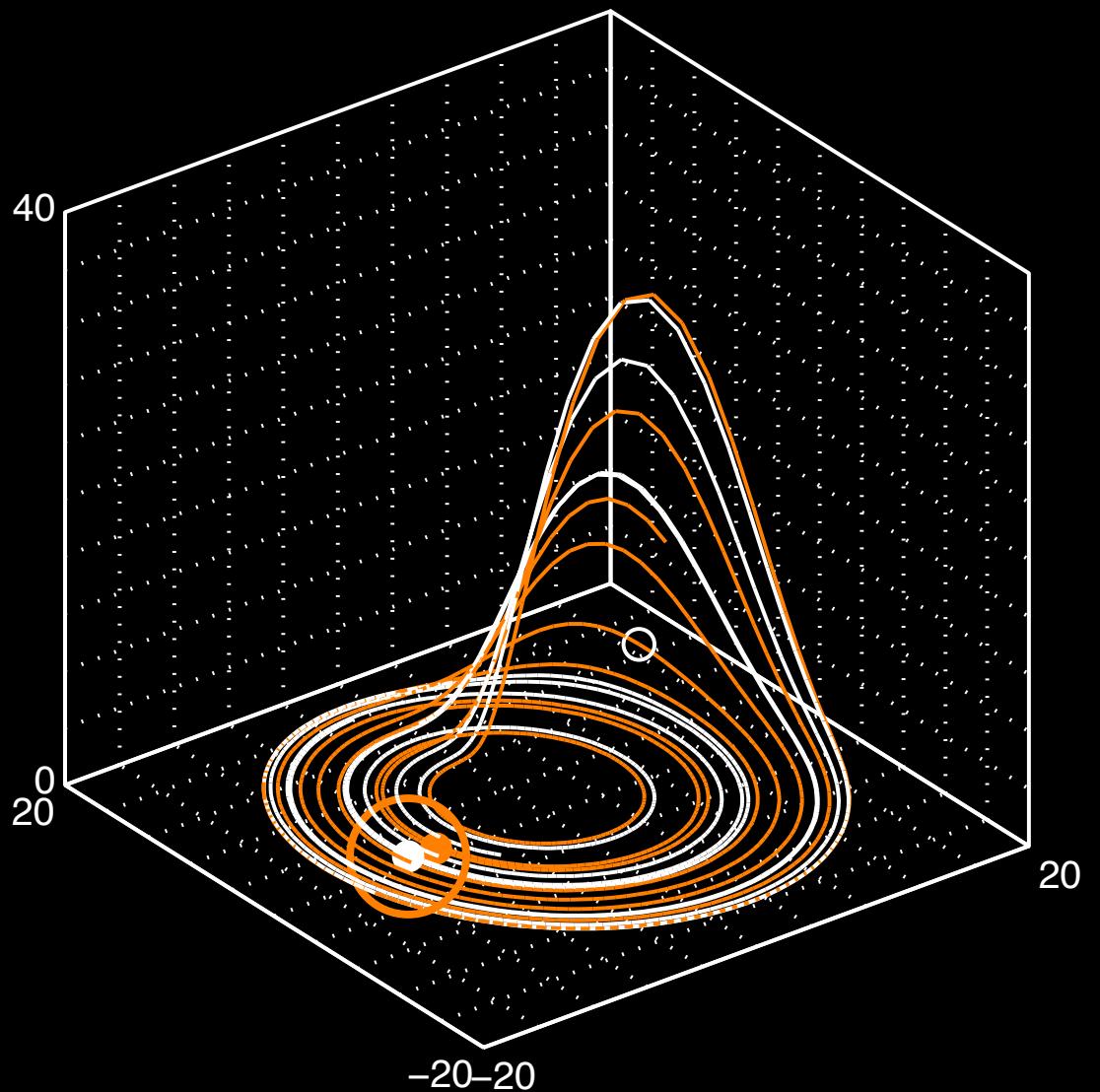


$$\text{CR}_{i,j}^{\vec{x},\vec{y}} = \Theta(\varepsilon - ||\vec{x}_i - \vec{y}_j||), \quad i = 1, \dots, N, j = 1, \dots, M$$

# CROSS RECURRENCE PLOT

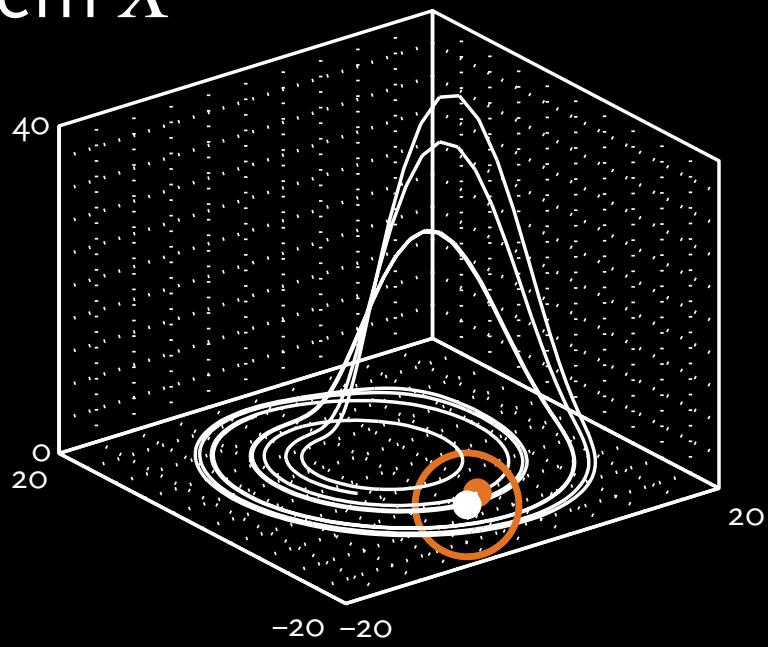
$$\text{CR}_{i,j}^{\vec{x},\vec{y}} = \Theta(\varepsilon - \|\vec{x}_i - \vec{y}_j\|), \quad i = 1, \dots, N, j = 1, \dots, M$$

- Occurrences of a similar state
- States should be comparable/  
similar; comparable dynamics
- Normalise time series

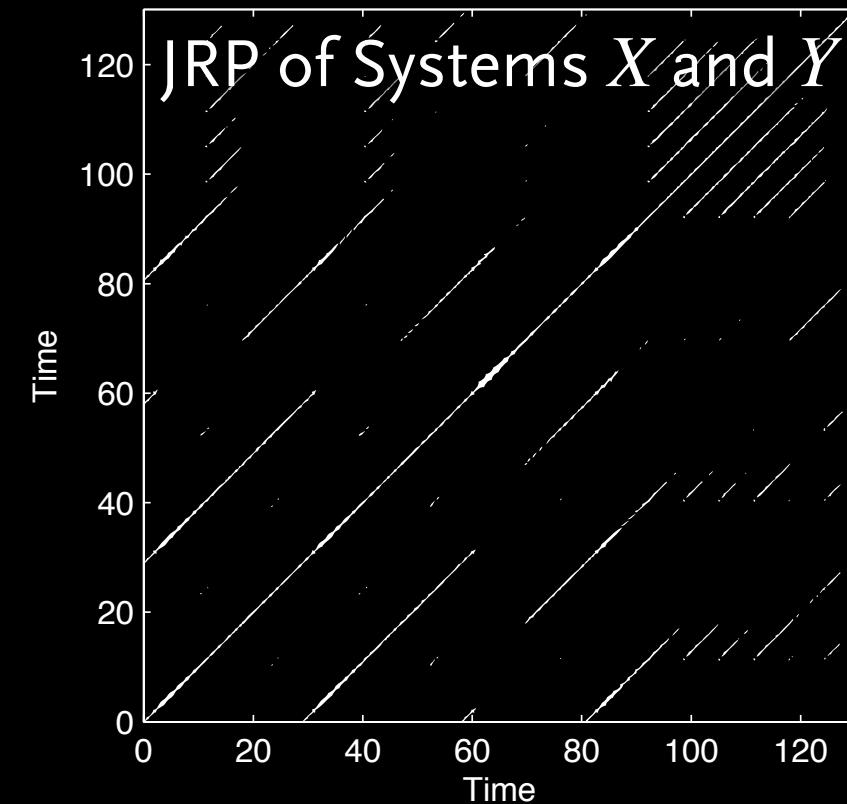
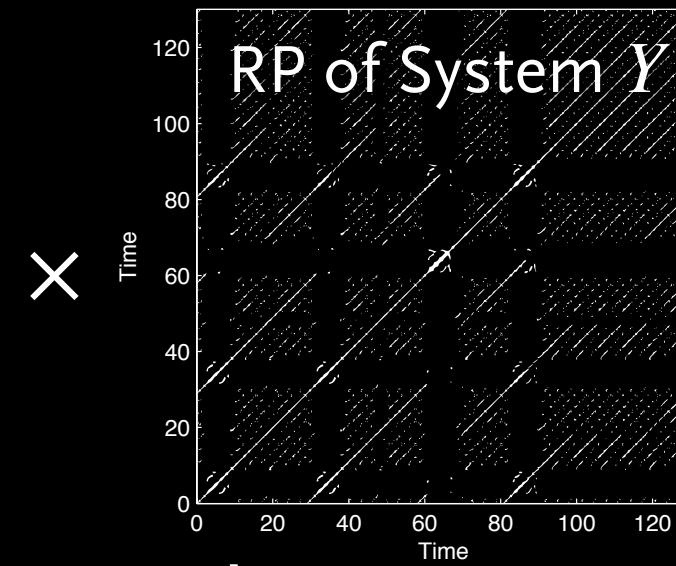
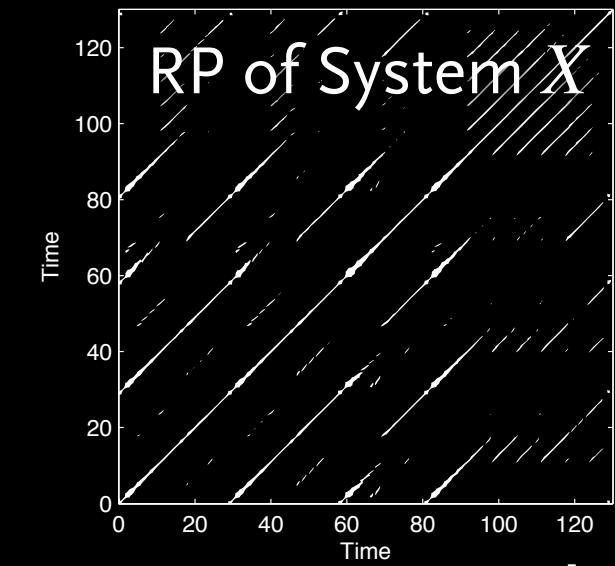
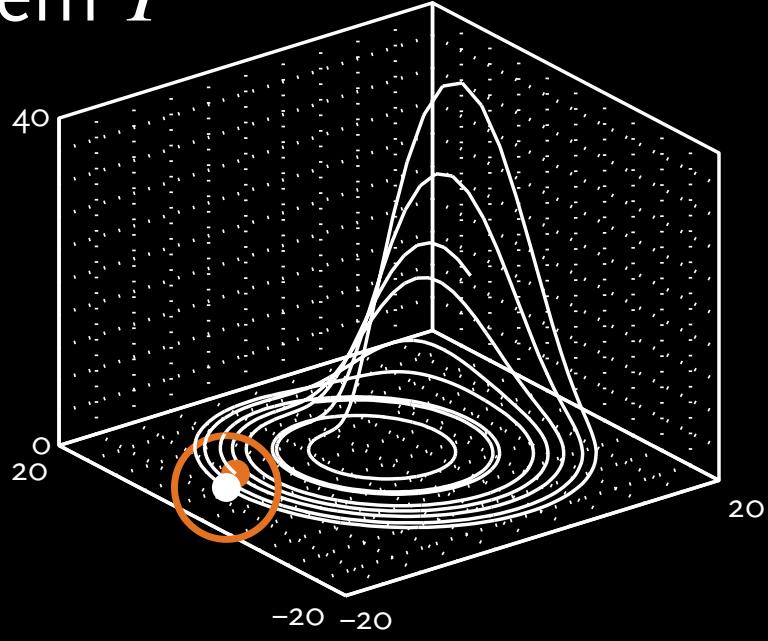


# JOINT RECURRENCE PLOT

System  $X$

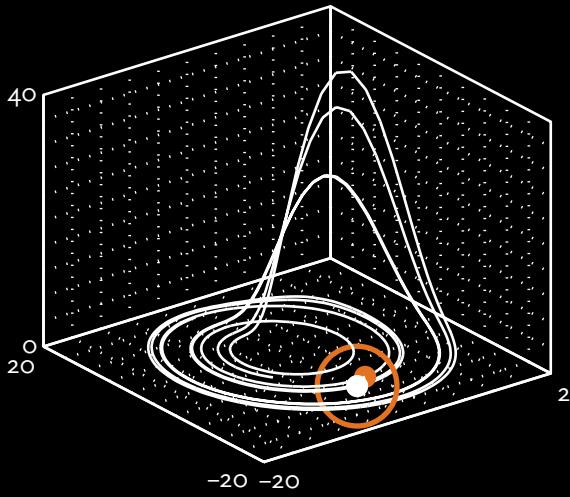


System  $Y$

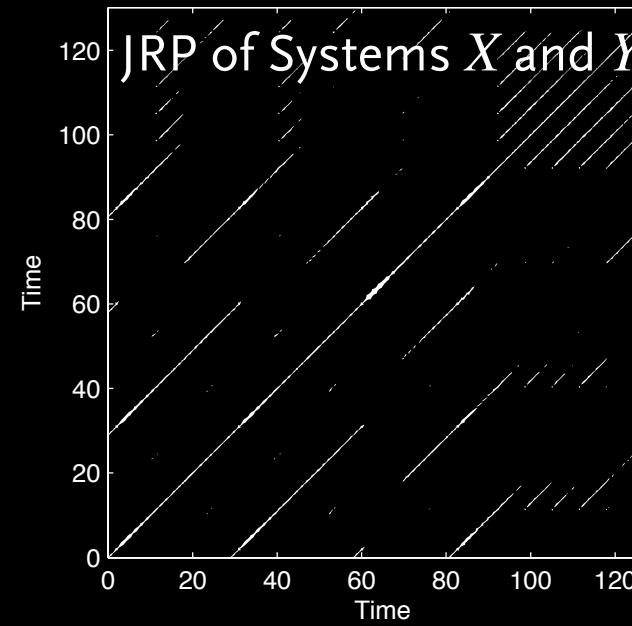
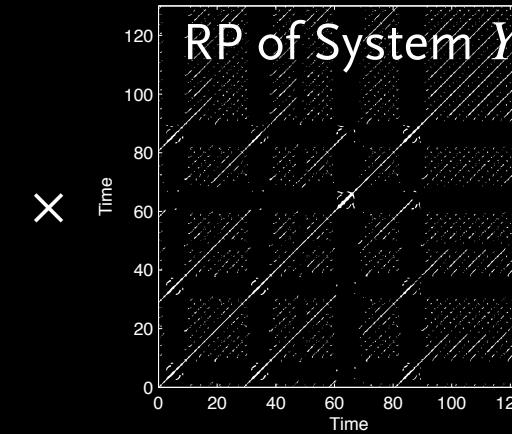
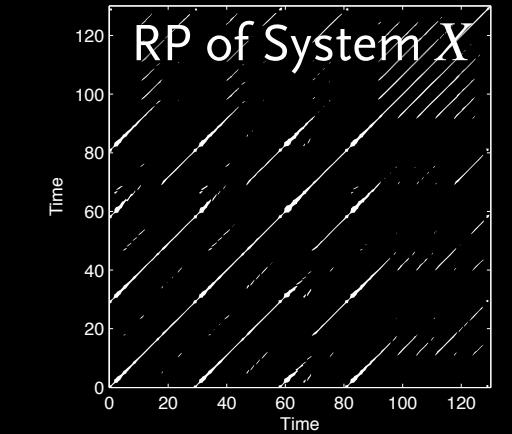
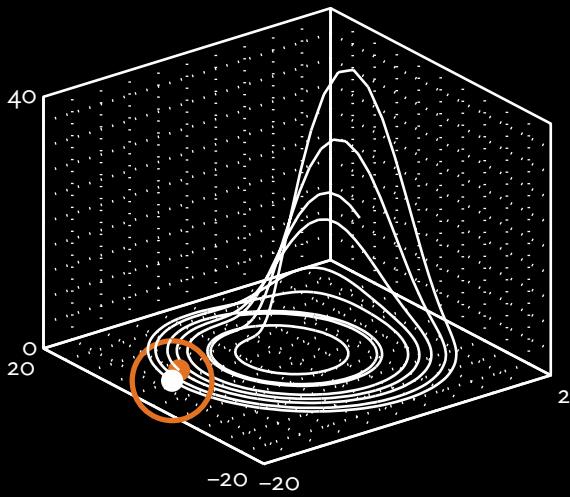


# JOINT RECURRENCE PLOT

System  $X$



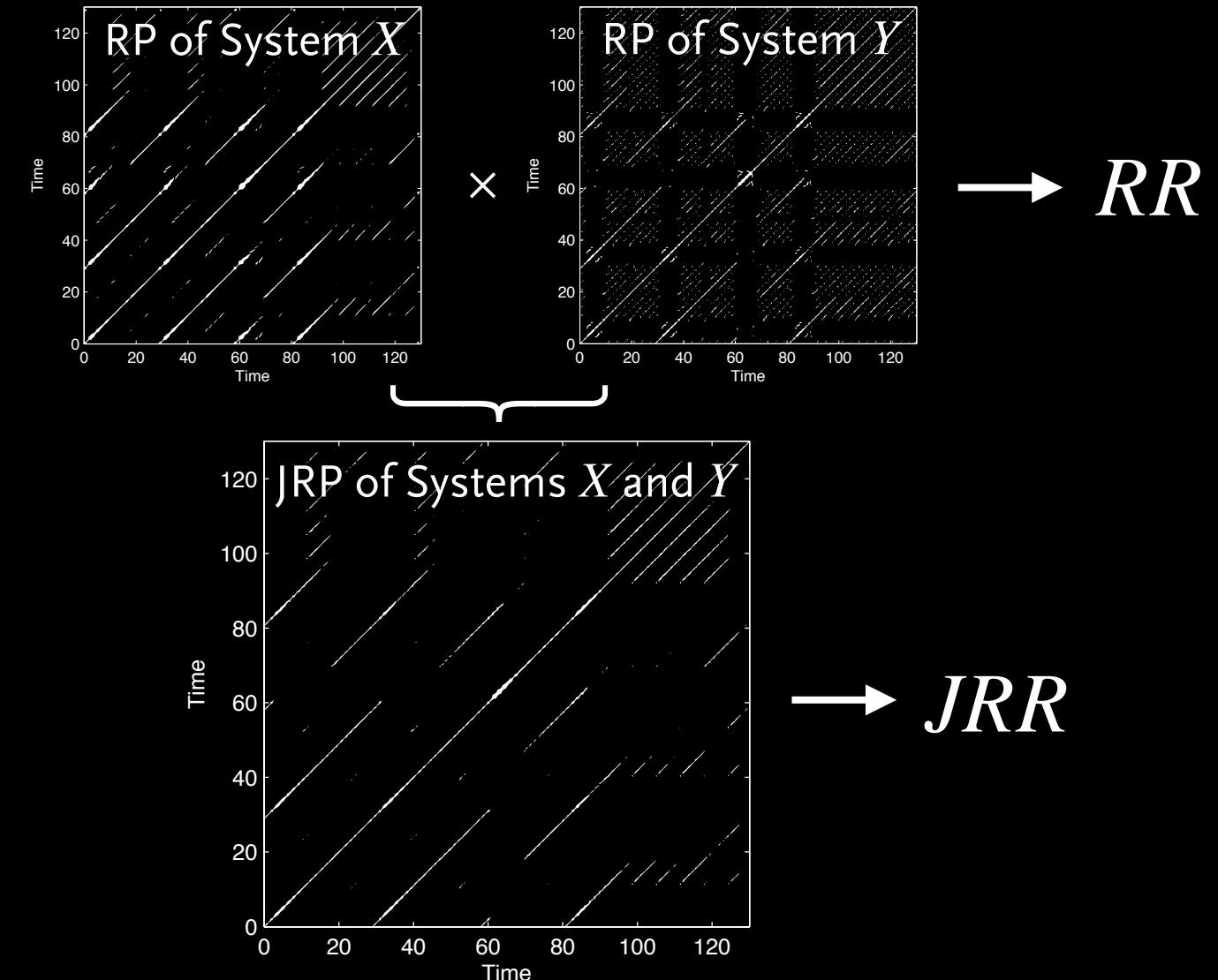
System  $Y$



$$\text{JR}_{i,j}^{\vec{x}_{(1,\dots,n)}} = \prod_{k=1}^n \mathbf{R}_{i,j}^{\vec{x}_{(k)}}, \quad i, j = 1, \dots, N$$

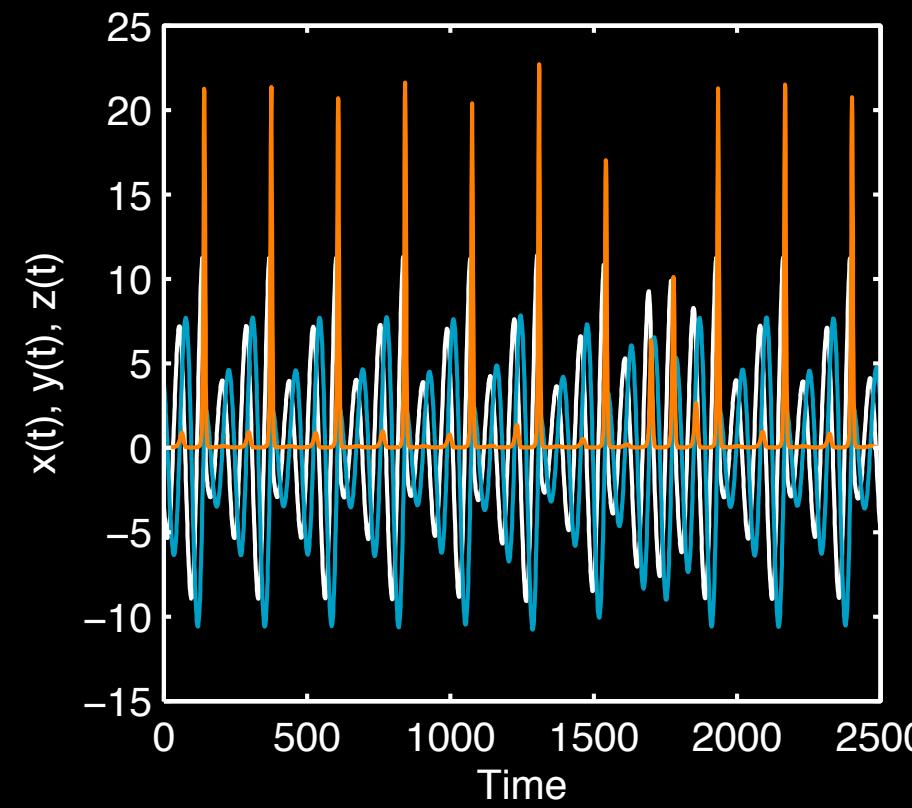
# JRP BASED SYNCHRONISATION MEASURE

- Similar recurrence pattern: generalised synchronisation (GS)
- Fix  $RR$  for system  $X$  and system  $Y$
- Synchronisation measure:  
$$S = \frac{JRR}{RR}$$
  
(0 – independent; 1 – GS)

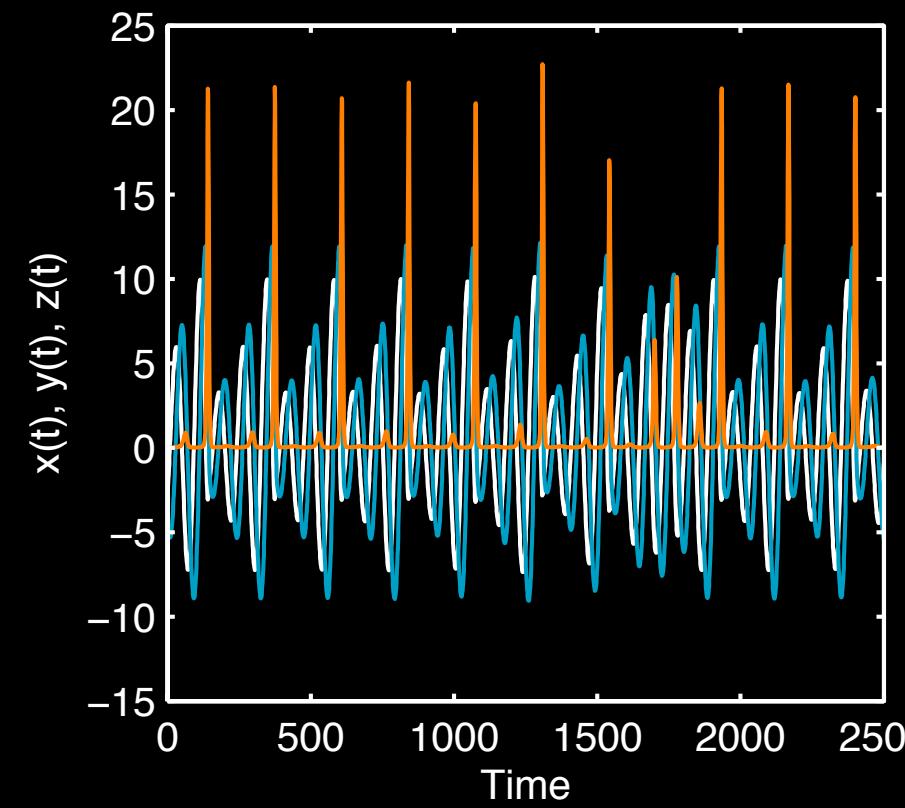


# DIFFERENCE CRP – JRP

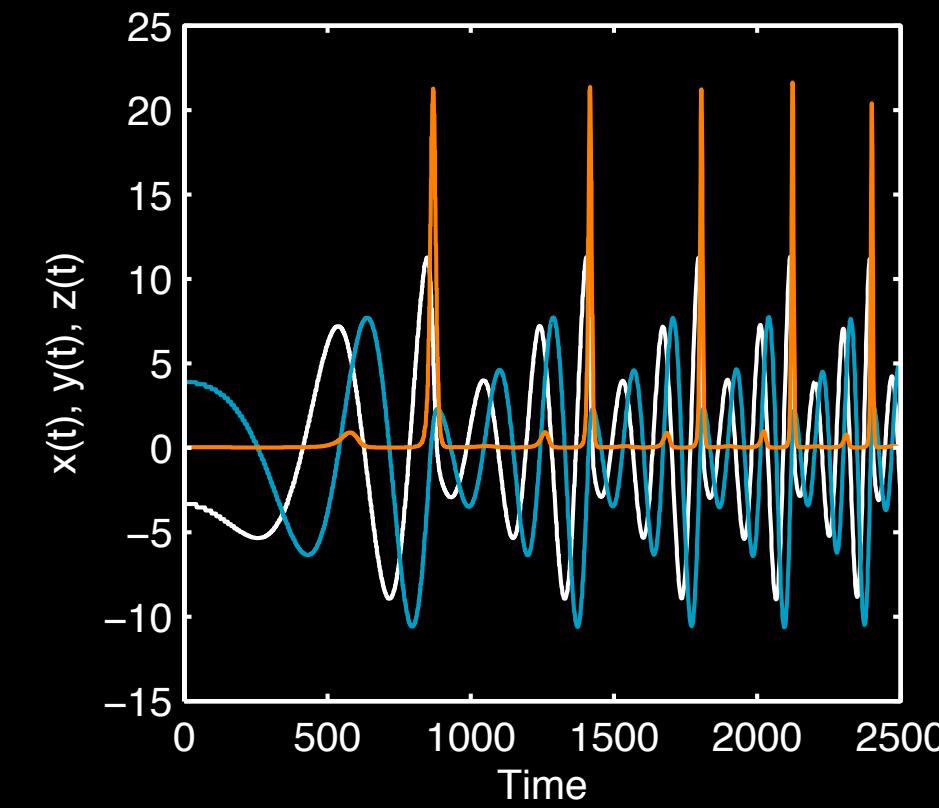
- Transformations of the Rössler system



original



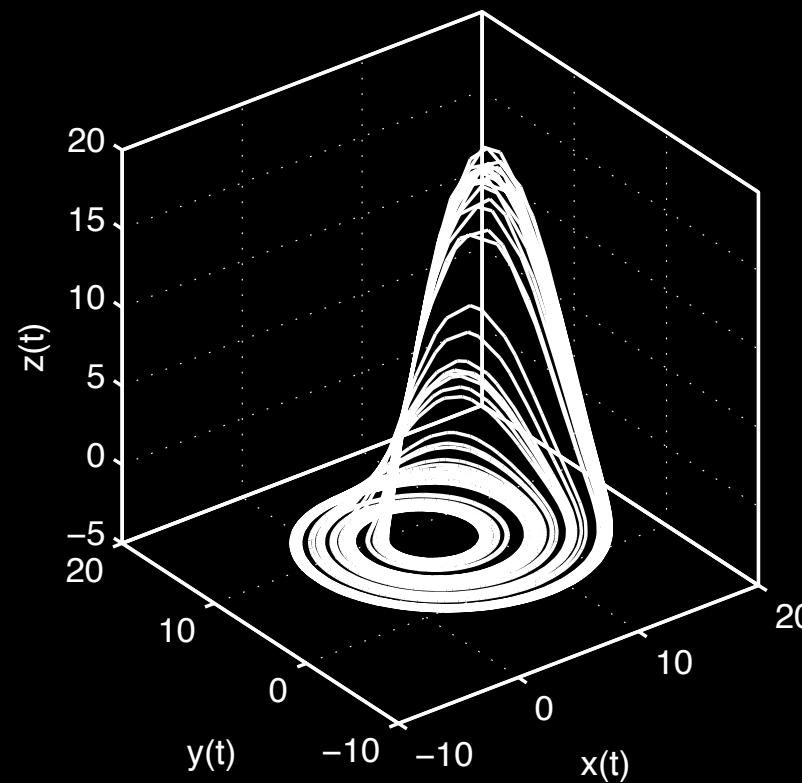
rotated



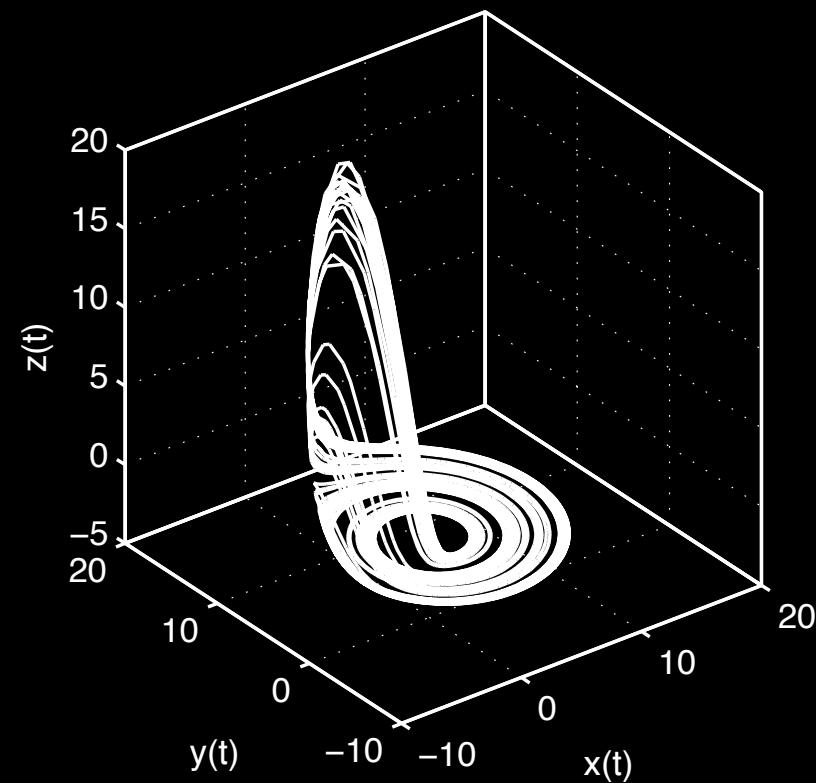
time compressed

# DIFFERENCE CRP – JRP

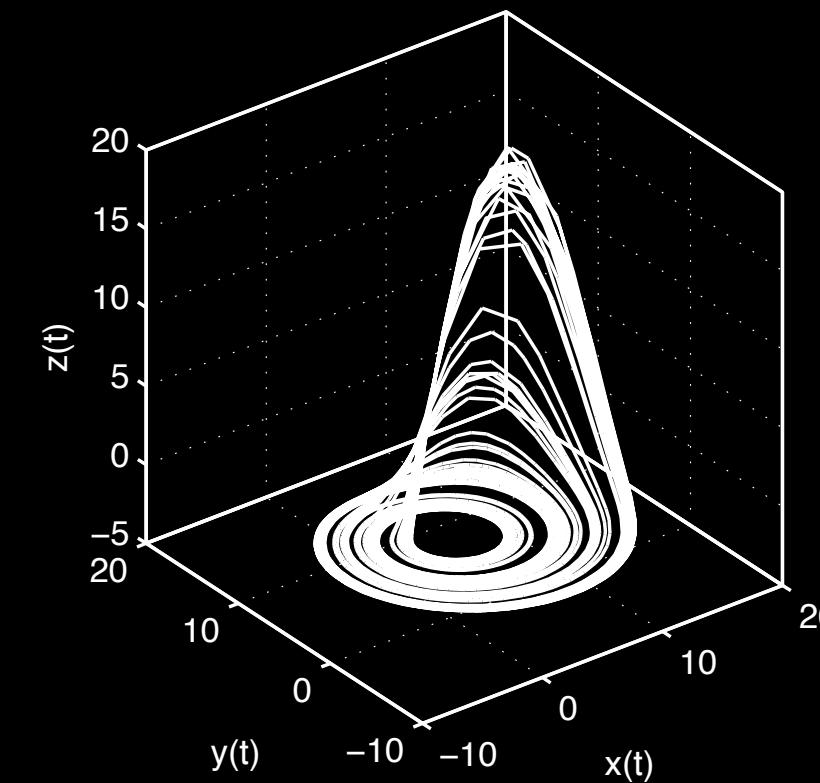
- Transformations of the Rössler system



original



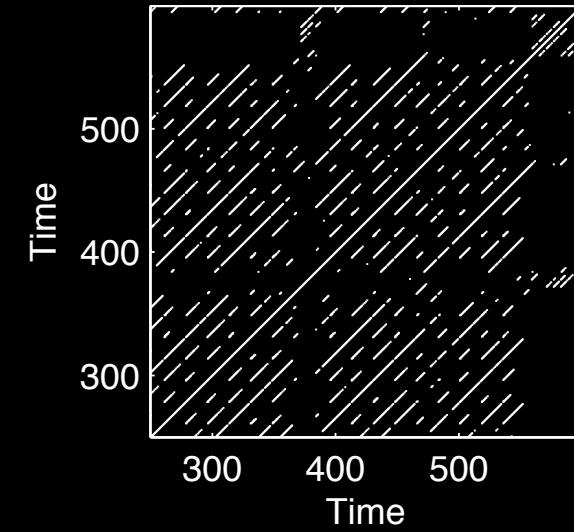
rotated



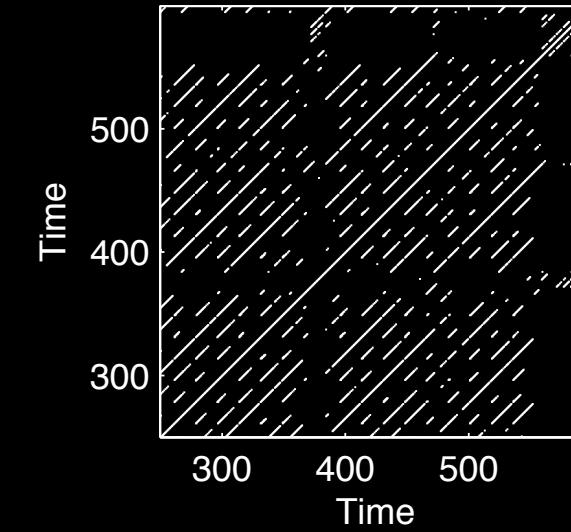
time compressed

# DIFFERENCE CRP – JRP

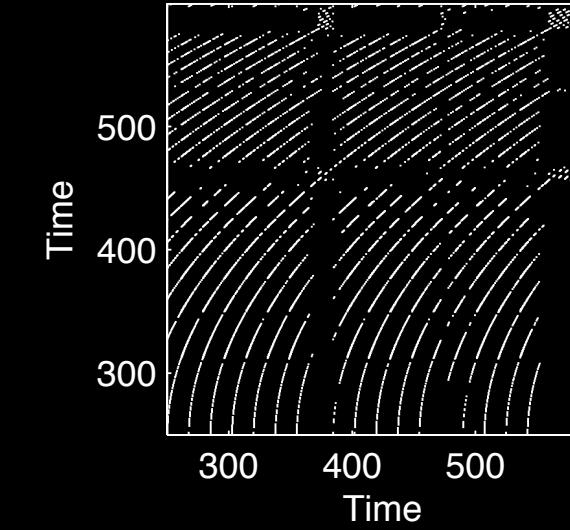
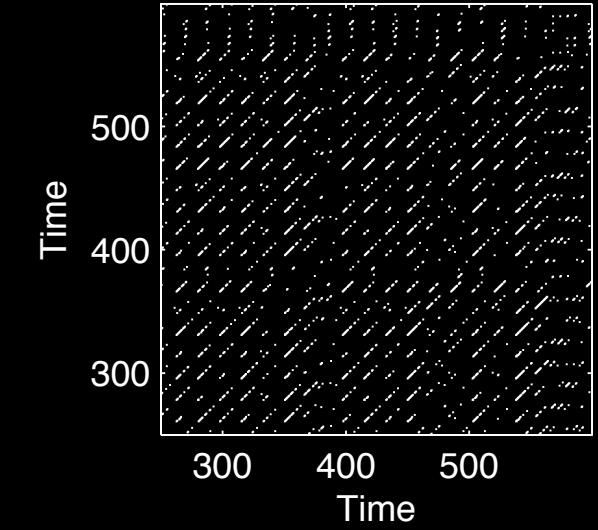
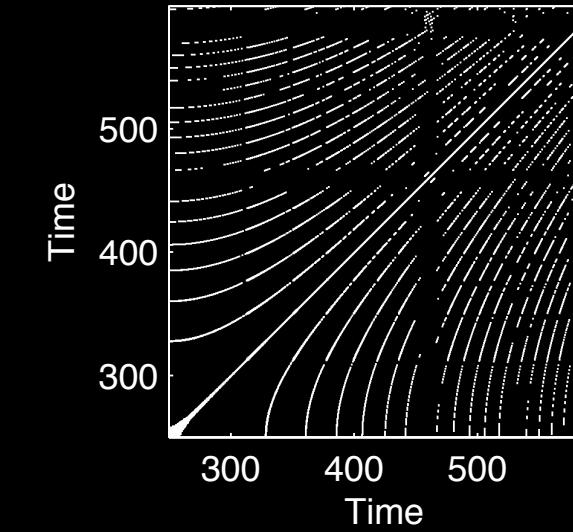
RP original



# RP rotated



# RP time compressed



A scatter plot with 'Time' on both the x-axis and y-axis, ranging from 300 to 550. A diagonal line represents the identity function. Data points are scattered around this line.

CRP

JRP

CRF

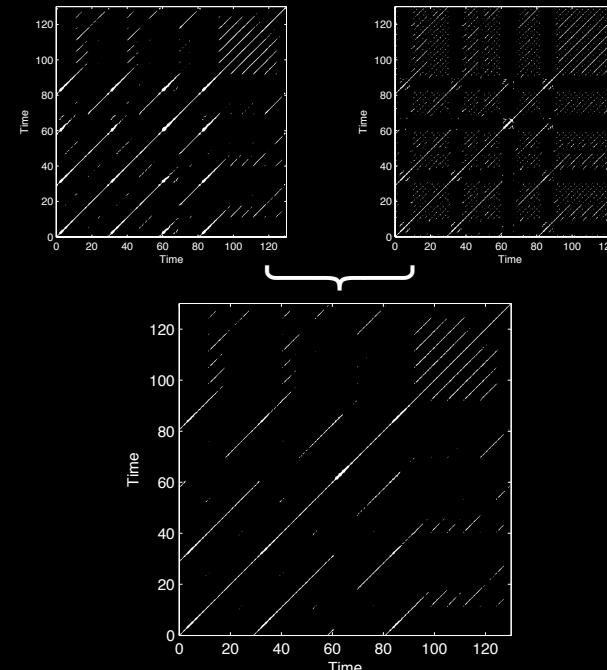
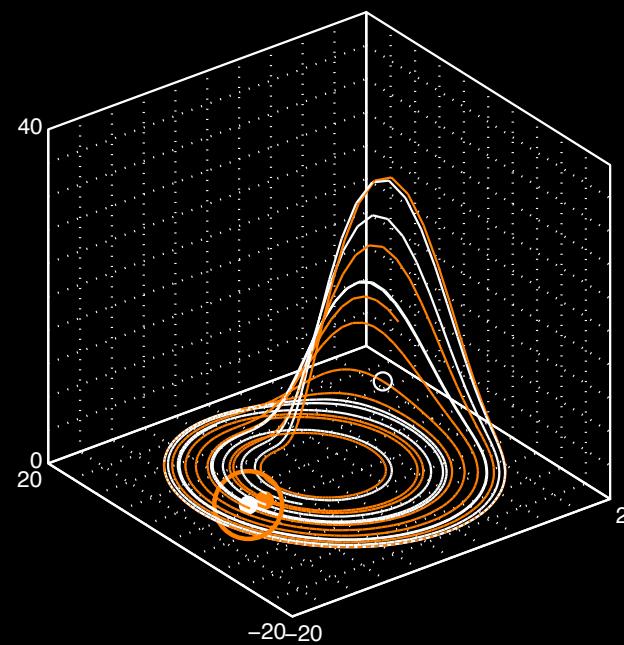
JRP

A horizontal orange arrow pointing to the right, spanning most of the page width.

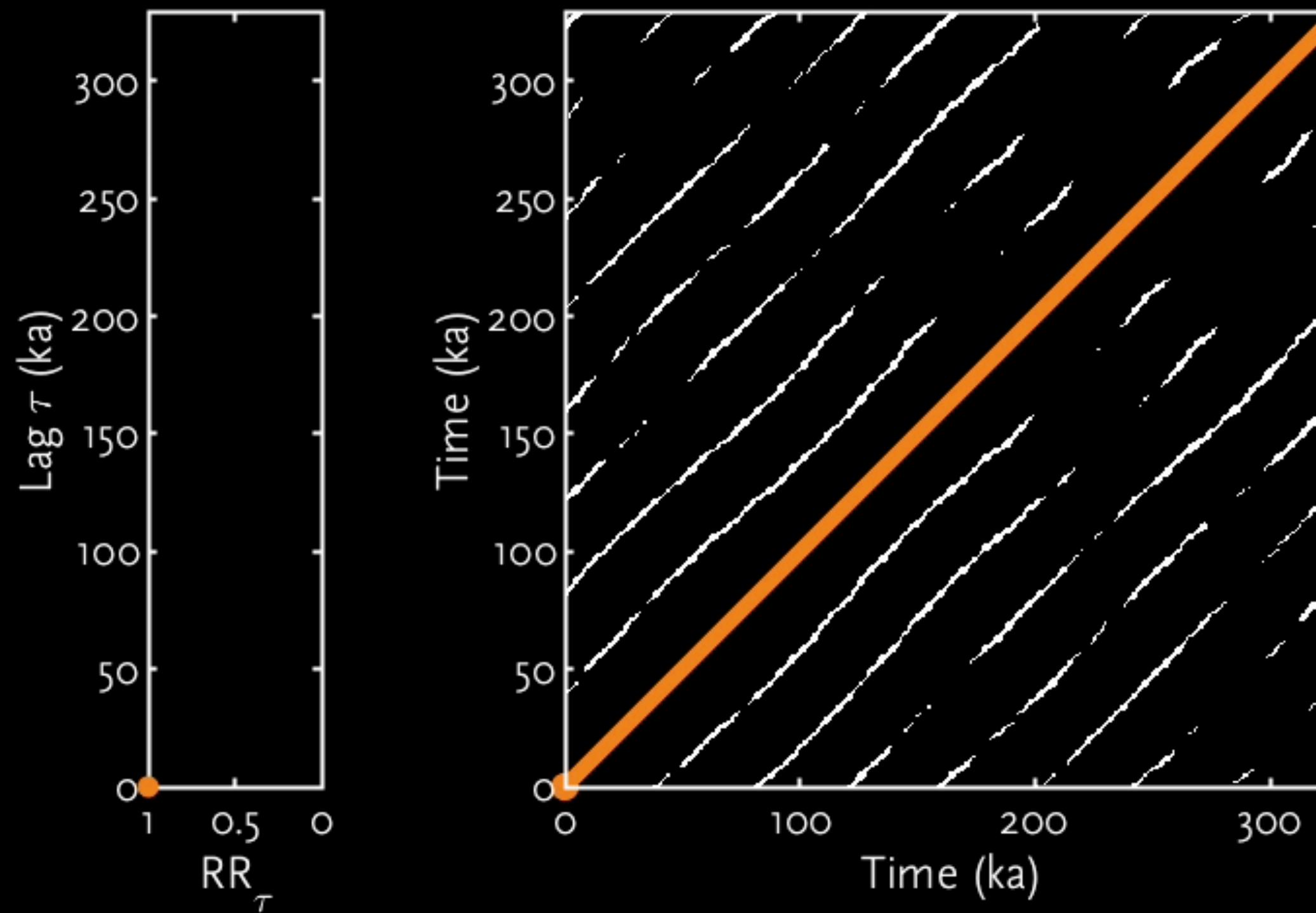
A horizontal orange double-headed arrow pointing to the right, indicating the direction of the next section.

# DIFFERENCE CRP–JRP

- CRP: simultaneous occurrence of a similar state
  - ➡ Comparable/ similar states
  - ➡ Differences in timing (time alignment)
- JRP: simultaneous occurrence of a recurrence
  - ➡ Different states
  - ➡ Same timing (generalised synchronisation)

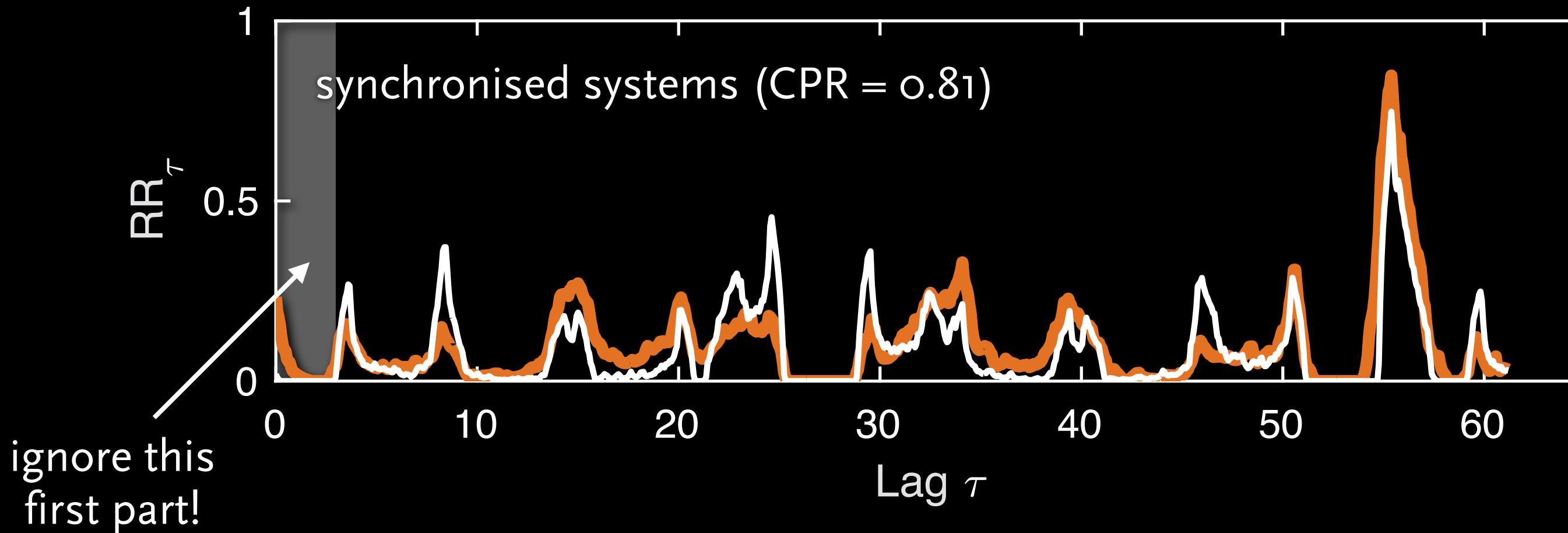


# PHASE SYNCHRONISATION USING $\tau$ -RECURRENCE RATE



# PHASE SYNCHRONISATION

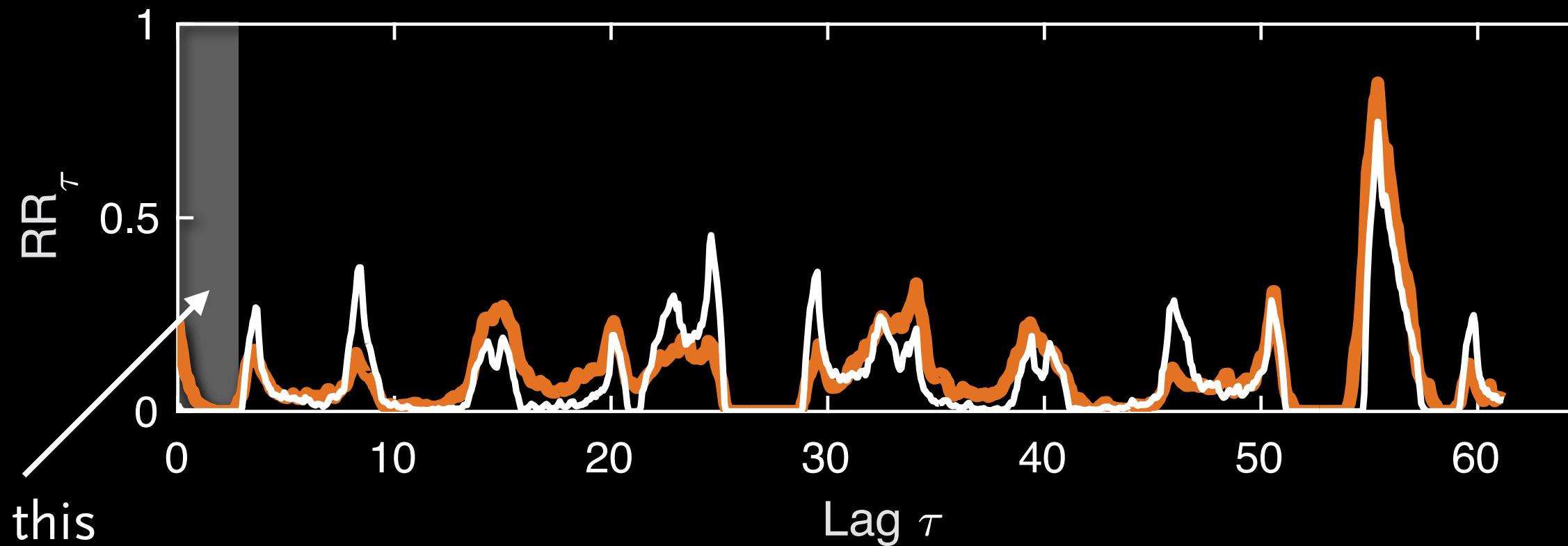
- Synchronisation index: correlation coefficient  $CPR = \langle RR_{\tau}^x \cdot RR_{\tau}^y \rangle$



Q u i z Z

# Quiz

- Why should we remove this first part from analysis?

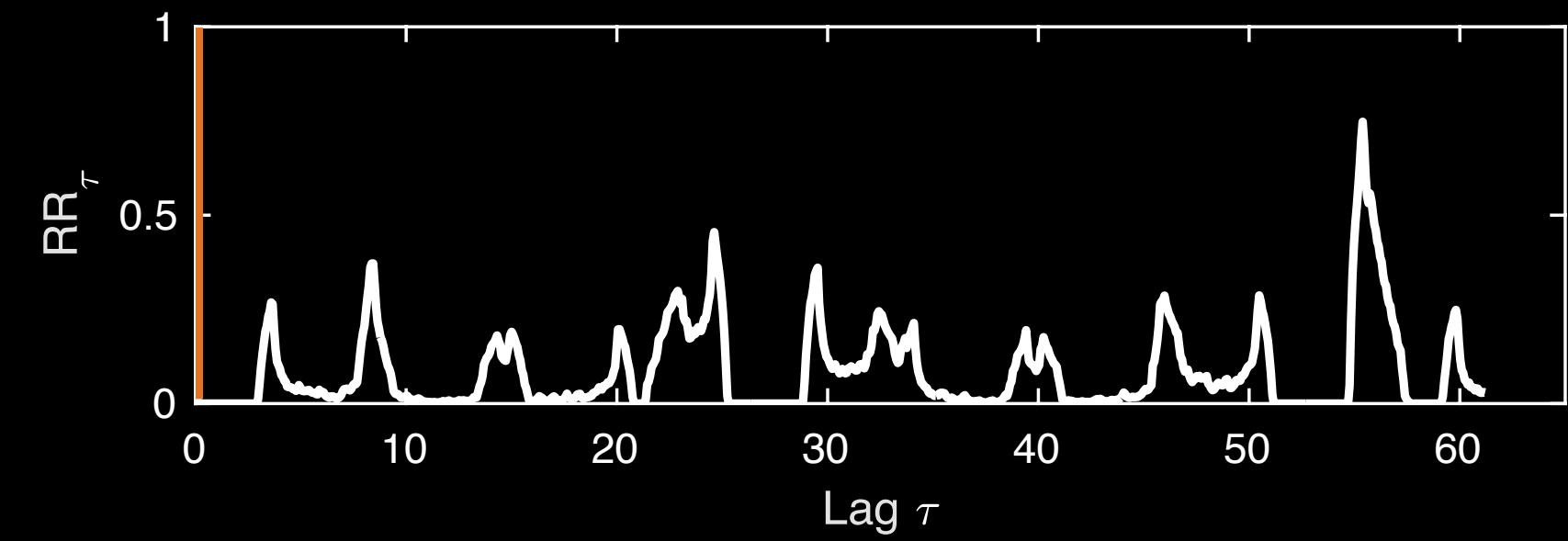
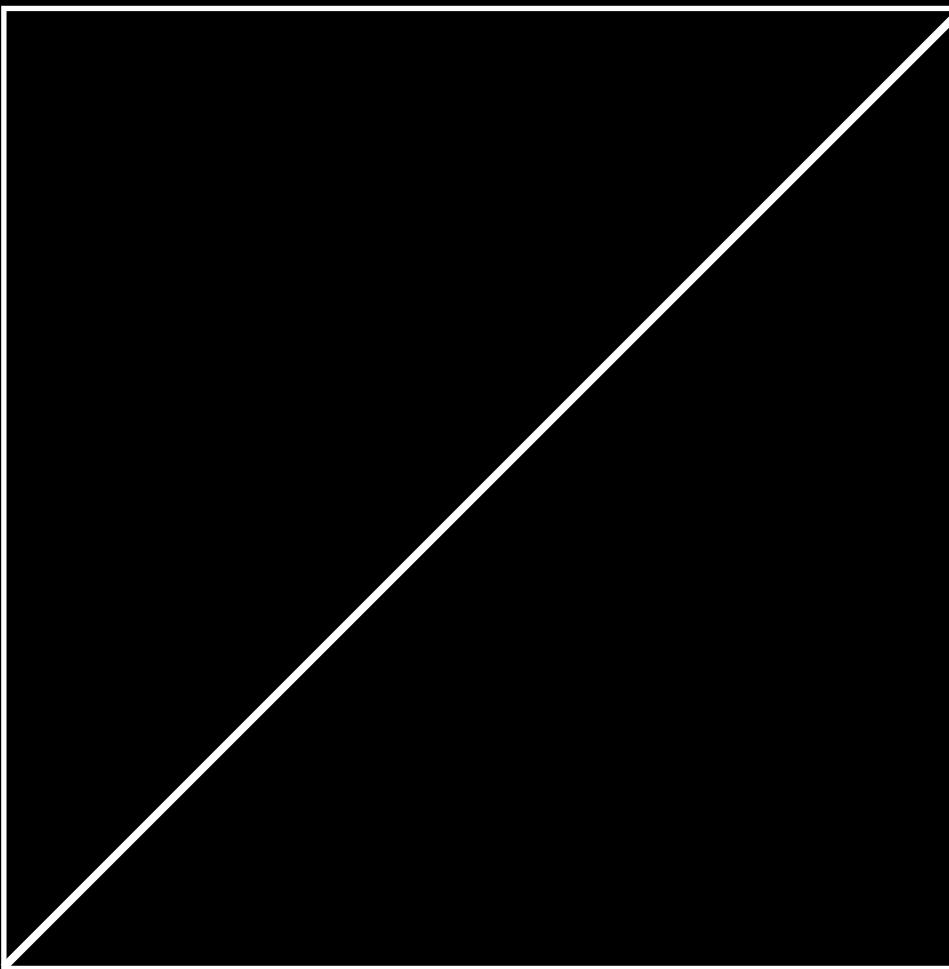


ignore this  
first part!



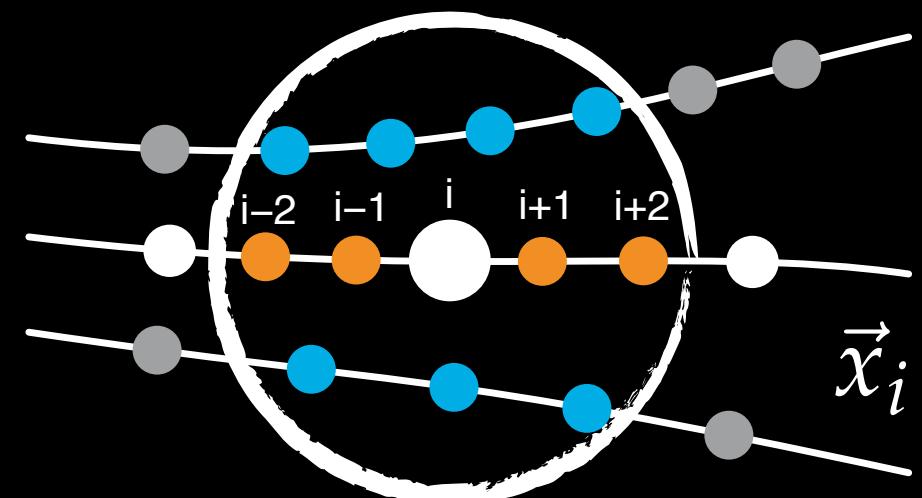
# PHASE SYNCHRONISATION

- We have the main diagonal (even when no other recurrences)!

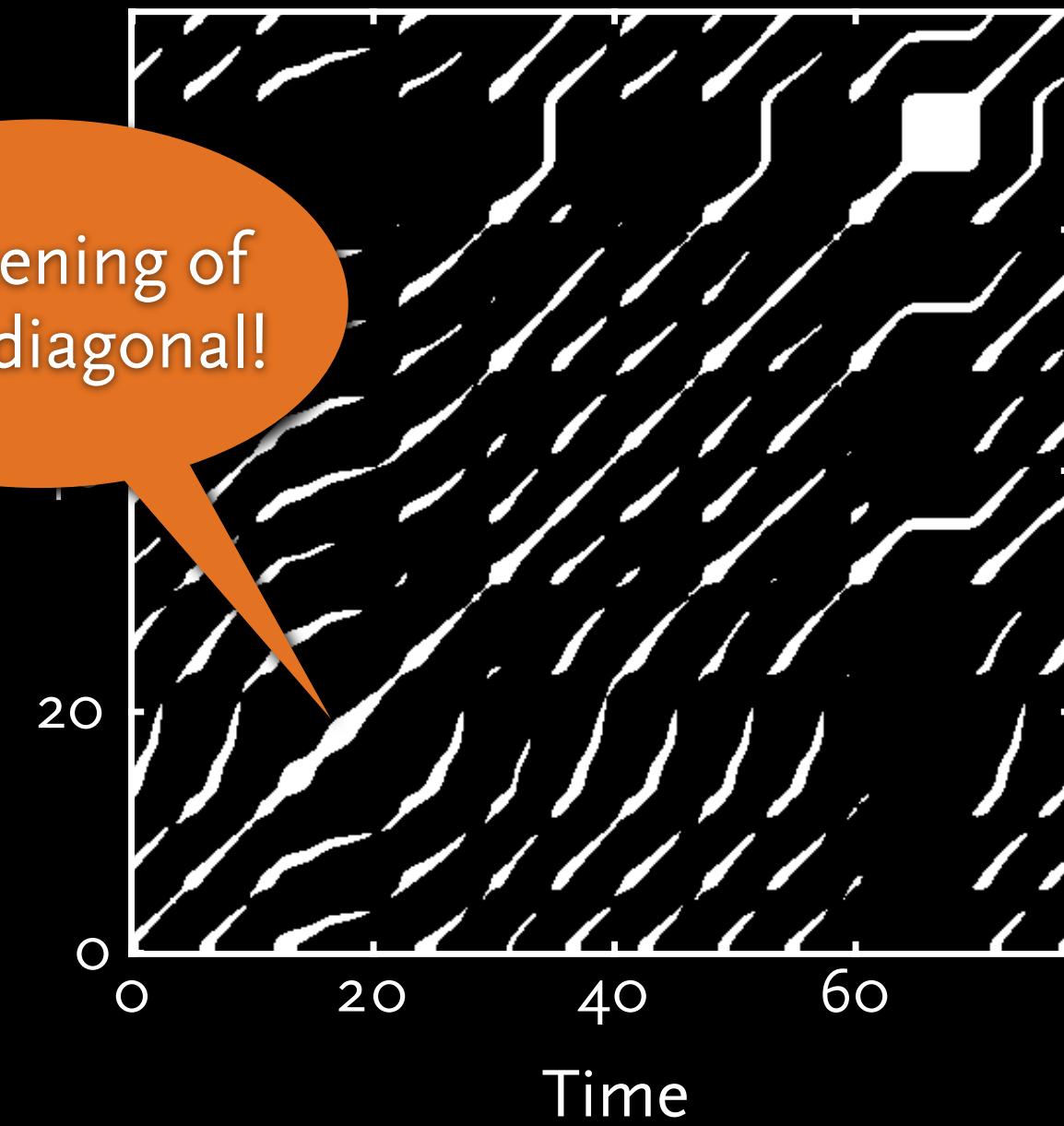




# PHASE SYNCHRONISATION



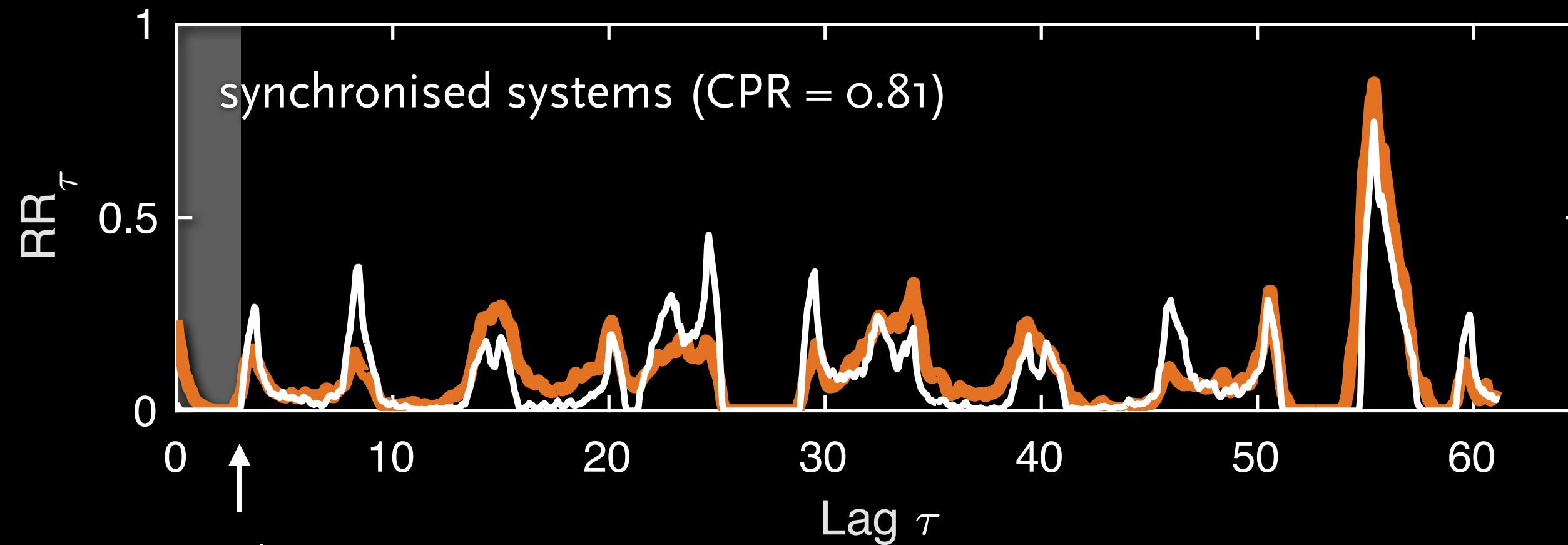
Thickening of  
main diagonal!





# PHASE SYNCHRONISATION

- Synchronisation index: correlation coefficient  $CPR = \langle RR_{\tau}^x \cdot RR_{\tau}^y \rangle$



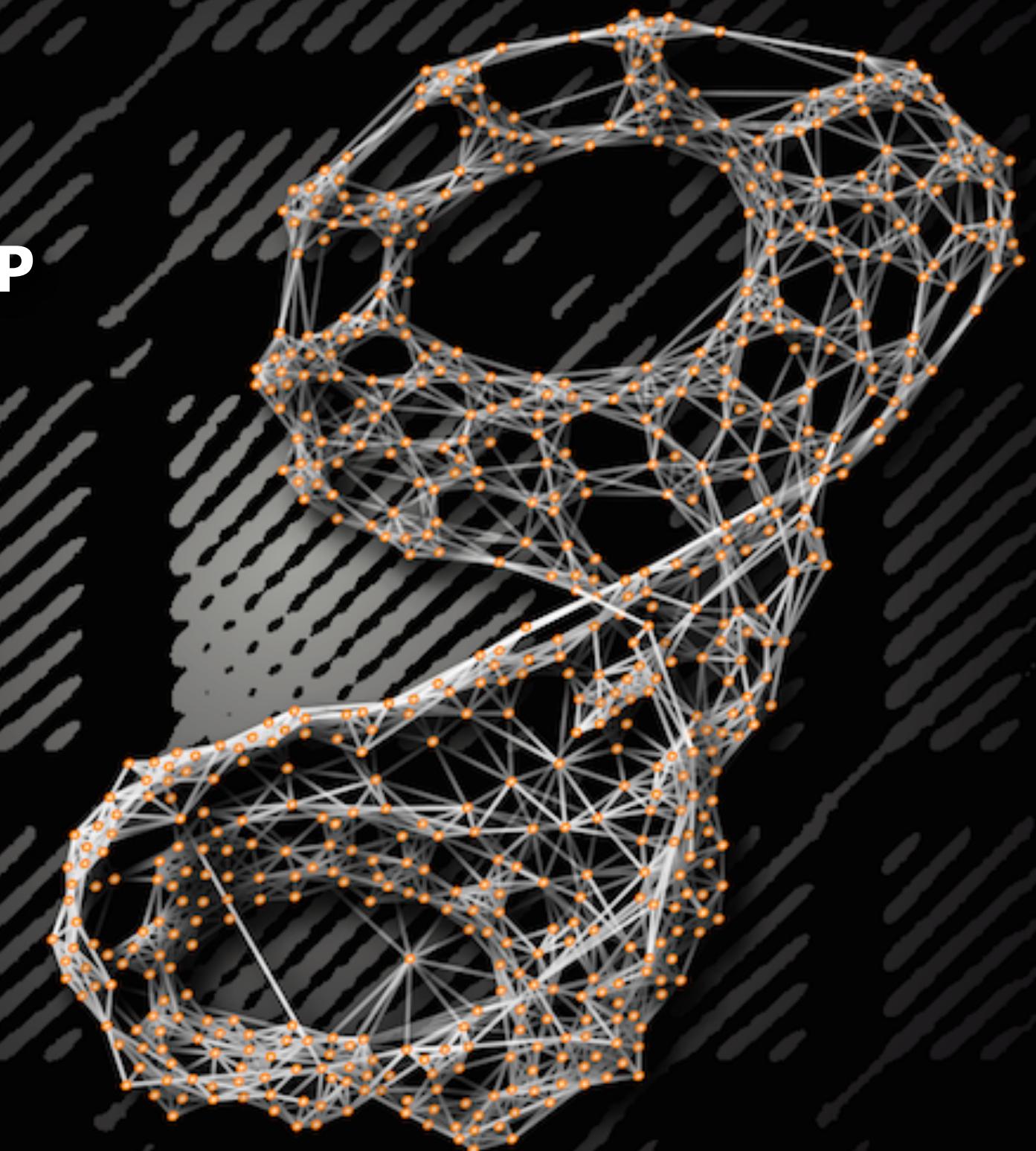
use auto-correlation time  
(where ACF falls below 1/e)



POTSDAM INSTITUTE FOR  
CLIMATE IMPACT RESEARCH

# RECURRENCE PLOTS WORKSHOP

- ① Phase space
- ② Visual interpretation
- ③ Sensitivity on Parameters
- ④ Recurrence quantification analysis
- ⑤ Coupling analysis

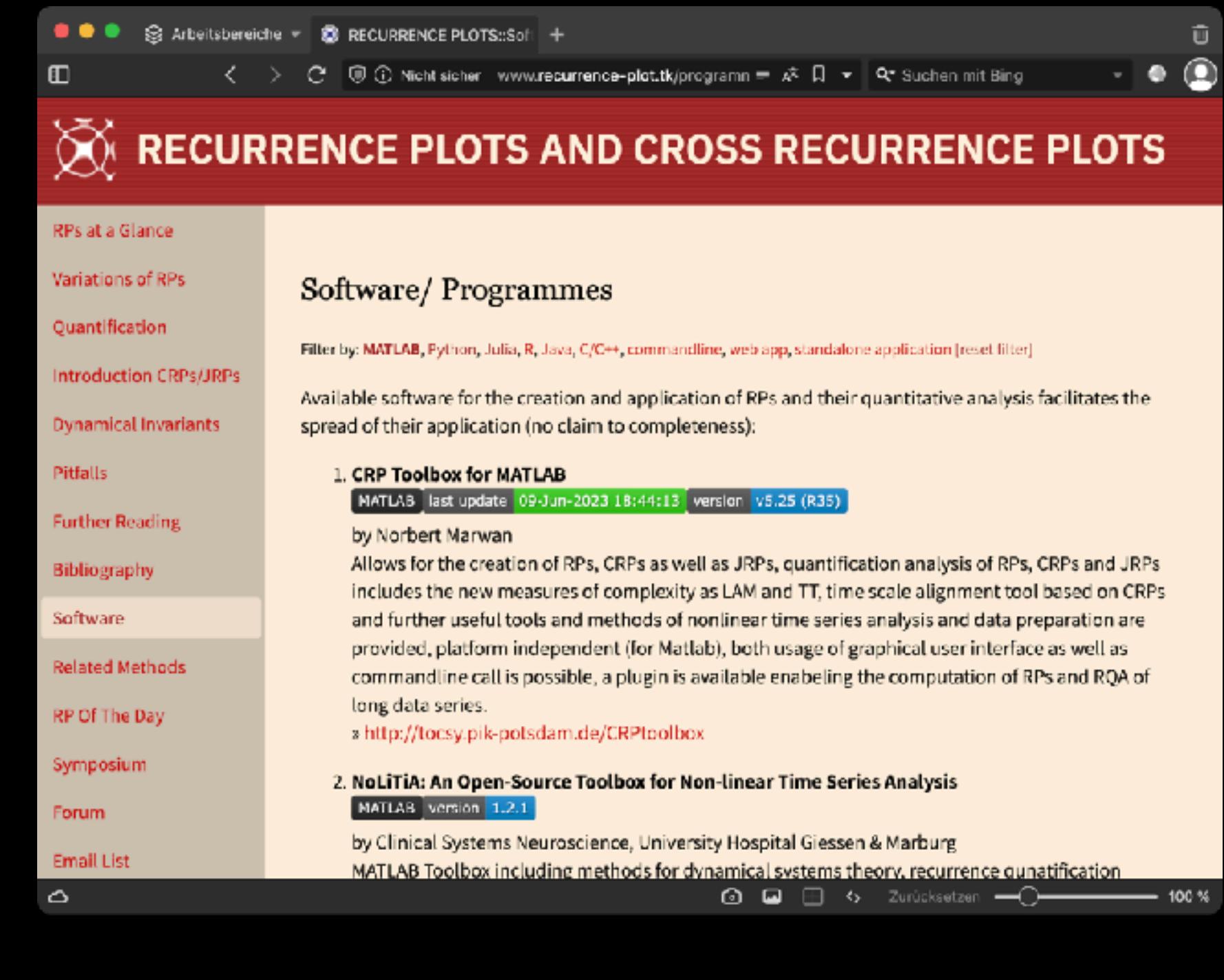


# SOFTWARE TOOLS

- MATLAB: CRP Toolbox
- Python: pyunicorn, igraph, networkx, pyRQA
- Julia: DynamicalSystems.jl

```
# dry run to pre-compile
x = embed(sol[1,1000:1500], 3, 6);
R = RecurrenceMatrix(x, 1.2, parallel=false);
Q = rqa(R, theiler = 1, onlydiagonal=false);

for (i,N_) in enumerate(N)
    x = embed(sol[1,1000:1000+N_], 3, 6);
    tRP_ = 0;
    tRQA_ = 0;
    for j in 1:K
        t1 = @elapsed R = RecurrenceMatrix(x, 1.2, parallel=false);
        t2 = @elapsed Q = rqa(R, theiler = 1, onlydiagonal=false);
        tRP_ = tRP_ + t1;
        tRQA_ = tRQA_ + t2;
        print(" " ,j, "\n")
    end
end
```



# CRP TOOLBOX FOR MATLAB

- Web site:  
<https://tocsy.pik-potsdam.de/CRPtoolbox> (CRP Toolbox)
- Access data:  
ID: **workshop**  
password: **potsdam**
- Download installation file **install.m**
- Call **install** from the MATLAB  
commandline

