

René L Warren

warrenlr@gmail.com 778.386.4192 http://renewarren.ca

scientist with 20+ years experience in biotechnology, genomics, bioinformatics

I have played a central role in managing the bioinformatics of large international collaborations to decrypt the genomes of *Rhodococus*, *Cryptococcus*, Bullfrog and Spruce

I developed the first software (SSAKE) for de novo genome assembly with then emerging short DNA sequences and pioneered the development of technologies that enabled the discovery of Fusobacterium in colon cancer, one of Time Magazine's top ten medical breakthrough of 2011

I am the BC Genome Sciences Centre's bioinformatics technology lab <u>Group Leader</u> In that role, I have conceptualized & led the development of genome analysis technologies I mentored over half a dozen undergraduate students, their work has led to first authorship

seeking additional challenges and leadership

PROFESSIONAL EXPERIENCE

Group Leader

2017 – present

BC Cancer Agency - Genome Sciences Centre, Vancouver

- Project leadership, expertise, guidance
- Coordinate research activities
- Interview, supervise, mentor COOP students and staff
- Conceptualize development of bioinformatics technologies
- Write research proposals and scientific articles

Coordinator

2002 -

BC Cancer Agency - Genome Sciences Centre, Vancouver

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- Lead bioinformatics software R&D (Python, PERL, R, unix)
- Published research (scientific journals, international conferences)
- Developed marketing (web portals, news release)
- Supervised a team of biologists and programmers
- Interviewed job candidates, taught and trained employees

Officer

2000 -

NRC - CNRC - Biotechnology Research Institute, Montréal

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- Engineered gene expression regulation technology (molec/cell biology)
- Designed, fabricated, tested components of the DNA "gene switch"
- Work led to a patent, technology sold to company

EDUCATION

2000 – 01	Certificate Computer Science	Concordia University
1997 – 99	MSc Biochemistry & Molecular Biology	UBC
1994 – 97	BSc Biochemistry Dean's Honours List	Université de Montréal
ADDITIONAL INFORMATION		
2015, 16	Recipient of the John Jambor Knowledge Fund travel award	
2011	Interviewed by NTN24 channel for Fusobacterium discovery colon cancer	
2009	Interviewed by Genome Technology to discuss next-generation sequencing	
2007	Interviewed by GenomeWeb for the development of SSAKE	
1998	UBC Graduate Fellowship awarded for MSc	
1997	Fonds de la Recherche en Santé Québec (FRSQ) awarded for BSc	
1996	Bursary from FRSQ for BSc honour's research project	
1995	Worked at NASA to coordinate the crystallization of proteins under microgravity: CMIX-4 payload, space shuttle <i>Endeavour</i>	
	PRESENTATIONS	
2017, 18	(selected from 16) Research in Computational Molecular Biology, Hong Kong / Paris – talks	
2017, 16	Intelligent Systems for Molecular Biology, Dublin UK / Orlando USA– talks	
2013, 10	Pacific Symposium on Biocomputing, Kona, Hawaii USA — posters	
2010	Sequencing, Finishing and Analysis in the Future, Santa Fe USA — talk	
2007	Synthetic Biology 3.0 conference, Zürich, Switzerland – talk	

PUBLICATIONS

(selected from 58 *co-first authors)

Warren RL. (2018) Visualizing genome synteny with xmatchview. *Journal of Open Source Software*. 3:497

Warren RL, *et al.* (2015) LINKS: Scalable, alignment-free scaffolding of draft genomes with long reads. *GigaScience* 4:35

Warren RL, et al. (2012) Derivation of HLA types from shotgun sequence datasets. Genome Med. 4:95 Castellarin M*, Warren RL*, et al. (2012) Fusobacterium nucleatum infection is prevalent in human colorectal carcinoma. Genome Research. 22:299-306

Warren RL, et al. (2007) Assembling millions of short DNA sequences using SSAKE. Bioinformatics. 23:500

E Allen-Vercoe, R Holt, R Moore, **R Warren.** Detection of fusobacterium in a gastrointestinal sample to diagnose gastrointestinal cancer. US Patent App. 13/877,421 / WO Patent 2,012,045,150

REFERENCES

Available upon request