Curriculum Vitae Daehwan Kim

Address: 1900 E. Monument St.

Welch Medical Library, Rm 101

Baltimore, MD 21205

Email: infphilo@gmail.com

<u>Home Page:</u> http://www.ccb.jhu.edu/people/infphilo<u>Citizenship:</u> South Korea (US Permanent Resident)

Professional Experience:

• Johns Hopkins University School of Medicine

June 2013 - Present

Post-doctoral Research Fellow in McKusick-Nathans Institute of Genetic Medicine Center for Computational Biology led by Dr. Steven L. Salzberg

• Johns Hopkins University School of Medicine

Sep 2011 - May 2013

Trainee in McKusick-Nathans Institute of Genetic Medicine Center for Computational Biology led by Dr. Steven L. Salzberg

• University of Maryland, College Park

June 2010 - May 2013

Graduate Research Assistant in Dr. Steven L. Salzberg's lab

- Working on TopHat, Cufflinks, and TopHat-Fusion

• University of Maryland, College Park

Sep 2008 - May 2010

Graduate Teaching Assistant (see homepage for more details)

• Nexon DD, Seoul, South Korea

Oct 2003 - Jul 2008

(a game development company)
Chief Technology Officer & Manager

- Led several projects as team manager and main developer
- Responsible for hiring and managing developers
- Set and maintained software development process for quality and accuracy of projects

Education:

• University of Maryland, College Park

Sep 2008 - May 2013

Ph.D. in Computer Science (GPA: 4.0/4.0) Supervised under Dr. Steven L. Salzberg • Chung-Ang University, Seoul, South Korea B.Eng. in Computer Science and Engineering Summa cum laude

Journal Publications:

- HISAT-genotype: practical approach for analyzing human variation on a personal computer <u>Daehwan Kim</u> and Steven L. Salzberg. *in preparation*.
- Centrifuge: rapid and accurate classification of metagenomic sequences.

 Daehwan Kim, Li Song, Florian Breitwieser, and Steven L. Salzberg. *in preparation*.
- OperonDB: comprehensive database of operons in 2759 bacterial and archaeal genomes. <u>Daehwan Kim</u>, Corina Antonescu, Mihaela Pertea, and Steven L. Salzberg. *in preparation*.
- The novel fusion transcript NR5A2-KLHL29 is generated by an insertion at the KLHL29 locus Zhenguo Sun, Steven L. Salzberg, <u>Daehwan Kim</u>, Valentin Antonescu, Yulan Cheng, Jee Hoon Song, Xiquan Ke, Binbin Huang, John M. Abraham, Sariat Ibrahim, and Stephen J. Meltzer *submitted*.
- Transcript-level expression analysis of RNA-seq experiments with HISAT, StringTie, and Ballgown Mihaela Pertea, <u>Daehwan Kim</u>, Jack Minyang Fu, Geo Pertea, Jeffrey T. Leek, and Steven L. Salzberg. *To appear in Nature Protocols* soon.
- HISAT: a fast spliced aligner with low memory requirements

 <u>Daehwan Kim</u>, Ben Langmead, and Steven L. Salzberg. *Nature Methods* 12, 357-360 (2015).
- Reconstruction and Estimation of Fusion Transcripts from RNA-Sequencing reads.
 <u>Daehwan Kim</u> and Steven L. Salzberg. *in preparation*.
- TopHat2: accurate alignment of transcriptomes in the presence of insertions, deletions and gene fusions.
 - <u>Daehwan Kim</u>, Geo Pertea, Cole Trapnell, Harold Pimentel, Ryan Kelley, and Steven L. Salzberg. *Genome Biology* 2013, 14:R36.
- Differential Gene and Transcript Expression Analysis of RNA-Seq Experiments with TopHat and Cufflinks.
 - Cole Trapnell, Adam Roberts, Loyal Goff, Geo Pertea, <u>Daehwan Kim</u>, David R. Kelley, Harold Pimentel, Steven L. Salzberg, John L. Rinn, and Lior Pachter. *Nature Protocols* 7, 562578 (2012).
- TopHat-Fusion: an algorithm for discovery of novel fusion transcripts.

 <u>Daehwan Kim</u> and Steven L. Salzberg. *Genome Biology* 2011, 12:R72.

Invited talks:

HISAT-genotype: practical approach for analyzing human variation on a personal computer.
 Korea Institute of Science and Technology (KIST) (May 2016)
 Mini-Symposium of Informatics, University of Colorado, Denver (Apr. 2016)

• Graph-based alignment of next-generation sequencing reads to a population of human genomes Broad Institute (Dec. 2015)

Seven Bridges Genomics (Dec. 2015)

Genome Informatics, Cold Spring Harbor Laboratory (Oct. 2015)

Honors and Awards:

- Best Poster Award Johns Hopkins' Annual Young Investigator Symposium on Genomics and Bioinformatics (Sep 2011)
- Block Grant Fellowship University of Maryland, College Park (2008 2010)
- Employee of the Year for 2005 and 2006 Nexon DD (awarded \$10,000 prize each year)

Academic Background:

- Computer Science (B.Eng. and Ph.D. degrees)
- Mathematics: self-taught most undergraduate mathematics courses (see homepage)
- Some biology and electrical engineering courses: self-taught (see homepage)

Skills:

• Programming Languages and Libraries

C/C++ (>10 years), Visual Basic, Java, Python, Perl, OCaml, ASP, Ruby, Lua STL, MFC, Boost, DirectX, OpenGL, CUDA, Hadoop, ACE (Adaptive Communication Environment), Gamebryo, OGRE, Torque

Tools

Emacs, VIM, gcc/g++/gdb, Make, CVS/SVN, LATEX, Visual Studio, Coq, MySQL, Matlab, Mathematica

• Operating Systems

Excellent working knowledge of Linux, Windows, Mac

• Embedded Systems

ARM9 Embedded Board, ATmega128 MCU, WinAVR Compiler, Ponyprog

References:

Steven L. Salzberg, Ph.D.

Professor

Departments of Biomedical Engineering, Computer Science, and Biostatistics McKusick-Nathans Institute of Genetic Medicine

Johns Hopkins University School of Medicine

Welch Medical Library, 1900 E. Monument St., Rm 107

Baltimore, MD 21205

(410) 614-6112

salzberg@jhu.edu

Ben Langmead, Ph.D.
Assistant Professor
Department of Computer Science
Johns Hopkins University
3400 North Charles St. (329 Malone)
Baltimore, MD 21218-2682
langmea@cs.jhu.edu

Michael Schatz, Ph.D.
Associate Professor
Departments of Computer Science and Biology
Johns Hopkins University
3400 North Charles St. (323 Malone)
Baltimore, MD 21218-2682
mschatz@cs.jhu.edu

Cole Trapnell, Ph.D.
Assistant Professor
Department of Genome Sciences
University of Washington
Foege Building Room S333C
3720 15th Ave NE
Seattle, WA 98105A
(206) 616-6898
coletrap@uw.edu

Mihai Pop, Ph.D.
Associate Professor
Department of Computer Science and Center for Bioinformatics and Computational Biology
University of Maryland, College Park
Biomolecular Sciences Building. Rm. 3120F
College Park, MD 20742
(301) 405-7245
mpop@umiacs.umd.edu