

Predicting ecological interactions across space

Kesem Abramov ¹, Barry Biton ¹, Geut Galai ¹, Rami Puzis ² and Shai Pilosof ^{1,3}



¹Department of Life Sciences; ²Department of Systems Information and Software Engineering; ³The Goldman Sonnenfeldt School of Sustainability and Climate Change Ben-Gurion University of the Negev, Be'er Sheva, Israel

What we observe in nature is never the full picture.

When we study ecological communities, many species interactions go undetected. This makes it difficult to fully understand how ecosystems function.

Link prediction tools can help fill these gaps,

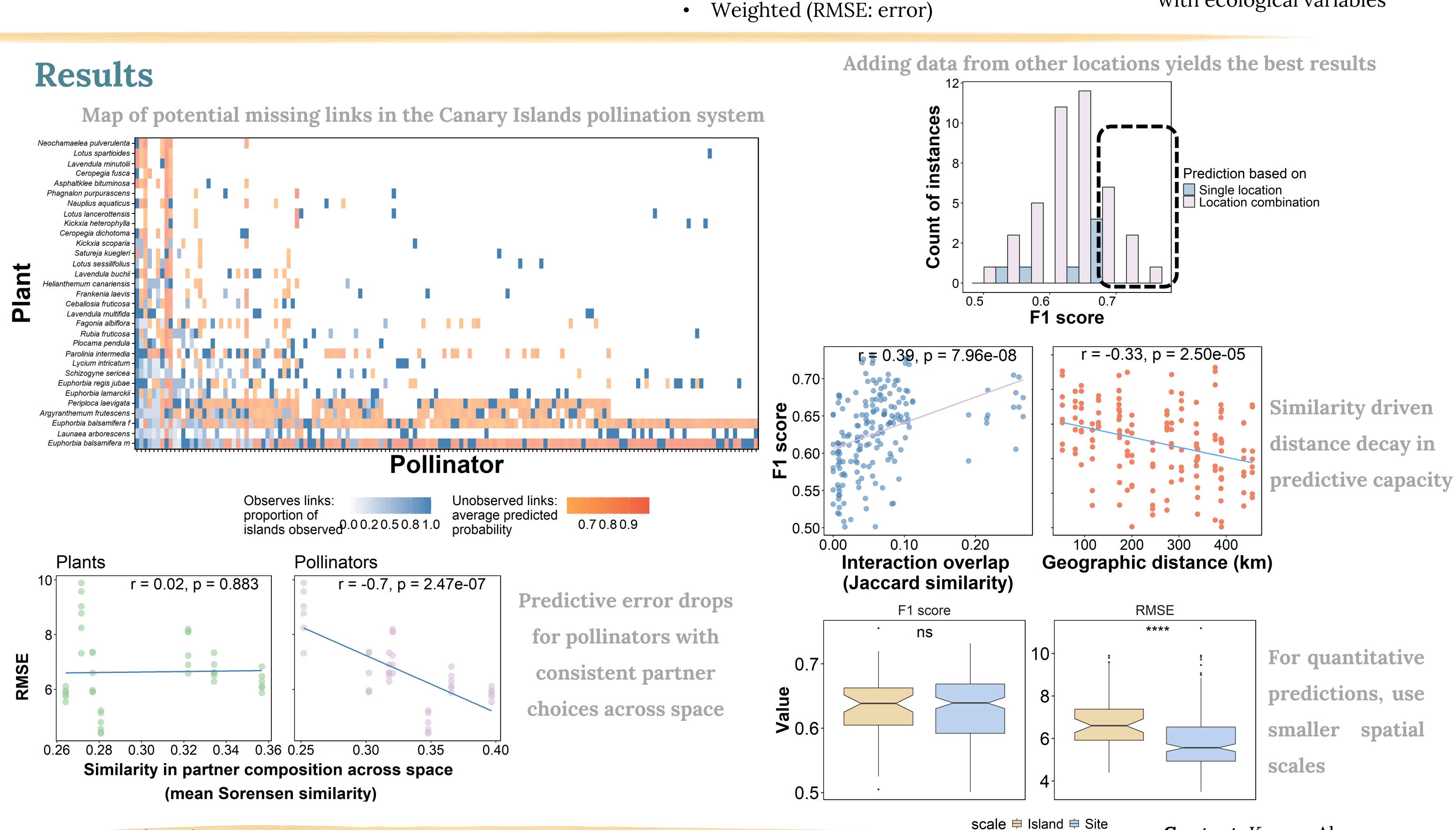
but they often depend on knowledge of species traits and ignore how interactions vary across space.

Goals

We set out to develop a method for predicting missing links in a target location using information from other locations, relying solely on network structure.

We then explored what factors influence how well we can predict these links across space.

Methods Developed predictive pipeline CANARY ISLANDS Pollination networks is 7 locations Removing links randomly Combining interactions • 2 sites in each location F1 score Tenerife Teno from a target location from two locations 0.65 Fuerteventura 0.60 **Combined matrix** Added location Target location 0.55 La Gomera Network Tenerife South size 800 Gran Canaria 700 El Hierro 600 Shared W 500 interactions 400 Unique to target site Unique to added site Western Sahara **E**cological inference **Evaluation** of Predicting links based on Singular prediction quality Contrasting prediction quality Value Decomposition (SVD) Binary (F1 score: quality) with ecological variables



Conclusions

We can make ecologically relevant predictions based solely on network structure

We are more likely to predict correctly:

- · By adding data from a similar, closer network
- Interactions that involve loyal partners

Contact: Kesem Abramov kesemk@post.bgu.ac.il