

SIGNAL BASED METHODS

THE MATRIX PROFILE

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May 8th 2025

LECTURE SUMMARY

1. Introduction

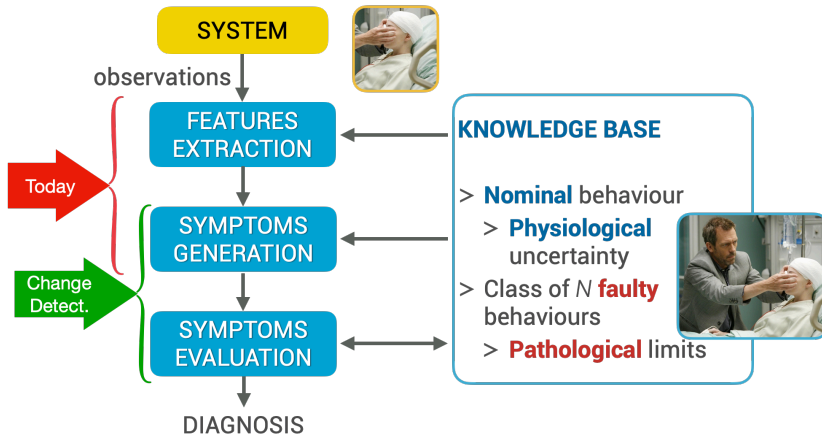
2. Definition of the MP

3. Conclusions

INTRODUCTION

INTRODUCTION

REMEMBER THE PARALLEL WITH MEDICAL DIAGNOSIS?



INTRODUCTION

THE MATRIX PROFILE: LOOKING FOR SELF SIMILARITIES IN TIME SERIES



Scene taken from the movie "The Matrix" (2003). Photo: Courtesy of Warner Bros

INTRODUCTION

THE MATRIX PROFILE: LOOKING FOR SELF SIMILARITIES IN TIME SERIES

Intuition: conservation is key

- ▶ If a pattern is conserved, there must be some mechanism that conserve it
- ▶ Question: what is conserved in a time series?
- ▶ Conservation \Rightarrow **healthy**

Examples

<i>Bengali:</i>	bābā	<i>Norwegian:</i>	papa
<i>Mandarin:</i>	baba	<i>Spanish:</i>	papá
<i>Indonesian:</i>	baba	<i>English:</i>	papa
<i>Turkish:</i>	baba	<i>Hindi:</i>	papa
<i>Polish:</i>	tata	<i>Xhosa:</i>	-tata

<https://www.cs.ucr.edu/~eamonn/MatrixProfile.html>

INTRODUCTION

THE MATRIX PROFILE: LOOKING FOR SELF SIMILARITIES IN TIME SERIES

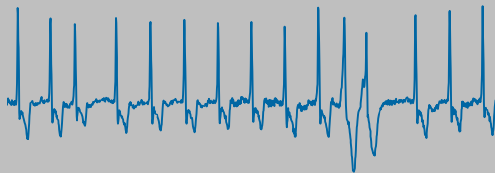
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Examples

 ECG signal



INTRODUCTION

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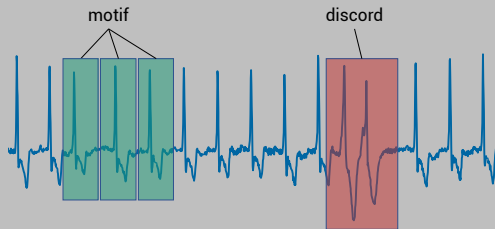
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Examples

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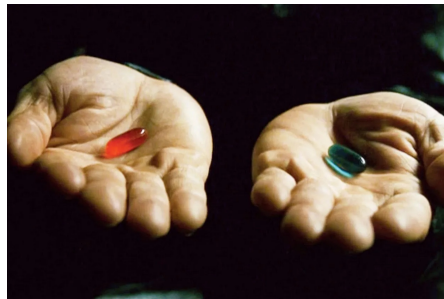


INTRODUCTION

THE MATRIX PROFILE (MP) IN PILLS

3 easy steps

- ✂ cut up the time series in pieces
- 🔍 compare each piece with every other piece
- 🧩 very similar \Rightarrow you found a pattern (**motif**)
very different \Rightarrow you found an anomaly (**discord**)

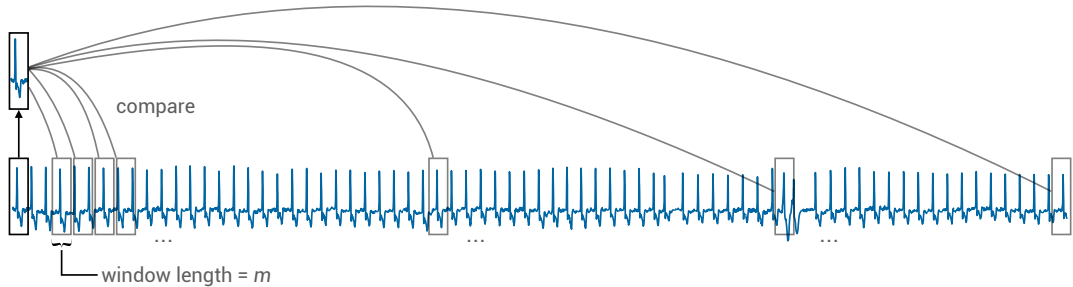


“Remember – all I am offering is the truth, nothing more.”

Photo: Courtesy of Warner Bros.”

INTRODUCTION

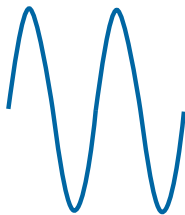
THE MATRIX PROFILE (MP) IN PILLS



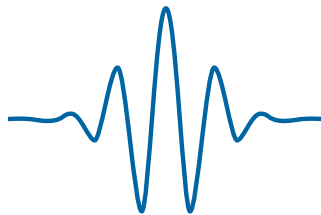
INTRODUCTION

THE MATRIX PROFILE (MP) AS A GENERALIZATION

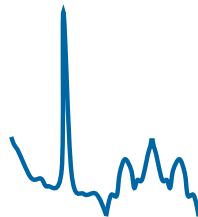
Sinusoid
(FFT)



Wavelet
(WT)



Shapelet
(MP)



INTRODUCTION

THE MATRIX PROFILE (MP): WHY?

PROS

- + domain agnostic
- + fast
- + only requires a single parameter.

CONS

- Basic assumption: repetition
- Basic assumption: steady-state
- Empirical choice of threshold

DEFINITION OF THE MP

DEFINITION OF THE MP

DISTANCE BETWEEN SUBSEQUENCES

- ▶ Let us consider two subsequences z_1 and z_2
 - ▶ Can be taken from same time series, or from different ones
- ▶ The **comparison** step in the MP is based on computing their distance

Definition (z-normalized Euclidean distance)

The **z-normalized Euclidean distance** of $z_1, z_2 \in \mathbb{R}^m$ is

$$d(z_1, z_2) \triangleq \sqrt{\sum_{i=1}^m (\bar{z}_1(i) - \bar{z}_2(i))^2} = \|\bar{z}_1 - \bar{z}_2\|_2$$

where the z-normalized values \bar{z}_1 and \bar{z}_2 are defined as

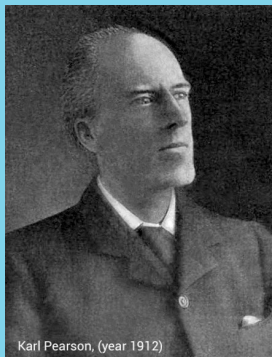
$$\bar{z}_1(i) \triangleq \frac{z_1(i) - \mu_1}{\sigma_1}, \quad \bar{z}_2(i) \triangleq \frac{z_2(i) - \mu_2}{\sigma_2}$$

and μ and σ denote, respectively, means and std. dev.

DEFINITION OF THE MP

DISTANCE BETWEEN SUBSEQUENCES

It is all Pearson's



Karl Pearson, (year 1912)

- ▶ It holds $d(z_1, z_2) = \sqrt{2m(1 - \text{corr}(z_1, z_2))}$
- ▶ where **corr** is the **Pearson's Correlation Coefficient**

$$\text{corr}(z_1, z_2) \triangleq \frac{\sum_{i=1}^m z_1(i)z_2(i) - m\hat{\mu}_1\hat{\mu}_2}{(m-1)\hat{\sigma}_1\hat{\sigma}_2}$$

- ▶ with $\hat{\mu}$ and $\hat{\sigma}$ being, as usual, sample means and std. devs.

DEFINITION OF THE MP

PAIRWISE DISTANCE COMPUTATION – EXAMPLE

Pairwise Euclidean Distance

0	1	3	2	9	1	14	15	1	2	2	10	7
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0	1	3	2	9	1	14	15	1	2	2	10	7
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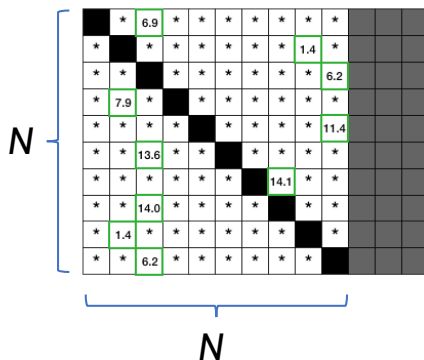
0.0	7.4	6.9	14.7	19.3	17.7	19.9	15.0					
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https://stumpy.readthedocs.io/en/latest/Tutorial_The_Matrix_Profile.html

DEFINITION OF THE MP

DISTANCE MATRIX COMPUTATION – EXAMPLE

Matrix Profile



6.9
1.4
6.2
7.9
11.4
13.6
14.1
14.0
1.4
6.2

Minimal
distances (MP)

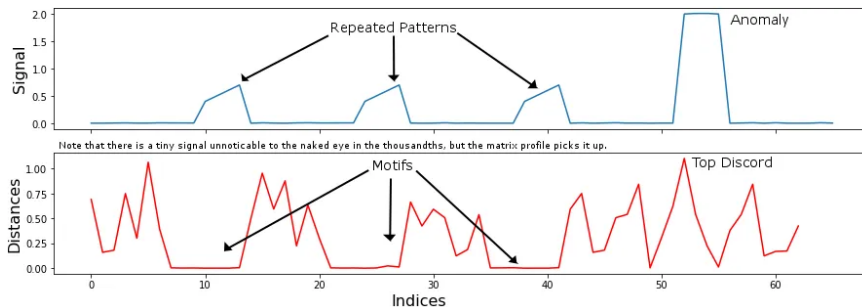
3
9
10
2
10
3
8
3
2
3

Index
profile (IP)

https://stumpy.readthedocs.io/en/latest/Tutorial_The_Matrix_Profile.html

DEFINITION OF THE MP

MATRIX PROFILE – EXAMPLE



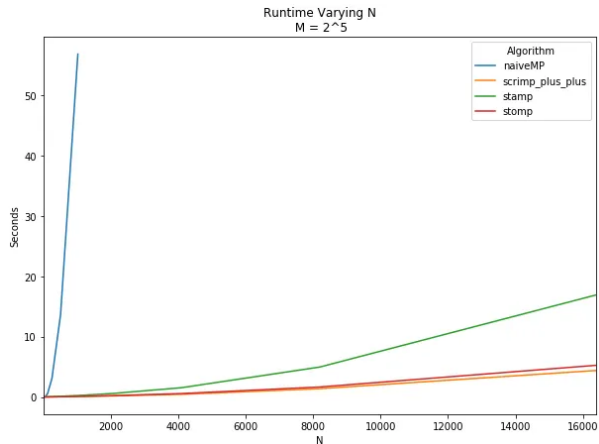
<https://towardsdatascience.com/introduction-to-matrix-profiles-5568f3375d90>

DEFINITION OF THE MP

MATRIX PROFILE – COMPUTATIONAL COMPLEXITY

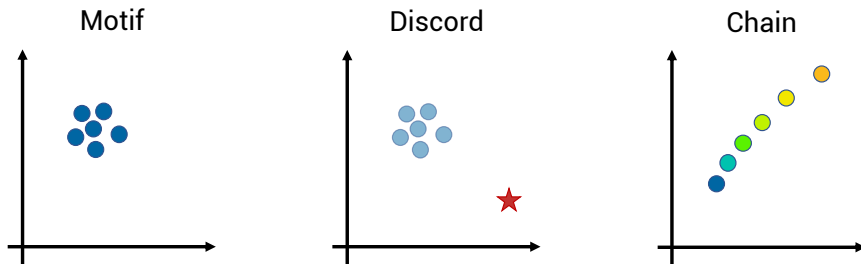
- ▶ Vanilla computation is $\mathcal{O}(n^2)$
- ▶ Efficient algorithms exist with $\mathcal{O}(n \log n)$ (similar to FFT)
- ▶ i.e. SCRIMP, STAMP, STOMP and GPU versions of that

<https://towardsdatascience.com/introduction-to-matrix-profiles-5568f3375d90>



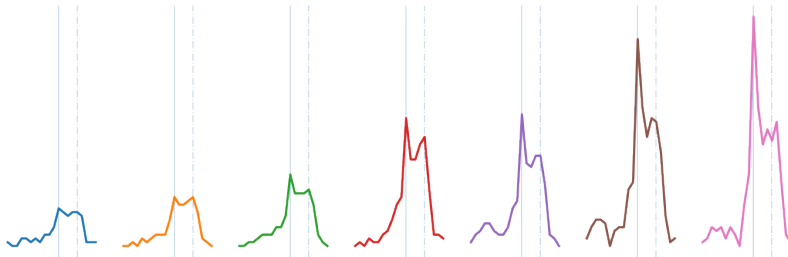
DEFINITION OF THE MP

MOTIFS, DISCORDS AND CHAINS



DEFINITION OF THE MP

CHAIN – EXAMPLE



https://stumpy.readthedocs.io/en/latest/Tutorial_Time_Series_Chains.html

CONCLUSIONS

CONCLUSIONS

IN THIS LECTURE WE COVERED

- ▶ **Introduction to MP**
- ▶ MP can compute very fast the self similarity of a signal to itself or to a nominal one
- ▶ Assuming your process is in/moving across a finite number of steady states, the MP can detect anomalies

Next lecture: **AI & ML for signal-based detection**

CONCLUSIONS

FURTHER READING

- ▶ STUMPY, a Python toolbox for MP
- ▶ MP page at UC Riverside
- ▶ Chin-Chia Michael Yeh et al. “Matrix Profile I: all pairs similarity joins for time series: a unifying view that includes motifs, discords and shapelets”. In: *2016 IEEE 16th international conference on data mining (ICDM)*. Ieee. 2016, pp. 1317–1322
- ▶ ...
- ▶ Sadaf Tafazoli and Eamonn Keogh. “Matrix Profile XXVIII: Discovering Multi-Dimensional Time Series Anomalies with K of N Anomaly Detection”. In: *Proceedings of the 2023 SIAM International Conference on Data Mining (SDM)*. SIAM. 2023, pp. 685–693

CONCLUSIONS

THANK YOU FOR YOUR ATTENTION!

For further information:
Course page on **Brightspace**
or
our **MS Team**