Samson Koelle

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I am a researcher and engineer with diverse skills in artificial intelligence, interpretable machine learning, and computational biology. I am looking for team-based work with real-world impact.

Experience

Uberduck	Cofounder	Seattle, WA	08/2021 - 03/2024		
Headed text-to-speech development for startup at 0-60k MRR					
Allen Institute for Brain Science	Graduate assistant	Seattle, WA	01/2020 - 12/2020		
Statistical methods development for modeling brain connectivity					
Amazon SCOT Forecasting Science	Research Intern	Seattle, WA	06/2017 - 09/2017		
Online training of fine-tuned forecasting models					
Fred Hutchinson CRC	Graduate assistant	Seattle, WA	09/2015 - 06/2017		
Computational analysis of single cell RNA sequencing					
National Heart, Lung, & Blood Inst.	IRTA	Bethesda, MD	07/2013 - 05/2015		
Studied distribution of clonality in hematopoiesis					

Education

Y Combinator	San Francisco, CA	01/2022 - 03/2022		
Ph.D. Statistics	University of Washington, Seattle, WA	09/2015 - 03/2022		
Thesis: Geometric algorithms for interpretable manifold learning.				
Committee: Marina Meila (advisor), Yen-Chi Chen, Zaid Harchaoui, John Lee				
B.A. Mathematics	Columbia University, New York, NY	09/2009 - 05/2013		

References

Zach Ocean	Uberduck, Inc.	z@uberduck.ai
Marina Meila	University of Washington Department of Statistics	mmp@uw.edu
Stefan Mihalas	Allen Institute for Brain Science	stefanm@alleninstitute.org
Cynthia Dunbar	National Heart Lung and Blood Institute	dunbarc@nhlbi.nih.gov

Service

Reviewer	AISTAT	2023
Reviewer	JMLR	2021
Reviewer	UAI	2020-2021
Organizer	UW Geometric Data Analysis reading group (uwgeometry.github.io)	2017-2021

Publications

Koelle, Samson et al. (Dec. 2023). "Modeling the cell-type-specific mesoscale murine connectome with anterograde tracing experiments". en. In: Netw Neurosci 7.4, pp. 1497–1512.

Zhang, Hanyu et al. (Feb. 2023). "Dictionary-based Manifold Learning". In: AISTATS 2024 (to appear). arXiv: 2302.00263 [cs.LG].

Koelle, Samson (2022). "Geometric Algorithms for Interpretable Manifold Learning". en. PhD thesis. Ann Arbor, United States: University of Washington.

Koelle, Samson J et al. (2022). "Manifold Coordinates with Physical Meaning". In: *J. Mach. Learn. Res.* 23.133, pp. 1–57.

- Buenfil, James et al. (2021). "Tangent Space Least Adaptive Clustering". en. In: 2021 ICML Unsupervised Learning for Reinforcement Learning Workshop.
- Espinoza, Diego A et al. (Apr. 2021). "Interrogation of clonal tracking data using barcodetrackR". en. In: Nat Comput Sci 1.4, pp. 280–289.
- Xu, Jason et al. (2019). "Statistical inference for partially observed branching processes with application to cell lineage tracking of in vivo hematopoiesis". In: Ann. Appl. Stat. 13.4, pp. 2091–2119.
- Meila, Marina et al. (Nov. 2018). "A regression approach for explaining manifold embedding coordinates".
- Paulson, K G et al. (Sept. 2018). "Acquired cancer resistance to combination immunotherapy from transcriptional loss of class I HLA". en. In: *Nat. Commun.* 9.1, p. 3868.
- Wu, Chuanfeng et al. (Nov. 2018a). "Clonal expansion and compartmentalized maintenance of rhesus macaque NK cell subsets". en. In: Sci Immunol 3.29.
- Wu, Chuanfeng et al. (Jan. 2018b). "Geographic clonal tracking in macaques provides insights into HSPC migration and differentiation". en. In: *J. Exp. Med.* 215.1, pp. 217–232.
- Yabe, Idalia M et al. (Dec. 2018). "Barcoding of Macaque Hematopoietic Stem and Progenitor Cells: A Robust Platform to Assess Vector Genotoxicity". en. In: *Mol Ther Methods Clin Dev* 11, pp. 143–154.
- Yu, Kyung-Rok et al. (Mar. 2018). "The impact of aging on primate hematopoiesis as interrogated by clonal tracking". en. In: *Blood* 131.11, pp. 1195–1205.
- Koelle, Samson J et al. (Mar. 2017). "Quantitative stability of hematopoietic stem and progenitor cell clonal output in rhesus macaques receiving transplants". en. In: *Blood* 129.11, pp. 1448–1457.
- Wu, Chuanfeng et al. (Apr. 2014). "Clonal tracking of rhesus macaque hematopoiesis highlights a distinct lineage origin for natural killer cells". In: Cell Stem Cell 14.4, pp. 486–499.