Computational Biologist ~ Software Engineer		Denver, Co	olorado 🔽	jdkrol351@gmail.com
			in	/in/jacob-krol-b3b784156
SUMMARY —		SKILLS —		
A scientific programmer specializing in machine learning (ML) with significant experience in front and back-end web development, High Performance Computing (HPC), and devops.		Languages:	R, Python, B Perl, C, Java	ash/Shell, JavaScript, HTML,
		Technologies	(HPC), PBS numpy, tidy	, PyTorch, Scikit-learn, Slurm 5 Torque (HPC), tidymodels, vverse, matplotlib, ggplot2, R er, Singularity, LaTeX, SQL.
PROJECTS —				
R,Shiny,Docker, Slurm,Bash	MolEvolvR: web-app for protein character Frontend, backend, and devops for the app			**
R,Docker	amR: R package for bacterial drug-resistance classification (ML) models R data files for drug resistance classification models, feature engineered omics datasets (training data) with supporting classes and functions. Publication in-prep.			
Python,Docker,CLI	Aibou: a python package for a turn-based, quick-time-event, CLI game A monster battle game a CLI user-interface and AI to play against. Quick time key press events are good dexterity training! Hosted on PyPI; installable via pip.			
R,Shiny	Company website for Krol Building			jakekrol/kRol_building
	Image showcase, mission statements, and on Deployed to shinyapps.io free hosting platform		n for Krol Bui	lding Company.
EDUCATION —				
9/2023 - 12/2023	Graduate (non-degree seeking) 4.0/4.0 BIOS 7747: Machine Learning for Biomedii School of Public Health (funded by employe			F Colorado Anschutz Medical Campus urse offered by the Colorado
9/2020 - 8/2022	Bachelor of Science, Neuroscience 3.89/4.0 Michigan State University Undergraduate research experience. Notable coursework: neural engineering, cognitive science, linguistics, genetics, statistics			
9/2017 - 5/2020	Math and Science transfer program 3.52/4 Notable coursework: Java programming, ca			Washtenaw Community College
11/2023 - present	Mathematics for Machine Learning Special Three course specialization: linear algebra, r Free registration as CU Anschutz employee	multivariate calcul		Coursera ple component analysis (PCA).
EXPERIENCE -				
•	Information Sciences Professional Machine learning prediction and statistical in Extensive hyperparameter tuning, model per classification models for drug classes. Developed and containerized a feature enging genomic sequences and executed on HPC to R / Python / Bash / Docker /	nference of drug- rformance evalua neering pipeline (a to generate trainin	resistance ac tion, and mod innotation, clu ig data from t	el selection to build ensemble ustering, and pangenomics) for housands of bacterial isolates.
•	Lorem ipsum dolor sit amet, consectetuer Developed machine learning models to pre Feature engineering omics datasets and state ponent Analysis Python / Scikit-learn / Pandas /	dict plant virus ho tistical inference o	f features: Fis	
MACHINE LEARN	IING EXPERIENCE	•		

convolutional neural networks, artificial neural networks, random forests, logistic regression, gradient boosting machines, adaboost, k-nearest-neighbors, OPTICS, DBSCAN, hierarchical clustering, k-means, linear/non-linear regression

github.com/jakekrol

Lab website

JACOB D KROL