

# René L Warren

warrenlr@gmail.com 778 . 386 . 4192 http://renewarren.ca

I'm a scientist with 20+ yrs experience in biotechnology, genomics, informatics who:

Developed the first software for *de novo* genome assembly with emerging short DNA sequences

Discovered *Fusobacterium* in colon cancer, one of *Time Magazine*'s 2011 top ten breakthrough

Coordinated the bioinformatics analysis of Rhodococus, Cryptococcus, Bullfrog, Spruce genomes

I am seeking new challenges and additional leadership

#### PROFESSIONAL EXPERIENCE

2017 - now

# **Group Leader**

BC Cancer Agency - Genome Sciences Centre, Vancouver

- Lead group, provide project expertise & guidance
- Conceptualize / lead development of genome analysis technologies
- Interview, supervise, mentor COOP students and staff

2002 - 17

## Coordinator

BC Cancer Agency – Genome Sciences Centre, Vancouver

- Lead bioinformatics software R&D (Python, PERL, R, unix)
- Published research (scientific journals, international conferences)
- Interviewed, taught, trained, supervised biologists and programmers

2000 - 01

#### Officer

NRC - CNRC - Biotechnology Research Institute, Montréal

- Engineered gene expression regulation technology (molec/cell biology)
- Designed, fabricated, tested components of DNA "gene switch"
- Collaborated with stakeholders, chemists, molecular and cell biologists

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# **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
2015, 16 2011 2009 2007 1998 1997 1996 1995	ADDITIONAL INFORMATION Recipient of the John Jambor Knowledge Fund travel award Interviewed by NTN24 channel for Fusobacterium discovery colon cancer Interviewed by Genome Technology to discuss next-generation sequencing Interviewed by GenomeWeb for the development of SSAKE UBC Graduate Fellowship awarded for MSc Fonds de la Recherche en Santé Québec (FRSQ) awarded for BSc Bursary from FRSQ for BSc honour's research project Worked at NASA to coordinate the crystallization of proteins under microgravity: CMIX-4 payload, space shuttle Endeavour	
2017, 18 2015, 16 2008, 12, 15 2010	PRESENTATIONS  (Selected from 16)  Research in Computational Molecular Biology, Hardligent Systems for Molecular Biology, Dublin Pacific Symposium on Biocomputing, Kona, Haw Sequencing, Finishing and Analysis in the Future	n UK / Orlando USA <b>talks</b> vaii USA <b>posters</b>

## **PUBLICATIONS**

Synthetic Biology 3.0 conference, Zürich, Switzerland

(Selected from 58 peer-reviewed \*co-first authors)

Warren RL. 2018. Visualizing genome synteny with xmatchview. J. 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. Bioinfo. 23:500

#### REFERENCES

Available upon request

talk

2007