

#### MAP syntax diagrams: an online tool to visualise, describe, and compare Ancient divine onomastic sequences

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SUMMARY

MAP syntax diagrams is an online R Shiny application for studying Ancient divine onomastic sequences. It uses the format defined by the "Mapping Ancient" Polytheisms" project to:

- represent sequences and
- describe semantic relationships between the parts of the names of gods (called "elements" in the MAP terminology).

The focus is on the **structure** of the sequences, not their content. These structures are modelled as graphs (a mathematical object composed by a set of nodes and a set of edges), so that they can be visualised, characterised, and compared using quantitative measurements.

### FROM SEQUENCES TO GRAPHS

In the MAP format, the sequence Δηώ καὶ Κόρη Ἐλευσινίαι is represented as  $[\Delta \eta \dot{\omega} + K \dot{\rho} \eta] # Ἐλευσίνιος, α, ον$ and coded as

[1+2]#3".



Έλευσίνιος, α, ον

In MAP syntax diagrams, this sequence is modelled by a graph. The [ and ] symbols enclose a syntagma,

+ indicates a relation of coordination between two elements, *#* indicates a qualification.

## WORKFLOW

### **APPLICATION MAIN FEATURES**

- Selection of subsets of sequences for comparison
- Summary statistics: diversity of the elements (Shannon-Weaver entropy) and



Online extraction from the MAP database<sup>1</sup>

Recoding of the relevant variables

https://base-map-polytheisms.huma-num.fr

# COMPARING GODS: APOLLO & APHRODITE

Greek sequences (up to 11 elements) from Cyprus related to Aphrodite (n=92) and Apollo (n=46) are compared. The statistics indicate that Apollo's sequences have more diverse elements (diversity index: 1.81 vs 1.19) and subsequences (median transition rate: 0.11 vs 0.01). However, using the diagrams to study the differences (coloured in red) we can see that, for example, the complex nested structures ("[[]]") observed in Aphrodite's sequences are **absent** in Apollo's sequences.



Input in *MAP* 

and analysis

syntax diagrams

structural properties (network metrics)

• Measurement of the difference between the subsets graph (Euclidean and "graph diffusion kernel" distances); display of a hierarchical clustering of the distances (for 4 and more subsets)

• Generation of **diagrams**:

1) to graphically summarise a set of sequences and

- 2) to highlight the differences between two subsets
- **Export** of statistics (in csv format) and diagrams (in svg format)

### STRUCTURAL CHANGE THROUGH TIME

Greek and Phoenician sequences from Cyprus are compared across three periods (see table). This demonstrates an increase in diversity and **complexity** over time, as reflected by:

total number of sequences	64	105	124
number of selected sequences	63	104	122
number of different sequences	8	13	17
sequence max. length	11	19	23
sequence median length	3	3	3
diversity of the elements	3.41	3.16	3.36
number of nodes	24	49	72
number of different symbols	5 / 9	6 / 9	9 / 9
number of terminal symbols	3	3	5

-500/-300 -300/-1

1/400

• an increase in the length of number of articulation points degree centralisation 0.03 0.04the sequences, in the numbers median transition rate 0.01 0.05 0.01 of different symbols, the different sequences, and the nodes in the structure. • a decrease in the centralisation value, meaning that the edges are less concentrated on some particular elements. However, the **diversity** of the **elements** composing the sequences (nouns, adjectives, etc.) is virtually **unchanging**, reflecting stability.

• MAP project: https://map-polytheisms.huma-num.fr • Application: https://analytics.huma-num.fr/Sebastien.Plutniak/map-syntax-diagrams • Code: https://github.com/sebastien-plutniak/map-syntax-diagrams DOI 10.5281/zenodo.4443133 • Manual: Plutniak. S. 2020. MAP Syntax diagrams : visualiser, décrire et comparer les séquences onomastiques divines de l'Antiquité. https://hal.archives-ouvertes.fr/hal-02532617