A Gromov-Wasserstein Approach for African Food Web Network Analysis RICE

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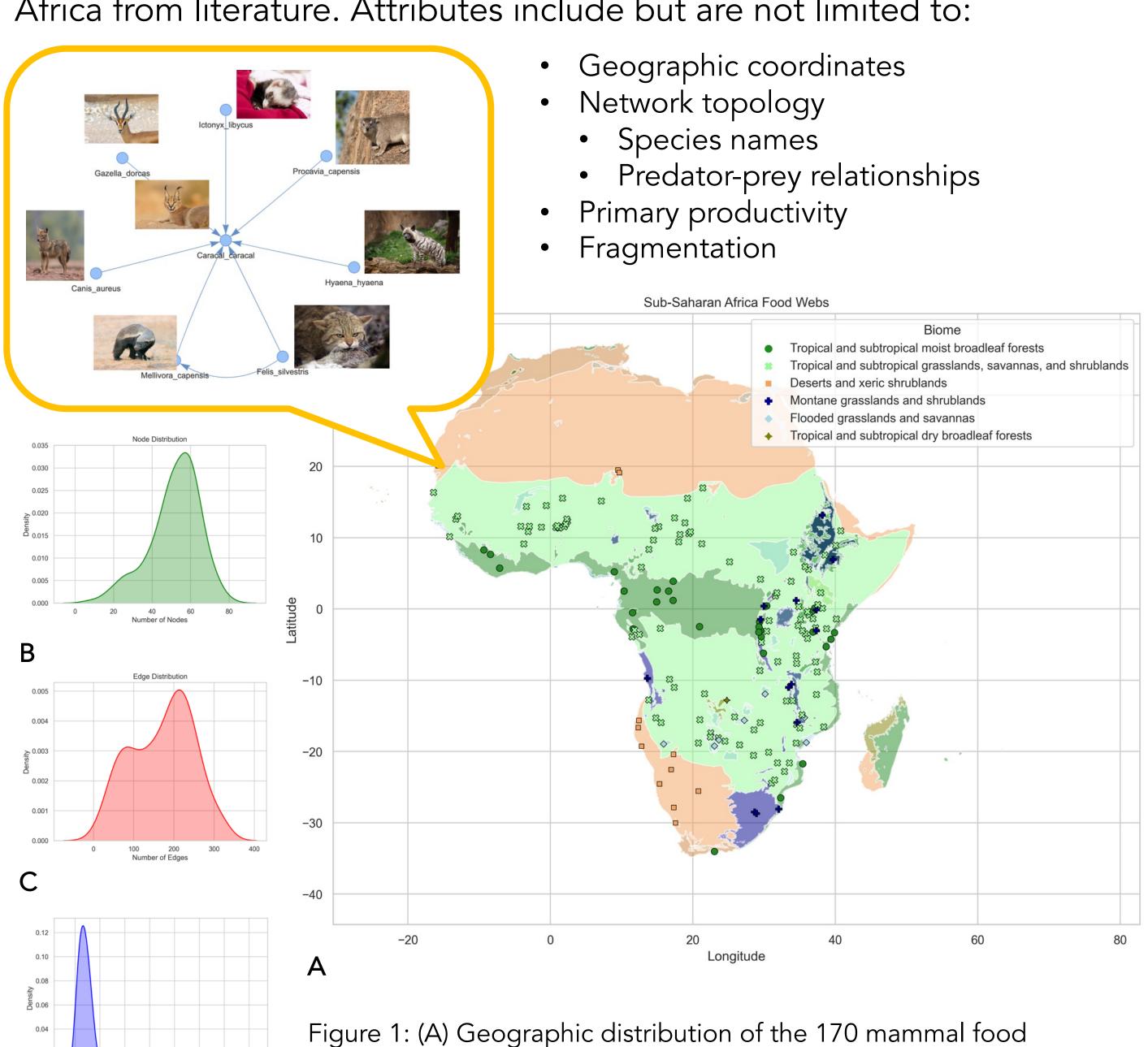


Background

- Predator-prey interactions in **food webs** play a critical role in maintaining biodiversity.
- Human-induced habitat loss threatens these interactions and overall ecosystem health.
- Understanding mammal food webs can inform conservation efforts.

Data

Our group curated 170 mammal food web networks of Sub-Saharan Africa from literature. Attributes include but are not limited to:



Problem of Interest

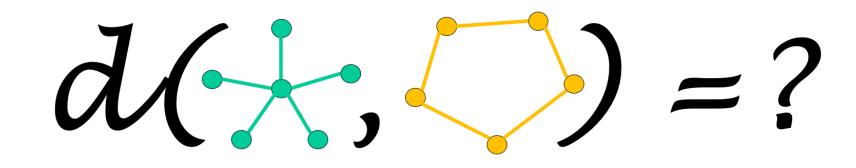
(D) The degree distribution across all networks.

webs colored by biome type. (B) The node distribution across all

food web networks. (C) The edge distribution across all networks.

A Topological Data Analysis Toolkit for Food Webs

But... how do we compare networks?



Existing methods (ex. graph kernels, graph neural networks) are either computationally intensive, difficult to interpret, applicable only to a singular network, or does not apply to networks with different sizes.

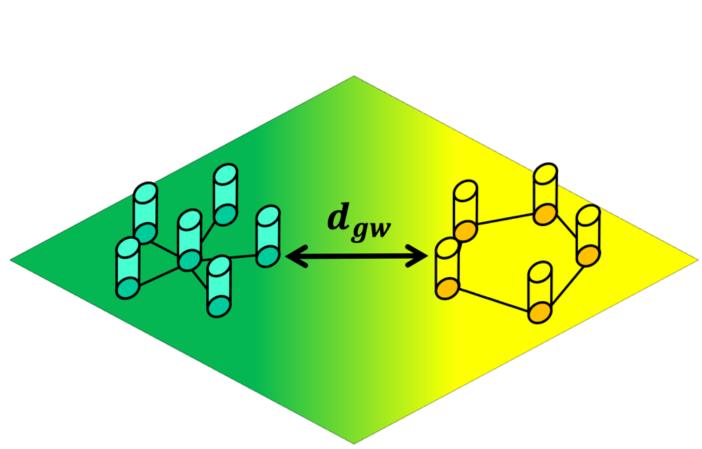
Optimal Transport

Definition 1: Let (X, w_x, μ_x) and (Y, w_Y, μ_Y) be two measure networks, we define the network Gromov-Wasserstein (GW) distance¹ as

$$d_{gw}(X,Y) := \min_{\mu \in \mathcal{L}(\mu_X,\mu_Y)} \int_{X \times X} \int_{Y \times Y} |w_X(x,x') - w_Y(y,y')| d\mu(x,y) d\mu(x',y')$$

where $\mathcal{L}(\mu_x, \mu_y)$ is the set of measure couplings between μ_x and μ_Y .

Theorem 1: The network GW distance is a pseudo-metric on the space of networks ${\mathcal N}$



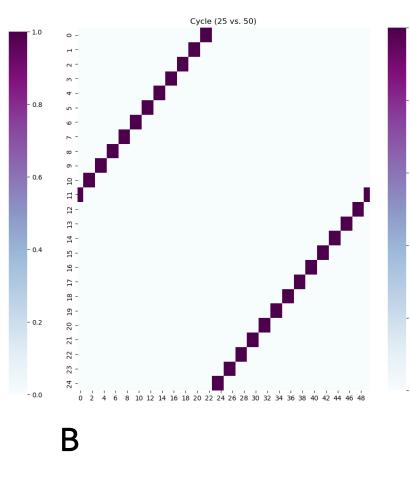
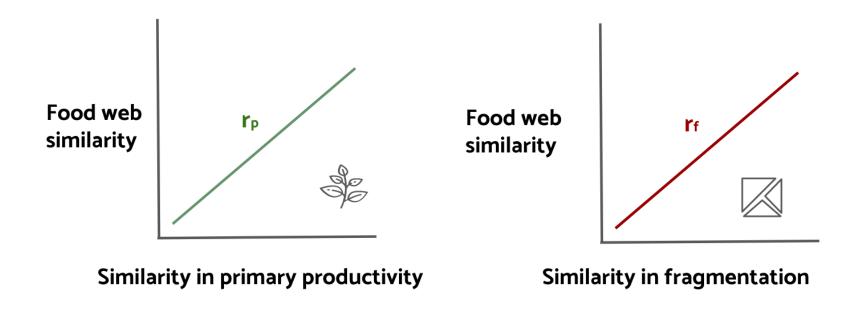
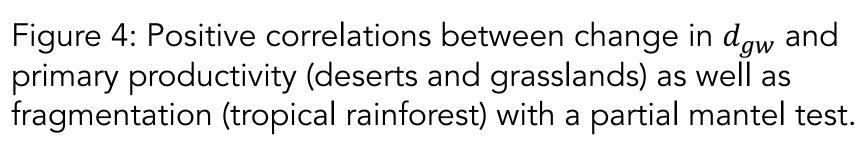


Figure 2: Conceptual illustration of the metric space of networks characterized by d_{aw} .

Figure 3: The optimal transport maps between (A) a 25-path and 50-path graph, and (B) a 25-cycle and 50-cycle graph.

Results: Gromov-Wasserstein Distance





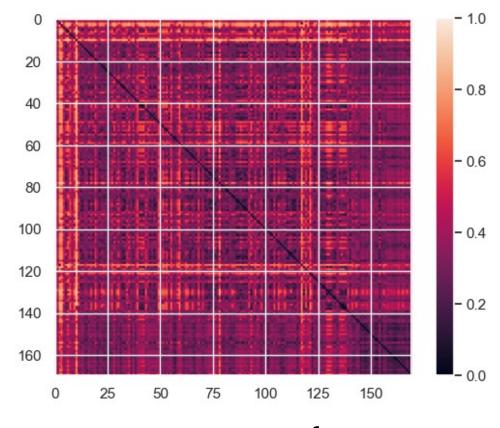


Figure 5: Heatmap of pairwise d_{qw}

Results: Gromov-Wasserstein Barycenter

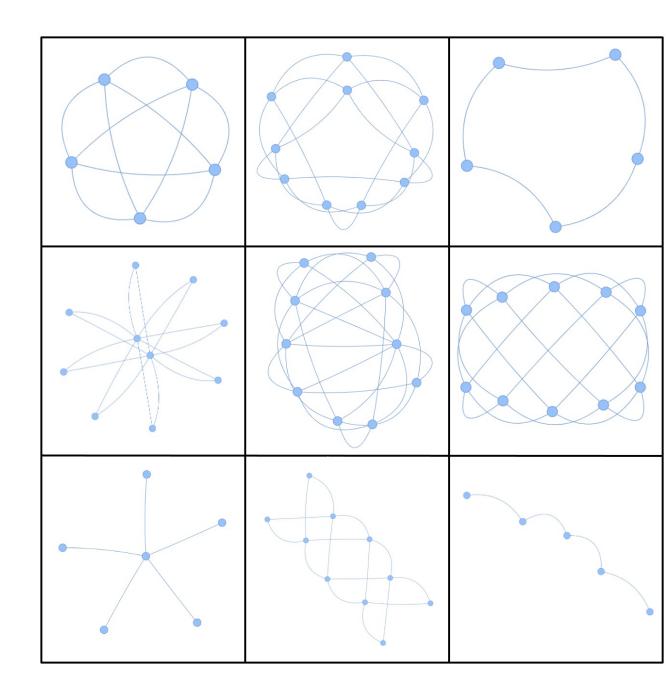


Figure 6: GW barycenter of complete graph (top-left), cycle graph (top-right), star graph (bottom-left), and path graph (bottom-right).

Definition 2: Let $\{C_s, p_s\}_{s=1}^S$ be S pairs of cost matrices C_s and node distribution p_s . The Gromov-Wasserstein Barycenter $^2\mathcal{B}$ of $\{C_s, p_s\}_{s=1}^S$ is defined as

$$\mathcal{B} := \min_{C \in \mathbb{R}^{n \times n}} \sum_{S} \lambda_{S} d_{gw}(C, C_{S}, p, p_{S})$$

where λ_s is the weight of (C_s, p_s) .

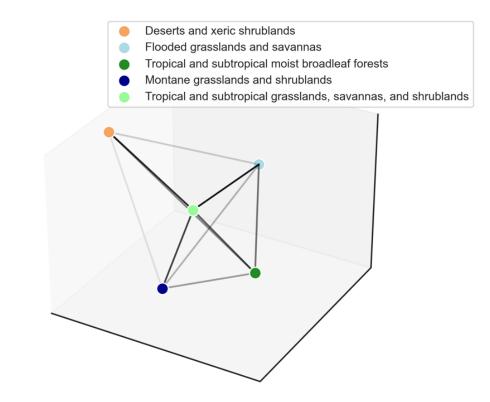
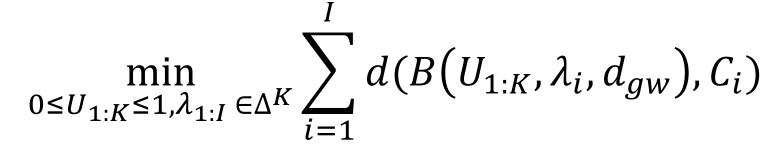


Figure 7: Distance between Barycenters of different biomes. Edge strength scaled with d_{qw} .

Results: Graph Factorization

Definition 3: Let $\{U_k, \lambda_k\}_{k=1}^K$ be K pairs of cost matrix and weight for the support networks. The Gromov-Wasserstein Factorization (GWF)³ framework for *I* observed networks is defined by



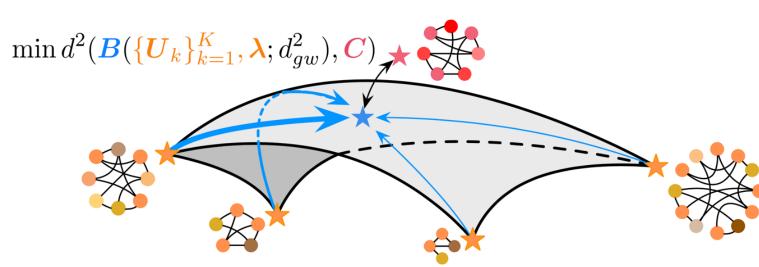


Figure 8: Conceptual illustration of the GWF framework. Courtesy of Xu, et al.

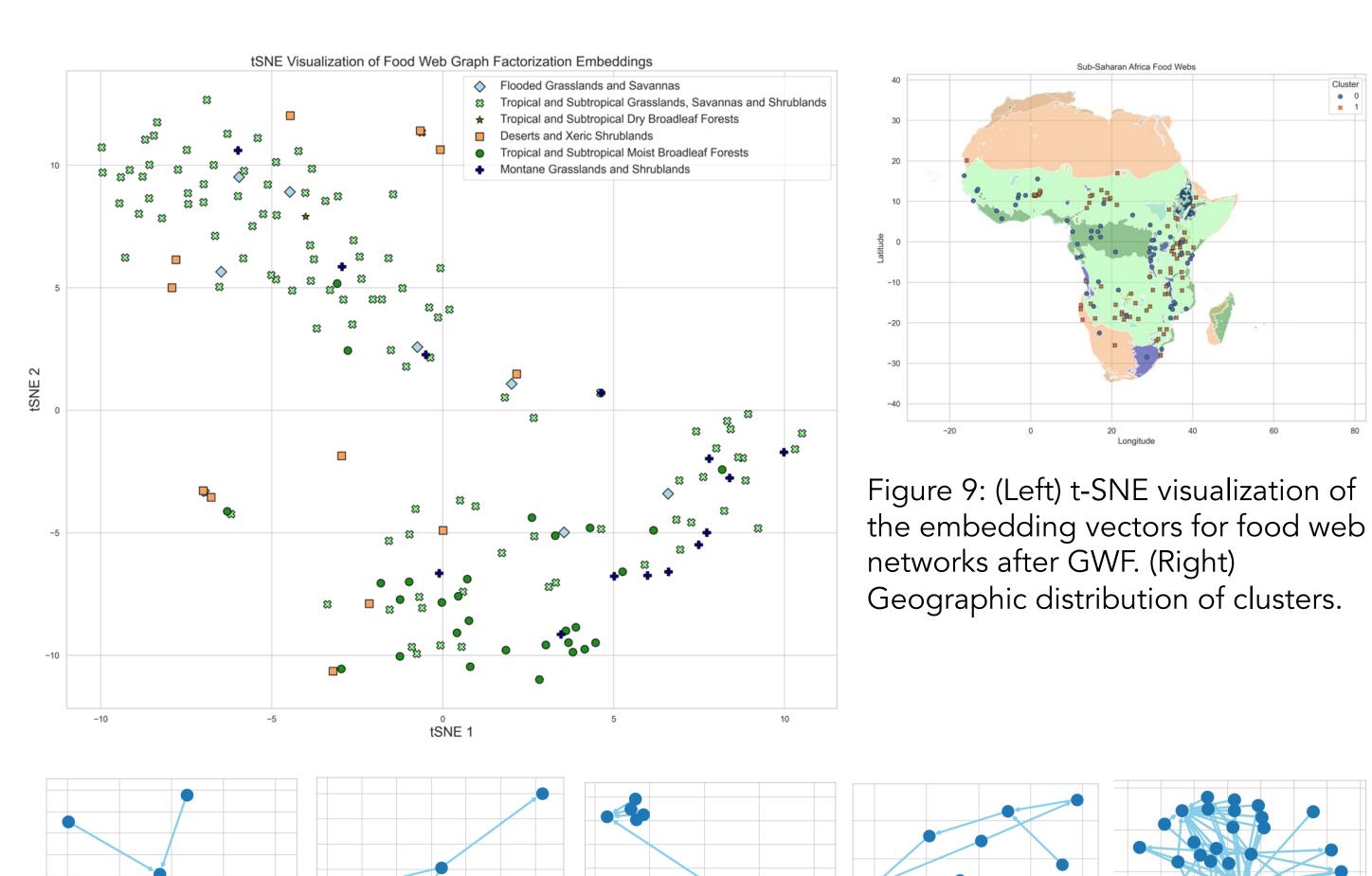


Figure 10: Sample of top-weighted learned support networks across the observed clusters.

Future Works

- Endow the food web networks with node distributions based on the flow of energy within an ecosystem.
- Incorporate features beyond network topology via the Fused Gromov-Wasserstein distance.
- Conduct a **theoretical analysis** of the growth in d_{aw} as a function of difference in network sizes.

References

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- Peyré, G., Cuturi, M., and Solomon, J. Gromov-Wasserstein averaging of kernel and distance matrices. In Proceedings of the 33rd International Conference on Machine Learning, 2016.
- Xu, H., Liu, J., Lu, D., and Carin, L. Representing graphs via gromov-wasserstein factorization. IEEE Transactions on Pattern Analysis and Machine Intelligence, 2022.



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