Sambit Panda

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SUMMARY

- Highly motivated professional with 10+ years of research experience; interests include machine learning, data science, statistics, cancer genomics, and neuroscience
- Author of 14 publications (h-index: 6, ~200 citations); see all at https://sampan.me/pdf/Sambit-Panda-CV.pdf
- 6+ years of experience using Python and R to develop data science solutions in academic and industry settings

SKILLS

Python (pandas, scikit-learn, TensorFlow, PyTorch, etc.), R, Cython, Cloud Services (AWS, Azure), Databases (SQL), Developer Tools (Git, Docker), Continuous Integration (CircleCI, Travis CI, etc.) MATLAB, Unix Shell Scripts, Familiarity with HTML/CSS, C/C++, Java, Generative AI, ML-Ops

RELEVENT EXPERIENCE

NeuroData Lab, Johns Hopkins   Jan 2019 – Present
Researcher   *Baltimore, MD*
- Developed multiple algorithms, notably KMERF (random forest-based hypothesis test), Nonparametric MANOVA (a nonparametric multivariate k-sample test), Fast Dcorr (fast approximation to the distance correlation test), and Causal Dcorr (distance correlation for causal inference)
- Authored 11 publications (5 first author, ~150 citations) related to early cancer detection, random forest, neural networks, causal inference, and hypothesis testing using **Python** packages like **TensorFlow**, **PyTorch**, etc.
- Created and maintained open-source **Python** packages like hyppo (~150 users, 200+ stars, ~100 forks) and treeple (50+ stars, ~20 forks); ported algorithms from these packages into SciPy.
- Developed and tested code using **Git**, **Docker**, Cloud Services (**AWS EC2/S3**, **Azure VM**), Continuous Integration (**CircleCI**, **Travis CI**, etc.), and **Python** packages (**pandas**, **scikit-learn**, etc.)
- Collaborated with Bert Vogelstein, a renowned scientist in cancer genomics, on the MIGHT algorithm that quantifies predictive information in liquid biopsy feature sets; used **Python** packages (treeple, **scikit-learn**, **pandas**, etc.); wrote manuscript in preparation for Science
- Served as SciPy symposium conference chair and reviewer; journal reviewer for SoftwareX; presented work at top conferences like the BRAIN PI meeting and GYSS
- Worked on a project annotating whole body CT scans using **Python**, **Unix shell scripts**

National Institutes of Environmental Health Sciences   May 2023 – Jul 2023
Data Scientist   *RTP, NC*
- Applied the KMERF algorithm (which I created) to discover relationships in neurological data using **Python** packages (**pandas**, **scikit-learn**, etc.) and **R**; won 1st place in poster competition
- Collaborated with researchers to publish two manuscripts: (1) neurotransmitter signaling from fear response in mice and (2) the development of a fiber photometry **R** package; developed tutorials interfacing **Python** and **MySQL**

PROJECTS

treeple (originally scikit-tree) | *Python, Cython*   2023 – Present
- Extends scikit-learn decision trees to do oblique splits, manifold learning, hypothesis testing, etc. (50+ stars, ~20 forks).
- Role: Core contributor and maintainer of this package.

hyppo (originally mgcpy) | *Python, Continuous Integration, AWS, Azure*   2018 – Present
- The first Python package for multivariate hypothesis testing, closing the gap with R (~150 users, 200+ stars, ~100 forks).
- Role: Creator and maintainer of this package.

scipy.stats.multiscale_graphcorr | *Python, Cython*   2019 – Present
- Multiscale Graph Correlation is a powerful multivariate test (the 1st and only multivariate test in SciPy).
- Role: Ported this algorithm from hyppo and maintain it.

FiPhA | *R*   2023
- A robust and user-friendly package for fiber photometry analysis.
- Role: Open-sourced this package and helped maintained it.

EDUCATION

Johns Hopkins Medical Institute
Baltimore, MD

PhD, Biomedical Engineering
Jul 2020 – Dec 2024
- Awards: Computational Biology Fellowship (2020)
- Service: A-Level Capital (VC Firm) Life Sciences Advisor, TA (Neurodata Design I & II)

Johns Hopkins University
Baltimore, MD

MSE, Biomedical Engineering
Aug 2018 – May 2020
- Awards: AWS IMAGINE Grant (2018)

NC State University & UNC Chapel Hill
Raleigh & Chapel Hill, NC

BS, Biomedical Engineering & Biology
Aug 2014 – May 2018
- Awards: Magna Cum Laude (2018), Honors Program (2018), Dean's List (2014 – 2018), Goodnight Scholarship (Full Ride, 2014), National Merit Scholarship (2014)

PUBLICATIONS (Highlighting 5 of 14)

1. **Panda, S.**, …, & Vogelstein, J. T. (2024*). hyppo: A Multivariate Hypothesis Testing Python Package*. Manuscript submitted for publication in JMLR.
2. **Panda, S.***, Shen, C.*, & Vogelstein, J. T. (2024). *Learning Interpretable Characteristic Kernels via Decision Forests*. Manuscript in preparation for ICLR 2024.
3. Curtis, S.*, **Panda, S.***, Li, A.*, …, Vogelstein, B., Vogelstein, J. T.^, & Douville, C.^ (2024). *Detecting and Combining Useful Sets of Predictive Variables*. Manuscript in preparation for Science.
4. **Panda, S.***, Shen, C.*, …, & Vogelstein, J. T. (2024). *Universally Consistent K-Sample Tests via Dependence Measures*. In Press in Stat Prob Letters.
5. Shen, C., **Panda, S.**, & Vogelstein, J. T. (2022). The Chi-Square Test of Distance Correlation. *Journal of Computational and Graphical Statistics*, *31*(1), 254–262. https://doi.org/10.1080/10618600.2021.1938585

PRESENTATIONS (Highlighting 3 of 21)

1. **Panda, S.**, …, & Cushman, J. D. (2023, July). *Elucidating Relationships within Neurological Screening Batteries via Random Forest-Based Hypothesis Testing* [Poster Presentation] RTP, NC, USA.
2. **Panda, S.**, …, & Vogelstein, J. T. (2022, January). *Nonparametric MANOVA via Independence Testing* [Oral Presentation]. Global Young Scientists Summit, Virtual. https://www.youtube.com/watch?v=rJyuTwkgfjQ
3. **Panda, S.**, …, & Vogelstein, J. T. (2021, June). *Nonparametric MANOVA via Independence Testing* [Poster Presentation] BRAIN Initiative Meeting, Virtual.