

# stemflow: A Python Package for Adaptive Spatio-Temporal Exploratory Model

Yangkang Chen<sup>1,2</sup>, Zhongru Gu<sup>1,3</sup>, and Xiangjiang Zhan<sup>1,2,3,4</sup>

<sup>1</sup> Key Laboratory of Animal Ecology and Conservation Biology, Institute of Zoology, Chinese Academy of Sciences, Beijing, China <sup>2</sup> University of Chinese Academy of Sciences, Beijing, China <sup>3</sup> Cardiff University-Institute of Zoology Joint Laboratory for Biocomplexity Research, Chinese Academy of Sciences, Beijing, China <sup>4</sup> Center for Excellence in Animal Evolution and Genetics, Chinese Academy of Sciences, Kunming, China

DOI: [10.xxxxxx/draft](https://doi.org/10.xxxxxx/draft)

## Software

- [Review](#) ↗
- [Repository](#) ↗
- [Archive](#) ↗

Editor: [Open Journals](#) ↗

## Reviewers:

- [@openjournals](#)

Submitted: 01 January 1970

Published: unpublished

## License

Authors of papers retain copyright and release the work under a Creative Commons Attribution 4.0 International License ([CC BY 4.0](#))

## Summary

Stemflow is a user-friendly Python package for Adaptive Spatio-Temporal Exploratory Model (AdaSTEM ([Fink et al., 2013](#))). AdaSTEM is a modeling framework that adopts “split-apply-combine” methodology ([Wickham, 2011](#)) – it adaptively splits data into spatiotemporal grids, train models for each grid, and combines the models for ensemble prediction. Models in stemflow follow the style of scikit-learn BaseEstimator class ([Pedregosa et al., 2011](#)). It provides one-line model creation, fitting, prediction, and evaluation. It implements spatio-temporal train-test-split and cross-validation functions. After model training, feature importance could be evaluated with spatio-temporal dynamics. Stemflow also provides functions for visualizing ensembles structured in model training and generating GIF file for predicted results to animate the spatio-temporal movement of animal population.

## Statement of need

Spatio-temporal big data is an increasingly valuable resource for modern ecological studies ([Farley et al., 2018](#)). A large amount of spatio-temporal big data is now derived from broad-scale surveys, such as citizen science projects ([Dickinson et al., 2010](#)). The intensity of survey activities grows rapidly as more people are involved in citizen science in recent years, resulted in exponential accumulation of observational data ([Di Cecco et al., 2021](#); [Sullivan et al., 2014](#)). However, daily species observation records uploaded by non-professionals in citizen science program are known to have larger bias than professionally structured research, both in terms of data veracity and spatio-temporal balance of the datasets, which necessitates elaborate modeling methods to provide insights ([Dickinson et al., 2010](#); [Farley et al., 2018](#)).

Some species distribution modeling (SDM) approaches were brought forward to adjust for bias in citizen science and model on the unobserved components ([Bird et al., 2014](#)). Still, many failed to account for the autocorrelation of space and time ([F. Dormann et al., 2007](#)), which is especially crucial in modeling inherently spatio-temporal biological events with variations at different scales ([Chave, 2013](#); [Levin, 1992](#)), such as seasonal migration. Adaptive Spatio-Temporal Exploratory Model (AdaSTEM) is a semi-parameterized machine learning model that leverages the spatio-temporal adjacency information of sample points to model occurrence or abundance of species ([Fink et al., 2013](#)). A QuadTree algorithm ([Samet, 1984](#)) is implemented to split data into smaller spatio-temporal grids (called stixels) conditional on the data abundance, with more abundant data allowing stixels to be divided into finer resolution (up to a maximum). Stixels with sample size less than a certain threshold will not be modeled; instead, these stixels will be labeled as unpredictable. This procedure controls the degree of model extrapolation (known as “long-distance prediction” problem in spatial settings) and reduces overfitting. A

base model is trained for each stixel, that is, targets are only modeled on their adjacent information in space and time. Splitting-training is carried out several times to generate multiple ensembles. Finally, prediction results were aggregated across these ensembles.

AdaSTEM shows the capacity of supporting large scale spatio-temporal ecological data modeling in many studies (Fink et al., 2020; Fuentes et al., 2023; La Sorte et al., 2022), especially for modeling animal abundance at different scales (Fink et al., 2013). One well-known application of AdaSTEM is the weekly abundance map of eBird Status and Trend product (Fink et al., 2022), which was widely used as data sources of abundance data of bird populations (Bird et al., 2014; Jarzyna & Stagge, 2023; Lin et al., 2022). The application of AdaSTEM could be extended to other fields with similar data structure and spatio-temporal dependence, for example, epidemiology. Despite the foreseeable significant role of spatio-temporal big data in the coming decades of scientific research, the development of tools has not necessarily kept pace.

Stemflow is positioned as a user-friendly Python package to meet the need of general application of modeling spatio-temporal large datasets. Scikit-learn style object-oriented modeling pipeline enables concise model construction with compact parameterization at the user end, while the rest of the modeling procedures are carried out under the hood. Once the fitting method is called, the model class recursively splits the input training data into smaller spatio-temporal stixels using QuadTree algorithm. For each of the stixels, a base model is trained only using data falls into that stixel. Stixels are then aggregated and constitute an ensemble. In the prediction phase, stemflow queries stixels for the input data according to their spatial and temporal index, followed by corresponding base model prediction. Finally, prediction results are aggregated across ensembles to generate robust estimations (see Fink et al. (2013) and stemflow documentation for details).

For survey projects that include abundance information like eBird (Sullivan et al., 2014), the targeted modeling values are often zero-inflated, owing to the fact of low observation probability in many species. Zero-inflation could lead to poor regression model performance (Campbell, 2021). In stemflow, we implement hurdle model classes that embed two sequential models: a classifier to classify the absence and presence states, followed by a regressor to model the abundance for prediction samples classified as presence. Hurdle model classes can be conjunctively used with AdaSTEM model classes in two ways: Use hurdle model as the base model for AdaSTEMRegressor (as in Johnston et al. (2015)), or use AdaSTEMClassifier and AdaSTEMRegressor as the classifier and regressor in hurdle model. We demonstrate the comparison of these two architectures in stemflow documentation.

One advantage of applying stemflow in scikit-learn style is that there is a variety of “base models” to choose from scikit-learn or scikit-learn-style repertoire. The choices vary from linear models to boosting and bagging tree-based models. Maxent model (C. B. Anderson, 2023) is also supported to play the role of “base model”, which largely expands the potential application for presence-only modeling (see documentation).

While there exists many open source packages for species distribution modeling (mostly in R, (Norberg et al., 2019); and currently one in Python (C. B. Anderson, 2023)), most of them solely leverage environmental variables and do not support integration of spatio-temporal information during model construction (but see C. B. Anderson (2023); S. C. Anderson et al. (2022); Dobson et al. (2023)). This disadvantage is usually noted along with the overconfidence of the model extrapolation capacity both for Maxent-based and ensemble-based models (A. Lee-Yaw et al., 2022). To our knowledge, stemflow is the first SDM package specifically crafted to address spatio-temporal dependencies in samples while also accounting for biases in sample distribution. With the rapid accumulation of data and development of machine learning techniques, stemflow will exhibit greater advantages in spatio-temporal modeling, and could be applied to other fields (e.g., epidemiology and weather prediction) in future.

## Acknowledgements

This project was based upon work supported by National Natural Science Foundation of China grants 32125005 (to X.Z.), 32270455 (to Z.G.), CAS Project for Young Scientists in Basic Research (YSBR-097 to X.Z. and Z.G.). We thank Dr. Daniel Fink at Cornell Lab of Ornithology for the R scripts of STEM model and QuadTree which inspired our implementation in Python.

## References

- A. Lee-Yaw, J., L. McCune, J., Pironon, S., & N. Sheth, S. (2022). Species distribution models rarely predict the biology of real populations. *Ecography*, 2022(6), e05877. <https://doi.org/10.1111/ecog.05877>
- Anderson, C. B. (2023). Elapid: Species distribution modeling tools for Python. *Journal of Open Source Software*, 8(84), 4930. <https://doi.org/10.21105/joss.04930>
- Anderson, S. C., Ward, E. J., English, P. A., & Barnett, L. A. K. (2022). *sdmTMB: An R package for fast, flexible, and user-friendly generalized linear mixed effects models with spatial and spatiotemporal random fields* [Preprint]. *Ecology*. <https://doi.org/10.1101/2022.03.24.485545>
- Bird, T. J., Bates, A. E., Lefcheck, J. S., Hill, N. A., Thomson, R. J., Edgar, G. J., Stuart-Smith, R. D., Wotherspoon, S., Krkosek, M., Stuart-Smith, J. F., Pecl, G. T., Barrett, N., & Frusher, S. (2014). Statistical solutions for error and bias in global citizen science datasets. *Biological Conservation*, 173, 144–154. <https://doi.org/10.1016/j.biocon.2013.07.037>
- Campbell, H. (2021). The consequences of checking for zero-inflation and overdispersion in the analysis of count data. *Methods in Ecology and Evolution*, 12(4), 665–680. <https://doi.org/10.1111/2041-210X.13559>
- Chave, J. (2013). The problem of pattern and scale in ecology: What have we learned in 20 years? *Ecology Letters*, 16(s1), 4–16. <https://doi.org/10.1111/ele.12048>
- Di Cecco, G. J., Barve, V., Belitz, M. W., Stucky, B. J., Guralnick, R. P., & Hurlbert, A. H. (2021). Observing the Observers: How Participants Contribute Data to iNaturalist and Implications for Biodiversity Science. *BioScience*, 71(11), 1179–1188. <https://doi.org/10.1093/biosci/biab093>
- Dickinson, J. L., Zuckerberg, B., & Bonter, D. N. (2010). Citizen Science as an Ecological Research Tool: Challenges and Benefits. *Annual Review of Ecology, Evolution, and Systematics*, 41(1), 149–172. <https://doi.org/10.1146/annurev-ecolsys-102209-144636>
- Dobson, R., Challinor, A. J., Cheke, R. A., Jennings, S., Willis, S. G., & Dallimer, M. (2023). dynamicSDM: An r package for species geographical distribution and abundance modelling at high spatiotemporal resolution. *Methods in Ecology and Evolution*, 14(5), 1190–1199. <https://doi.org/10.1111/2041-210X.14101>
- F. Dormann, C., M. McPherson, J., B. Araújo, M., Bivand, R., Bolliger, J., Carl, G., G. Davies, R., Hirzel, A., Jetz, W., Daniel Kissling, W., Kühn, I., Ohlemüller, R., R. Peres-Neto, P., Reineking, B., Schröder, B., M. Schurr, F., & Wilson, R. (2007). Methods to account for spatial autocorrelation in the analysis of species distributional data: A review. *Ecography*, 30(5), 609–628. <https://doi.org/10.1111/j.2007.0906-7590.05171.x>
- Farley, S. S., Dawson, A., Goring, S. J., & Williams, J. W. (2018). Situating Ecology as a Big-Data Science: Current Advances, Challenges, and Solutions. *BioScience*, 68(8), 563–576. <https://doi.org/10.1093/biosci/biy068>

- 137 Fink, D., Auer, T., Johnston, A., Ruiz-Gutierrez, V., Hochachka, W. M., & Kelling, S. (2020).  
 138 Modeling avian full annual cycle distribution and population trends with citizen science  
 139 data. *Ecological Applications*, 30(3), e02056. <https://doi.org/10.1002/eap.2056>
- 140 Fink, D., Auer, T., Johnston, A., Strimas-Mackey, M., Ligocki, S., Robinson, O., Hochachka,  
 141 W., L., Jaromczyk, Rodewald, A., Wood, C., Davies, I., & Spencer., A. (2022). *eBird status*  
 142 *and trends, data version: 2021; released: 2022*. Cornell Lab of Ornithology, Ithaca, New  
 143 York. <https://doi.org/10.2173/ebirdst.2021>
- 144 Fink, D., Damoulas, T., & Dave, J. (2013). Adaptive Spatio-Temporal Exploratory Models:  
 145 Hemisphere-wide species distributions from massively crowdsourced eBird data. *Proceedings*  
 146 *of the AAAI Conference on Artificial Intelligence*, 27(1), 1284–1290. <https://doi.org/10.1609/aaai.v27i1.8484>
- 148 Fuentes, M., Van Doren, B. M., Fink, D., & Sheldon, D. (2023). BirdFlow: Learning seasonal  
 149 bird movements from eBird data. *Methods in Ecology and Evolution*, 14(3), 923–938.  
 150 <https://doi.org/10.1111/2041-210X.14052>
- 151 Jarzyna, M. A., & Stagge, J. H. (2023). Decoupled spatiotemporal patterns of avian taxonomic  
 152 and functional diversity. *Current Biology*, 33(6), 1153–1161.e4. <https://doi.org/10.1016/j.cub.2023.01.066>
- 154 Johnston, A., Fink, D., Reynolds, M. D., Hochachka, W. M., Sullivan, B. L., Bruns, N.  
 155 E., Hallstein, E., Merrifield, M. S., Matsumoto, S., & Kelling, S. (2015). Abundance  
 156 models improve spatial and temporal prioritization of conservation resources. *Ecological*  
 157 *Applications*, 25(7), 1749–1756. <https://doi.org/10.1890/14-1826.1>
- 158 La Sorte, F. A., Horton, K. G., Johnston, A., Fink, D., & Auer, T. (2022). Seasonal associations  
 159 with light pollution trends for nocturnally migrating bird populations. *Ecosphere*, 13(3),  
 160 e3994. <https://doi.org/10.1002/ecs2.3994>
- 161 Levin, S. A. (1992). The Problem of Pattern and Scale in Ecology: The Robert H. MacArthur  
 162 Award Lecture. *Ecology*, 73(6), 1943–1967. <https://doi.org/10.2307/1941447>
- 163 Lin, H.-Y., Binley, A. D., Schuster, R., Rodewald, A. D., Buxton, R., & Bennett, J. R. (2022).  
 164 Using community science data to help identify threatened species occurrences outside of  
 165 known ranges. *Biological Conservation*, 268, 109523. <https://doi.org/10.1016/j.biocon.2022.109523>
- 167 Norberg, A., Abrego, N., Blanchet, F. G., Adler, F. R., Anderson, B. J., Anttila, J., Araújo,  
 168 M. B., Dallas, T., Dunson, D., Elith, J., Foster, S. D., Fox, R., Franklin, J., Godsoe,  
 169 W., Guisan, A., O'Hara, B., Hill, N. A., Holt, R. D., Hui, F. K. C., ... Ovaskainen, O.  
 170 (2019). A comprehensive evaluation of predictive performance of 33 species distribution  
 171 models at species and community levels. *Ecological Monographs*, 89(3), e01370. <https://doi.org/10.1002/ecm.1370>
- 173 Pedregosa, F., Varoquaux, G., Gramfort, A., Michel, V., Thirion, B., Grisel, O., Blondel,  
 174 M., Prettenhofer, P., Weiss, R., Dubourg, V., Vanderplas, J., Passos, A., Cournapeau,  
 175 D., Brucher, M., Perrot, M., & Duchesnay, Édouard. (2011). Scikit-learn: Machine  
 176 Learning in Python. *Journal of Machine Learning Research*, 12(85), 2825–2830. <http://jmlr.org/papers/v12/pedregosa11a.html>
- 178 Samet, H. (1984). The Quadtree and Related Hierarchical Data Structures. *ACM Computing*  
 179 *Surveys*, 16(2), 187–260. <https://doi.org/10.1145/356924.356930>
- 180 Sullivan, B. L., Aycrigg, J. L., Barry, J. H., Bonney, R. E., Bruns, N., Cooper, C. B.,  
 181 Damoulas, T., Dhondt, A. A., Dietterich, T., Farnsworth, A., Fink, D., Fitzpatrick, J.  
 182 W., Fredericks, T., Gerbracht, J., Gomes, C., Hochachka, W. M., Iliff, M. J., Lagoze,  
 183 C., La Sorte, F. A., ... Kelling, S. (2014). The eBird enterprise: An integrated approach  
 184 to development and application of citizen science. *Biological Conservation*, 169, 31–40.  
 185 <https://doi.org/10.1016/j.biocon.2013.11.003>

<sup>186</sup> Wickham, H. (2011). The split-apply-combine strategy for data analysis. *Journal of Statistical*  
<sup>187</sup> *Software*, 40, 1–29. <https://doi.org/10.18637/jss.v040.i01>

DRAFT