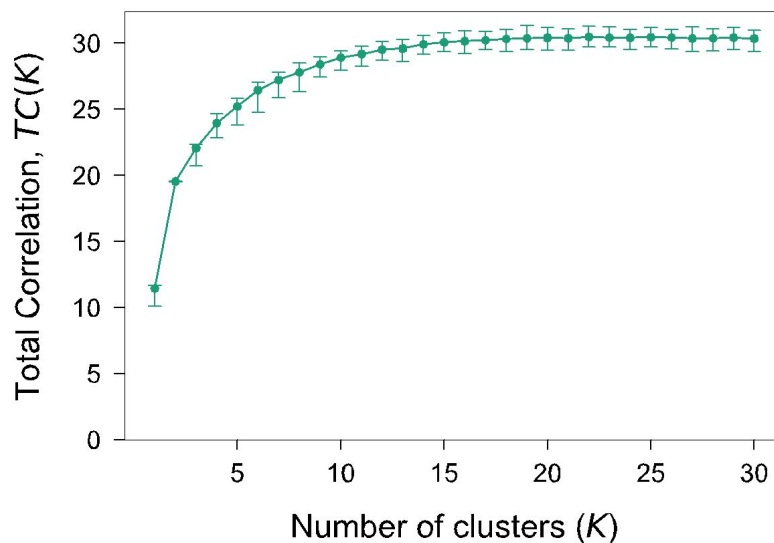


Results for the new data in “maps of myths”

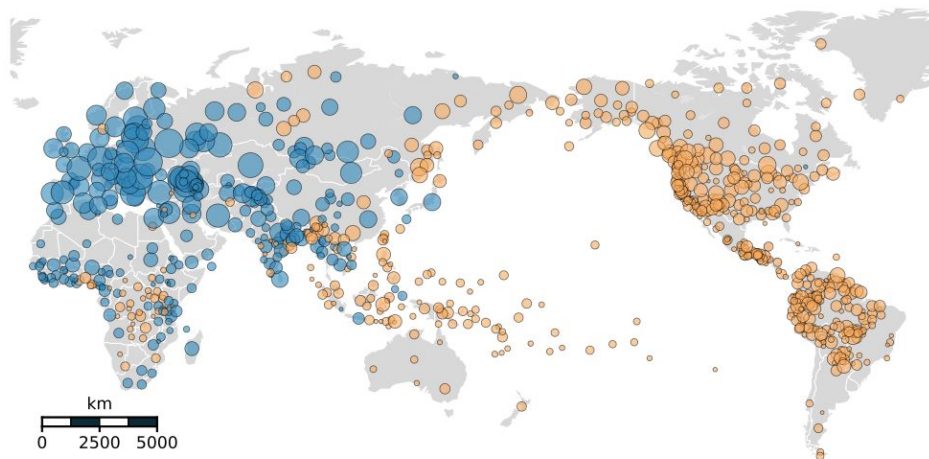
- 958 traditions with their geographical locations (longitude and latitude)
- 2583 motifs
 - “I120a” is duplicated. It has two ids, 103927 / 103940
 - 103940 is taken as it has more traditions than 103927
- Out of 958 traditions, 901 traditions that have at least ten motifs are selected for CorEx analysis. Thus, the size of the new binary matrix is 901 by 2583.
- Previously, we use the data of 683 traditions and 1638 motifs.

1. Total correlation still saturates after ten clusters

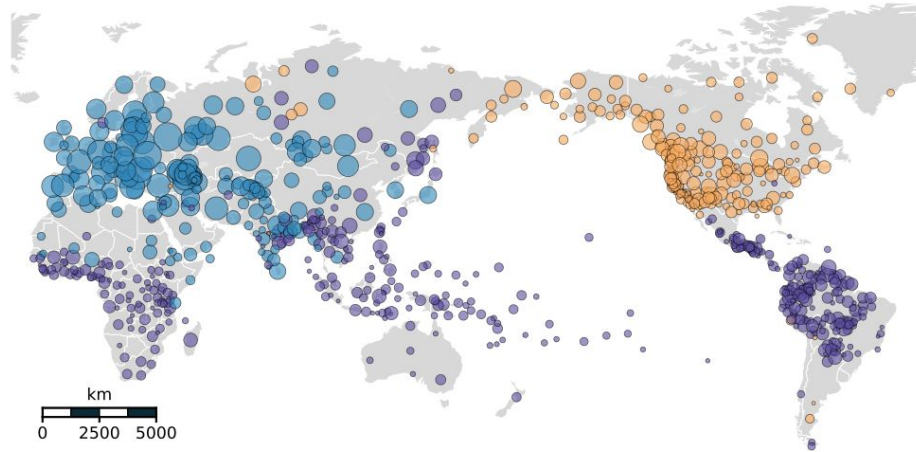


2. CorEx clusterings ($K=2, 3, 4, \dots, 12$)

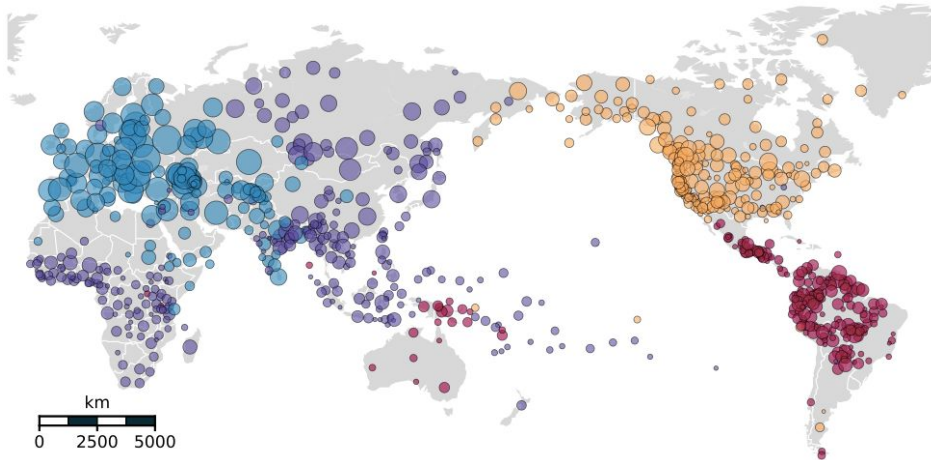
- $K=2$



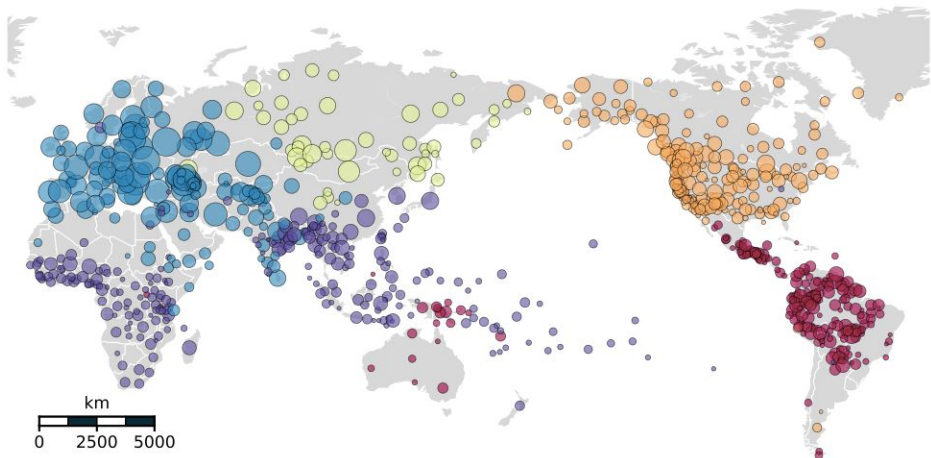
- K=3



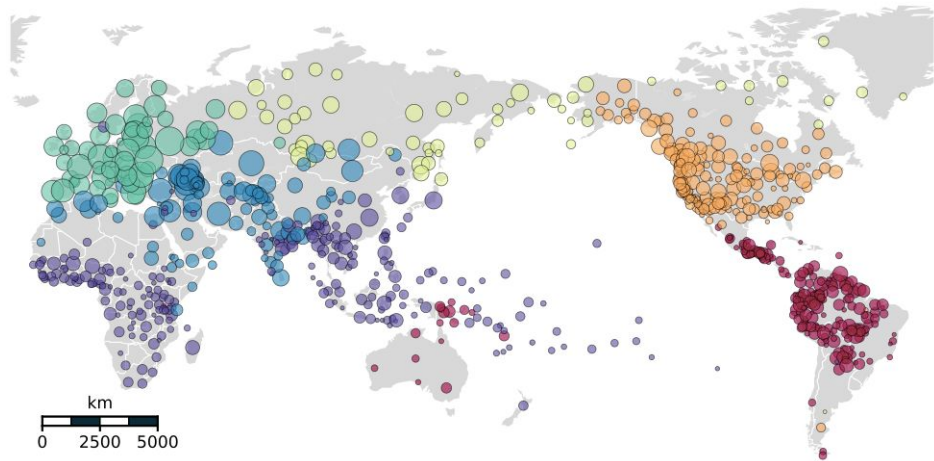
- K=4



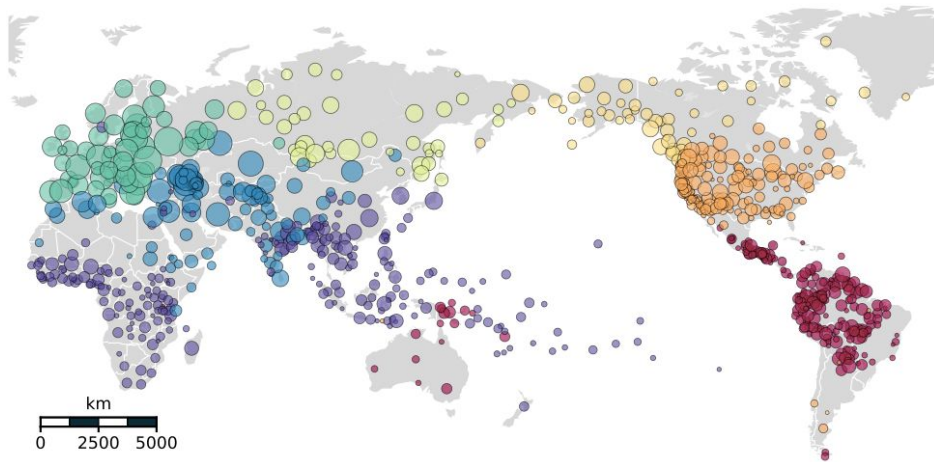
- K=5



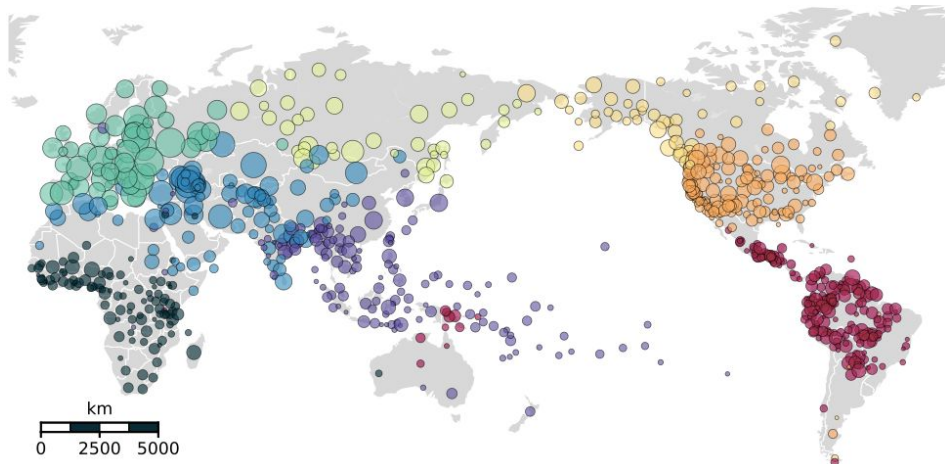
- K=6



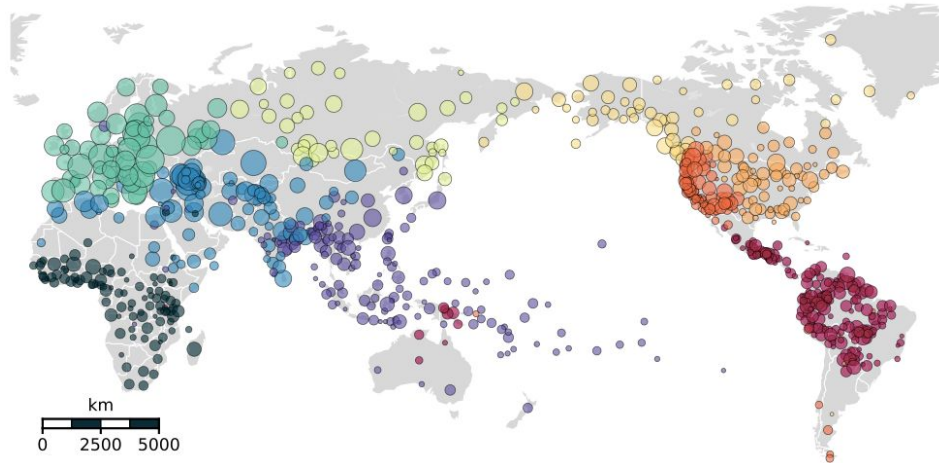
- K=7



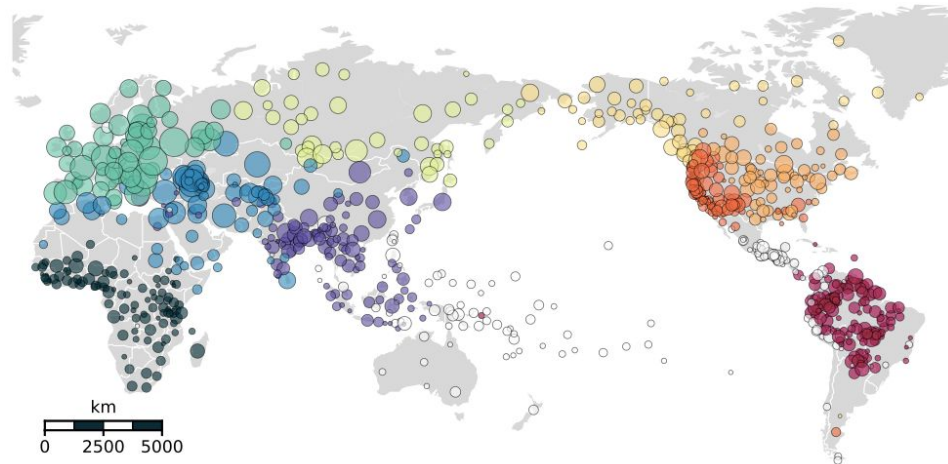
- K=8



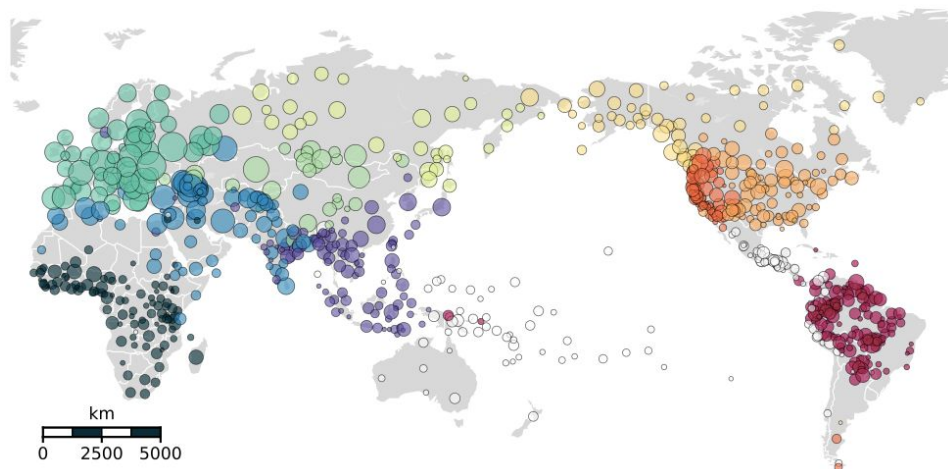
- K=9



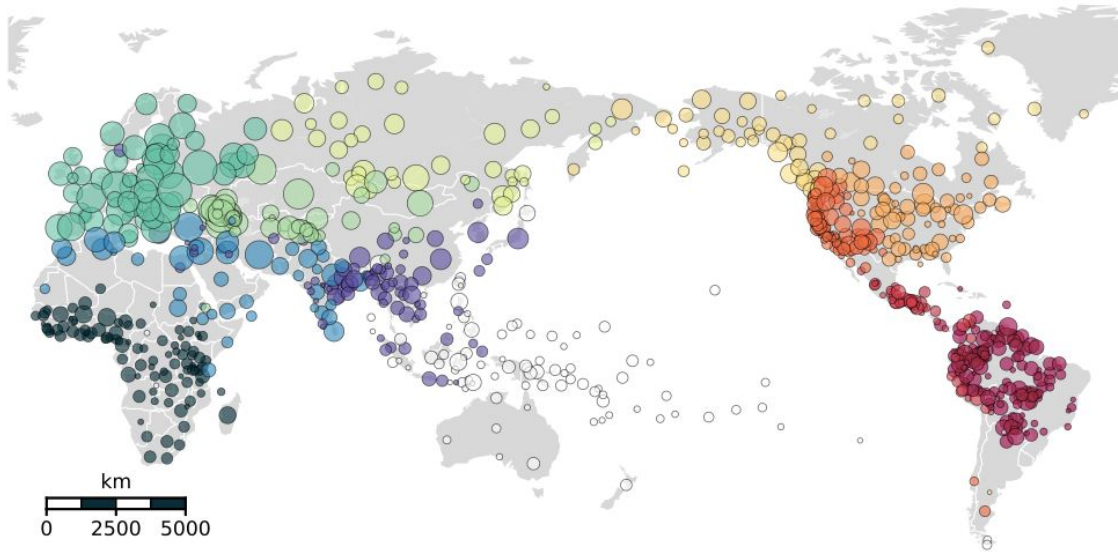
- K=10



- K=11



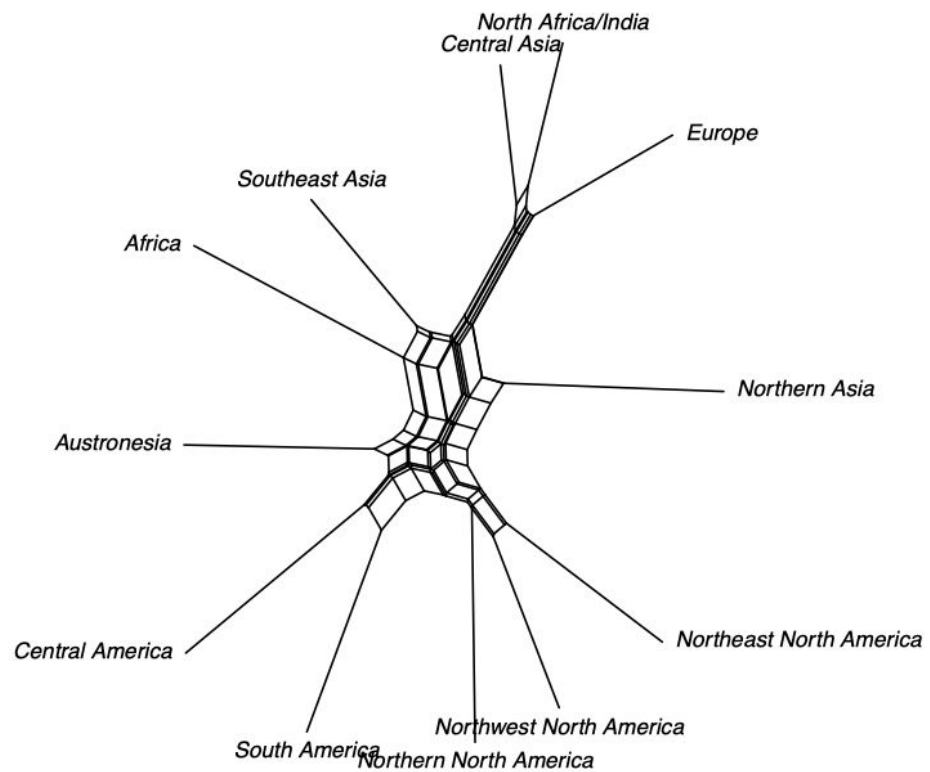
- K=12



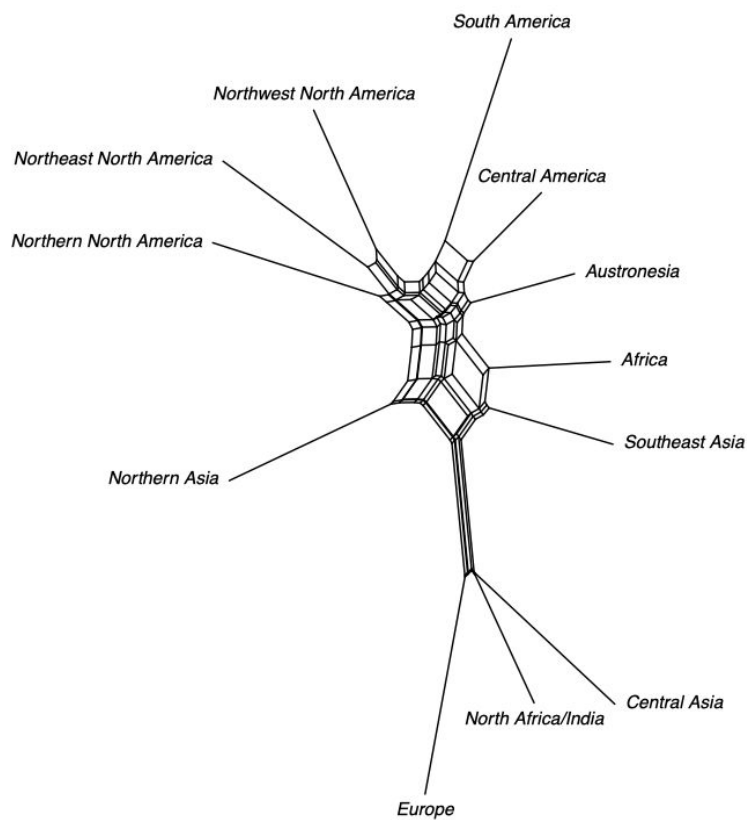
- Clusters
 - A new cluster that covers North Africa and India emerges
 - Two clusters in Western North America become a cluster in the new data
- Things to check
 - Phylogenetic network
 - AMOVA, NMI with bioregions (Needs to collect language family affiliation for new traditions)
 - Similarity decaying
 - Spatial patterns (Motif classification was changed by Berezkin, <http://mapsofmyths.com/motif-types>)
 - Multiscale Backbone

3. Phylogenetic network

- Cosine distance

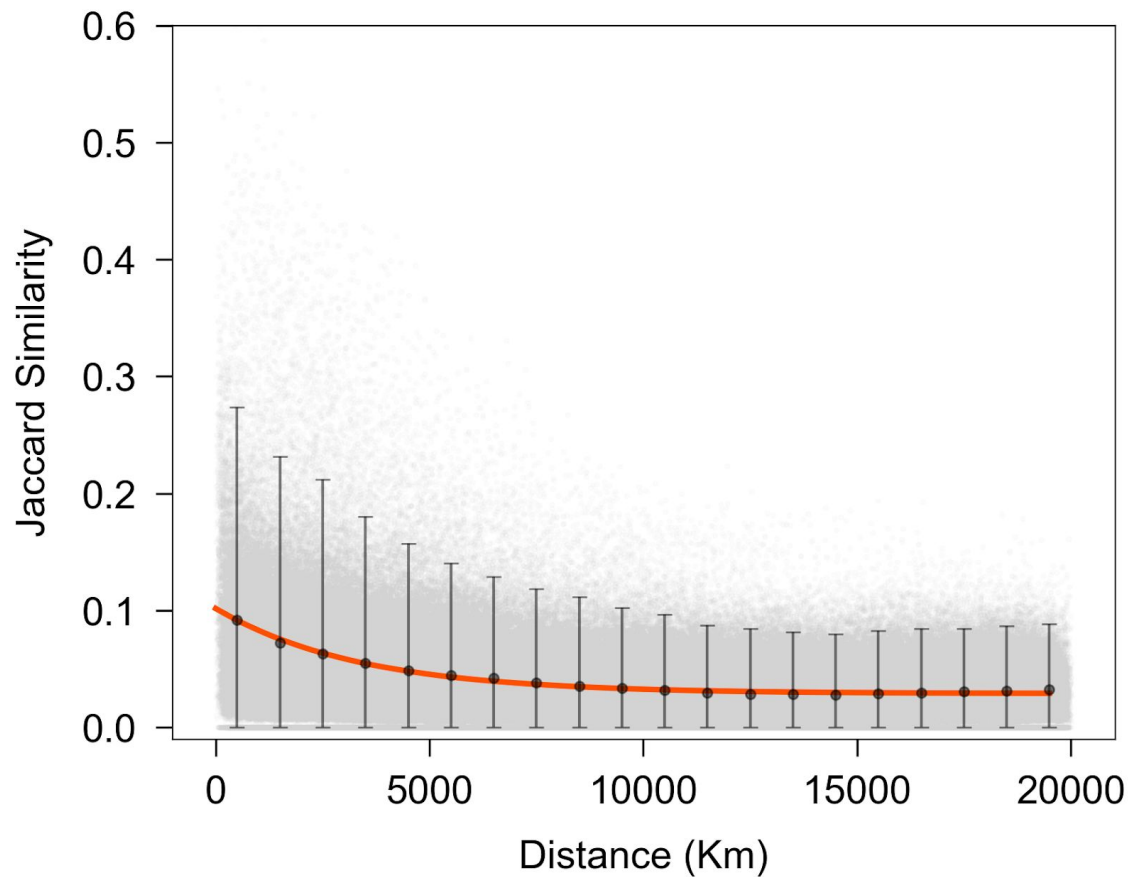


- Hamming distance



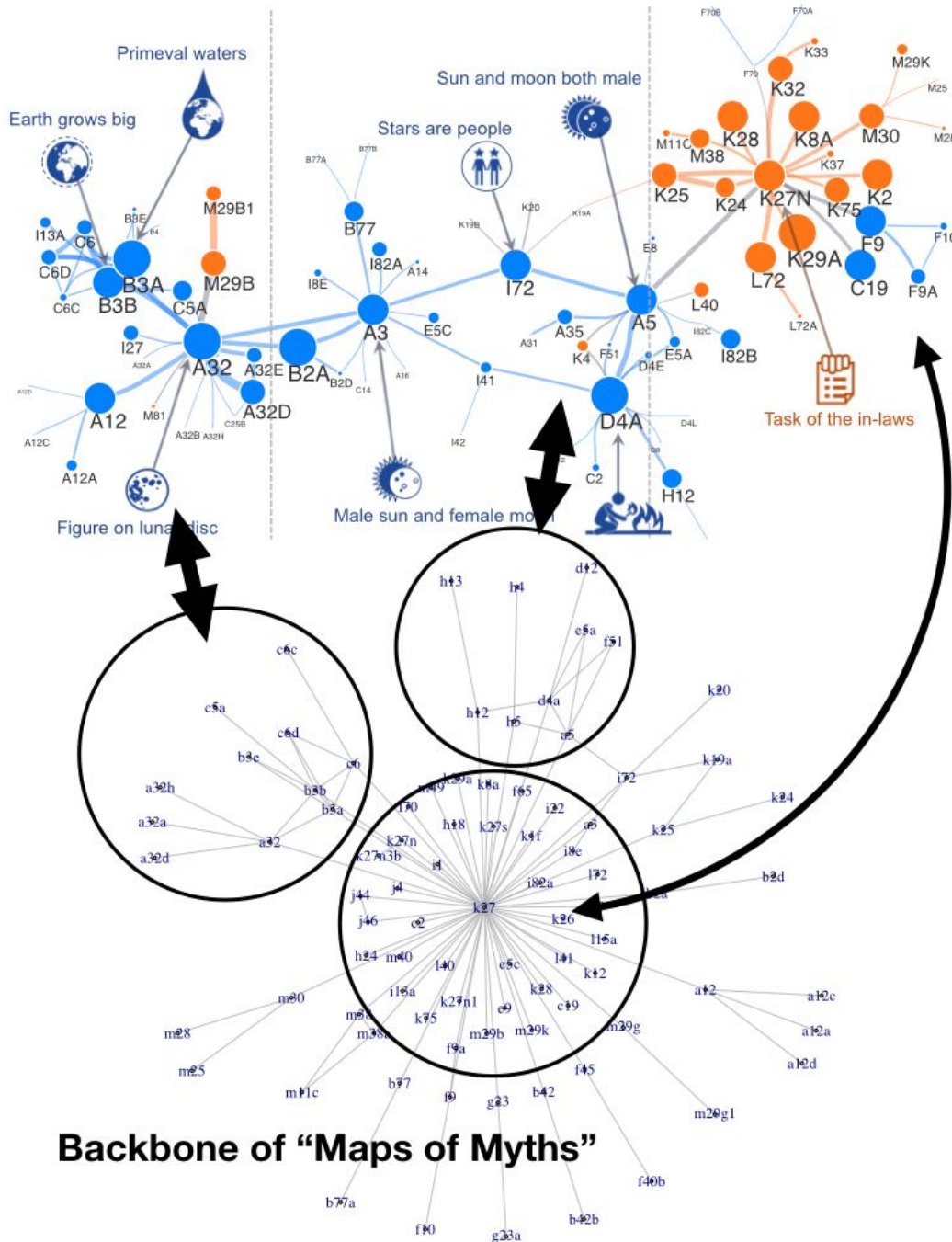
- Network structures of two distances are similar
 - Africa – Austronesia
 - Africa – Southeast Asia – Central Asia, ...

4. Similarity decaying



Characteristic distance is 3364 km, which is larger than the previous characteristic distance 2572 km.

5. Motif backbone



6. Summary

- CorEx clusterings, phylogenetic networks, and similarity decaying patterns are similar in both data, the previous data we used and the new data from “maps of myths”
- However, due to the increase of data coverage and the changes in motif classification, additional works would be needed to obtain figures for our manuscript, if we use the new data.