Robert Lesurf

Senior Bioinformatician

Personal Information

Address: Available upon request

Phone: +1 647-679-8446

Email: robert.lesurf@gmail.com

Website: lesurf.org

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l'échnical Skills		
Bioinformatics		Developed machine learning pipeline to increase accuracy of diagnostic and prognactic biomerkors in prostate capacity
Genomics		and prognostic biomarkers in prostate cancer.
Data Analysis		 Led and co-analyzed several other cancer genomics research projects.
Machine Learning	Sep 2014 -	Postdoctoral Research Associate
Statistical Modeling	Sep 2016	McDonnell Genome Institute, Washington University, St. Louis, MO, USA
e e		 Led genomics analysis for clinical trial of breast cancer, computationally
Data Visualization		identified genomic and transcriptional features predictive of drug response.
Cluster Computing		• Designed a 'regulome' capture targets in partnership with Roche.
Version Control		Built data visualization functions for the GenVisR R package.
Fluent in English & French		
		 Mentored students and junior employees.

Senior Bioinformatician

Experience

Jan 2019 -

Soft Skills

2000-2014	FILD MICGIII OIIIVEISILY, MOIILIEAI, QC, Callaua
	Biochemistry (Bioinformatics option)
	Used machine learning and microarray data to identify and predict early
	stage breast cancer patients who may be safely spared therapy.
	• Developed visualization algorithms for genomic signatures across tumours.
2006-2008	M.Sc McGill University, Montreal, QC, Canada
2000-2008	• • • •
	Computer Science (Bioinformatics option)
	Identified genomic features of mouse models for human cancer.
2002 2006	R Sc. Honours, Queen's University Kingston ON Canada
	2008-2014

Programming Languages

••	
Python	
Perl	
HTML	
Java	
SQL	
Unix	

R

Organized professional with over a decade of research experience in genomics, bioinformatics, and machine learning. Distinguished leadership resulting in the completion and publication of eighteen peer-reviewed scientific studies. I have a passion for data analysis, visualization, problem solving, and summarizing results to broad audiences.

Jan 2019 -	
Present	The Hospital for Sick Children (SickKids), Toronto, ON, Canada
	 Leading internal and international bioinformatics team collaborations.
	 Supervising students and their research projects.
	 Advising hospital migration to cloud compute environments.
	• Identifying genomic variants that cause heart disease in children, and the
	influence of these variants on disease severity and outcome.
Sep 2016 -	Bioinformatician, Data Scientist
Jan 2019	Ontario Institute for Cancer Research, Toronto, ON, Canada
	• Led team development for genomics data analysis pipeline, bringing
	software tools into a unified framework for automated quality control and
	analysis of sequencing data.
	• Developed machine learning pipeline to increase accuracy of diagnostic
	and prognostic biomarkers in prostate cancer.
	• Led and co-analyzed several other cancer genomics research projects.
Sep 2014 -	Postdoctoral Research Associate
Sep 2016	McDonnell Genome Institute, Washington University, St. Louis, MO, USA
•	• Led genomics analysis for clinical trial of breast cancer, computationally
	identified genomic and transcriptional features predictive of drug response.
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Education	
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