

# Kamil Slowikowski

149 13th Street Charlestown, MA 02129

✉ [kslowikowski@mgh.harvard.edu](mailto:kslowikowski@mgh.harvard.edu)

🔗 [slowkow.com](http://slowkow.com)

## EDUCATION

---

### PhD in Bioinformatics and Integrative Genomics

2012–2019

Harvard University. Boston, Massachusetts.

### BS in Bioinformatics

2007–2011

Loyola University Chicago. Chicago, Illinois.

## GRANT SUPPORT

---

### Predoctoral Individual National Research Service Award (F31-AR070582)

2016–2018

National Institute of Arthritis and Musculoskeletal and Skin Diseases (NIAMS).

Title: *Transcriptomics in synoviocytes defines pathogenesis of rheumatoid arthritis.*

## PROFESSIONAL ACTIVITIES

---

### Postdoctoral Associate. Massachusetts General Hospital.

2019 – Present

[Dr. Chloé Villani](#). Center for Immunology and Inflammatory Diseases.

[Dr. Bo Li](#). Center for Immunology and Inflammatory Diseases.

### Graduate Research Assistant. Harvard Medical School.

2013 – 2019

[Dr. Soumya Raychaudhuri](#). Department of Biomedical Informatics.

[Dr. Michael B. Brenner](#). Division of Rheumatology, Immunology and Allergy, Brigham and Women's Hospital.

### Undergraduate Research Assistant. Loyola University Chicago.

Winter – Summer 2012

[Dr. Sushma Reddy](#). Department of Biology.

[Dr. Catherine Putonti](#). Department of Biology.

### NSF Research Fellow. University of California Santa Cruz.

Summer 2011

[Dr. Todd M. Lowe](#). Department of Biomolecular Engineering.

### Undergraduate Research Assistant. Loyola University Chicago.

2009–2011

[Dr. Howard M. Laten](#). Department of Bioinformatics.

### NSF Research Fellow. The Field Museum. Chicago, Illinois.

Summer 2010

[Dr. Scott Lidgard](#). Integrative Research Center.

### Independent Software Developer. Chicago, Illinois.

2006–2010

Developed computer game extensions for 17 international clients.

## EDUCATIONAL ACTIVITIES

---

### Mentor for Harvard-Amgen Scholars. Harvard University.

2017

Student: Gopal Vashishtha. Harvard College.

Title: *Transcriptional dynamics of synovial fibroblasts in rheumatoid arthritis.*

### Mentor for Research Science Institute (RSI). Harvard University.

2016 – 2017

Student: Sushil Upadhyayula.

Title: *Dissecting the heterogeneity of CD4+ T cell activation with single cell RNA-seq.*

### Instructor. Grades 3-8. The Innovation Institute. Newton, MA.

*Computer Programming for Future Entrepreneurs*

2016 – 2017

*Computer Programming for Future Scientists and Engineers*

2015 – 2016

## SOFTWARE AND TUTORIALS

---

🔗 [Immunogenomics.io](#) (R, HTML, Javascript). Websites for RNA-seq data visualization.

🔗 [ggrepel](#) (R, C++). Repel overlapping text labels away from each other. **Over 3 million downloads.**

🔗 [SNPSEA](#) (C++). Identify cell types and pathways affected by genetic risk loci.

🔗 [snakefiles](#) (Python). Tutorial and Snakefiles for reproducible and scalable RNA-seq data analysis.

🔗 [harmony](#) (Python). Harmony is a data integration algorithm (Korsunsky *et al.* 2019).

## SCHOLARSHIPS AND AWARDS

---

<b>Fifty for the Future Award.</b> Illinois Technology Foundation.	2011
<b>Mulcahy Research Scholar.</b> Loyola University Chicago.	2010–2011
<b>Loyola Presidential Scholarship.</b> Loyola University Chicago.	2007–2011
<b>Dean's List.</b> Loyola University Chicago.	2007–2011

## VOLUNTEER ACTIVITIES

---

<b>Software Developer.</b> <a href="http://cuSTEMized.org">cuSTEMized.org</a> in Boston, MA.	2013–Present
<b>Computer Programming Tutor.</b> Loyola University Chicago (LUC).	2010–2011
<b>Emergency Medical Technician - Basic.</b> LUC Emergency Medical Services.	2008–2010

## TALKS

---

6. Identifying transcriptional regulators central to rheumatoid arthritis: transcriptomics of IL-17 dose-response and time series in stromal cells.
  - *Federation of Clinical Immunology Societies (FOCIS)*. Boston, MA. 2019-06-20
5. Transcriptomics reveals the role of synovial fibroblasts in rheumatoid arthritis.
  - *Leena Peltonen School of Human Genomics*. Les Diablerets, Switzerland. 2018-08-19
4. An introduction to ggrepel.
  - *Partners R User Group Meeting*. Massachusetts General Hospital, Boston, MA. 2018-07-19
3. Insights into RA pathogenesis from single cell transcriptomics of synovial tissue.
  - *Federation of Clinical Immunology Societies (FOCIS)*. Chicago, IL. 2017-06-14
2. AMP rheumatoid arthritis phase 1: single-cell RNA-seq with CEL-seq2.
  - *Accelerating Medicines Partnership (AMP)*. Houston, TX. 2017-02-15
1. Single-cell transcriptomics reveals disease-associated fibroblast subsets in rheumatoid arthritis.
  - *Joint Biology Consortium*. Brigham and Women's Hospital, Boston, MA. 2017-01-27
  - *Medical and Population Genetics Seminar Series*. Broad Institute, Boston, MA. 2016-04-28
  - *Immunogenomics 2015*. HudsonAlpha Institute, Huntsville, AL. 2015-09-30
  - *Cell Circuits and Epigenomics Program Seminar Series*. Broad Institute, Boston, MA. 2015-06-22

## POSTERS

---

6. Introduction to ggrepel.
  - *rstudio::conf2019*. Austin, TX. 2019-01-17
5. Identifying transcriptional regulators central to rheumatoid arthritis: transcriptomics of IL-17 dose-response and time series in stromal cells.
  - *MGH Division of Rheumatology, Allergy, and Immunology Annual Retreat*. North Falmouth, MA. 2019-10-18
  - *Federation of Clinical Immunology Societies (FOCIS) 2019*. Boston, MA. 2019-06-20
  - *Broad Institute Annual Retreat*. Cambridge, MA. 2018-12-17
  - *ISCB Conference on Regulatory and Systems Genomics*. New York, NY. 2018-12-10
  - *American College of Rheumatology (ACR) Annual Meeting*. San Diego, CA. 2017-11-04
  - *Immune Profiling in Health and Disease*. Seattle, WA. 2016-10-03

- |    |   |            |
|----|---|------------|
| 4. | Single-cell transcriptomics identifies pathogenic synovial fibroblasts in rheumatoid arthritis. |            |
|    | • <i>National Human Genome Research Institute (NHGRI) Annual Meeting</i> . Bethesda, MD.        | 2016-04-07 |
|    | • <i>4th Annual Single Cell Analysis Investigators Meeting</i> . Bethesda, MD.                  | 2016-03-02 |
|    | • <i>Harvard Program in Quantitative Genomics Conference</i> . Boston, MA.                      | 2015-11-05 |
| 3. | SNPSEA: an algorithm to identify cell types, tissues, and pathways affected by risk loci.       |            |
|    | • <i>Harvard Graduate Women in Science and Engineering (HGWISE) Symposium</i> . Cambridge, MA.  | 2015-04-25 |
|    | • <i>Harvard Biological and Biomedical Sciences (BBS) Retreat</i> . Provincetown, MA.           | 2014-08-05 |
| 2. | Reverse transcriptase SuperScript III adds non-template bases during cDNA polymerization.       |            |
|    | • <i>Summer Undergraduate Research Symposium</i> . Santa Cruz, CA.                              | 2011-08-12 |
| 1. | Retrotransposon-associated minisatellites in the soybean genome.                                |            |
|    | • <i>Great Lakes Bioinformatics Conference</i> . Athens, OH.                                    | 2011-05-03 |
|    | • <i>Loyola Undergraduate Research Symposium</i> . Chicago, IL.                                 | 2011-04-16 |
|    | • <i>Chicago Area Undergraduate Research Symposium</i> . Chicago, IL.                           | 2011-04-02 |

## SELECTED PEER REVIEWED ARTICLES

15. [Fast, sensitive and accurate integration of single-cell data with Harmony](#)  
Ilya Korsunsky, Nghia Millard, Jean Fan, **Kamil Slowikowski**, Fan Zhang, Kevin Wei, Yuriy Baglaenko, Michael Brenner, Po-ru Loh, Soumya Raychaudhuri  
*Nature Methods*, 2019. [🔗 Learn about Harmony](#)
14. [Tubular Cell and Keratinocyte Single-cell Transcriptomics Applied to Lupus Nephritis Reveal Type I IFN and Fibrosis Relevant Pathways](#)  
Evan Der, Hemant Suryawanshi, Pavel Morozov, Manjunath Kustagi, Beatrice Goilav, Saritha Ranabathou, Peter Izmirly, Michael Belmont, Robert Clancy, Mordecai Koenigsberg, Michele Mokrzycki, Helen Rominieki, Jay Graham, Juan Rocca, Nicole Bornkamp, Nicole Jordan, Emma Schulte, Ming Wu, James Pullman, **Kamil Slowikowski**, Soumya Raychaudhuri, Joel Guthridge, Judith James, Jill Buyon, Thomas Tuschl  
*Nature Immunology*, 2019.
13. [The immune cell landscape in kidneys of patients with lupus nephritis](#)  
Arnon Arazi, Deepak A. Rao, Celine C. Berthier, Anne Davidson, Yanyan Liu, Paul J. Hoover, Adam Chicoine, Thomas M. Eisenhaure, A. Helena Jonsson, Shuqiang Li, David J. Lieb, Fan Zhang, **Kamil Slowikowski**, Edward P. Browne, Akiko Noma, Danielle Sutherby, Scott Steelman, Dawn E. Smilek, Patti Tosta, William Apruzzese, Elena Massarotti, Maria Dall'Era, Meyeon Park, Diane L. Kamen, Richard A. Furie, Fernanda Payan-Schober, William F. Pendergraft III, Elizabeth A. McInnis, Jill P. Buyon, Michelle A. Petri, Chaim Putterman, Kenneth C. Kalunian, E. Steve Woodle, James A. Lederer, David A. Hildeman, Chad Nusbaum, Soumya Raychaudhuri, Matthias Kretzler, Jennifer H. Anolik, Michael B. Brenner, David Wofsy, Nir Hacohen, Betty Diamond the Accelerating Medicines Partnership in SLE network  
*Nature Immunology*, 2019. [🔗 View the data](#)
12. [Lymphocyte innateness defined by transcriptional states reflects a balance between proliferation and effector functions](#)  
Maria Gutierrez-Arcelus, Nikola Teslovich, Alex R. Mola, Rafael B. Polidoro, Aparna Nathan, Hyun Kim, Susan Hannes, **Kamil Slowikowski**, Gerald F. M. Watts, Ilya Korsunsky, Michael B. Brenner, Soumya Raychaudhuri, Patrick J. Brennan  
*Nature Communications*, 2019. [🔗 View the data](#)
11. [Discovering in vivo cytokine-eQTL interactions from a lupus clinical trial](#)  
Emma E. Davenport, Tiffany Amariuta, Maria Gutierrez-Arcelus, **Kamil Slowikowski**, Harm-Jan Westra, Yang Luo, Ciyue Shen, Deepak A. Rao, Ying Zhang, Stephen Pearson, David von Schack, Jean S. Beebe, Nan Bing, Sally John, Michael S. Vincent, Baohong Zhang and Soumya Raychaudhuri  
*Genome Biology*, 2018.
10. [Mixed-effects association of single cells identifies an expanded effector CD4+ T cell subset in rheumatoid arthritis](#)  
Chamith Y. Fonseka\*, Deepak A. Rao\*, Nikola C. Teslovich, Ilya Korsunsky, Susan K. Hannes, **Kamil Slowikowski**, Michael F. Gurish, Laura T. Donlin, James A. Lederer, Michael E. Weinblatt, Elena M. Massarotti, Jonathan S. Coblyn, Simon M. Helfgott, Derrick J. Todd, Vivian P. Bykerk, Elizabeth W. Karlson, Joerg Ermann, Yvonne C. Lee, Michael B. Brenner, and Soumya Raychaudhuri  
*Science Translational Medicine*, 2018.
9. [Methods for high-dimensional analysis of cells dissociated from cryopreserved synovial tissue](#)  
Laura T. Donlin\*, Deepak A. Rao\*, Kevin Wei, **Kamil Slowikowski**, Mandy J. McGeachy, Jason D. Turner, Nida Meednu, Fumitaka Mizoguchi, Maria Gutierrez-Arcelus, David J. Lieb, Joshua Keegan, Kaylin Muskat, Joshua Hillman, Cristina Rozo, Edd Ricker, Thomas M. Eisenhaure, Shuqiang Li, Edward P. Browne, Adam Chicoine, Danielle Sutherby, Akiko Noma, Accelerating Medicines Partnership RA/SLE Network, Chad Nusbaum, Stephen Kelly, Alessandra B. Pernis, Lionel B. Ivashkiv, Susan M. Goodman, William H. Robinson, Paul J. Utz,

James A. Lederer, Ellen M. Gravallese, Brendan F. Boyce, Nir Hacohen, Costantino Pitzalis, Peter K. Gregersen, Gary S. Firestein, Soumya Raychaudhuri, Larry W. Moreland, V. Michael Holers, Vivian P. Bykerk, Andrew Filer, David L. Boyle, Michael B. Brenner and Jennifer H. Anolik  
*Arthritis Research and Therapy*, 2018.

8. [Heritability enrichment of specifically expressed genes identifies disease-relevant tissues and cell types](#)  
 Hilary K. Finucane, Yakir A. Reshef, Verner Anttila, **Kamil Slowikowski**, Alexander Gusev, Andrea Byrnes, Steven Gazal, Po-Ru Loh, Caleb Lareau, Noam Shores, Giulio Genovese, Arpiar Saunders, Evan Macosko, Samuela Pollack, The Brainstorm Consortium, John R. B. Perry, Jason D. Buenrostro, Bradley E. Bernstein, Soumya Raychaudhuri, Steven McCarroll, Benjamin M. Neale, Alkes L. Price  
*Nature Genetics*, 2018.
7. [Refining the role of de novo protein-truncating variants in neurodevelopmental disorders by using population reference samples](#)  
 Jack A Kosmicki, Kaitlin E Samocha, Daniel P Howrigan, Stephan J Sanders, **Kamil Slowikowski**, Monkol Lek, Konrad J Karczewski, David J Cutler, Bernie Devlin, Kathryn Roeder, Joseph D Buxbaum, Benjamin M Neale, Daniel G MacArthur, Dennis P Wall, Elise B Robinson, Mark J Daly  
*Nature Genetics*, 2017.
6. [Pathologically expanded peripheral T helper cell subset drives B cells in rheumatoid arthritis](#)  
 Deepak A. Rao, Michael F. Gurish, Jennifer L. Marshall, **Kamil Slowikowski**, Chamith Y. Fonseka, Yanyan Liu, Laura T. Donlin, Lauren A. Henderson, Kevin Wei, Fumitaka Mizoguchi, Nikola C. Teslovich, Michael E. Weinblatt, Elena M. Massarotti, Jonathan S. Cobllyn, Simon M. Helfgott, Yvonne C. Lee, Derrick J. Todd, Vivian P. Bykerk, Susan M. Goodman, Alessandra B. Pernis, Lionel B. Ivashkiv, Elizabeth W. Karlson, Peter A. Nigrovic, Andrew Filer, Christopher D. Buckley, James A. Lederer, Soumya Raychaudhuri, Michael B. Brenner  
*Nature*, 2017.
5. [A method to decipher pleiotropy by detecting underlying heterogeneity driven by hidden subgroups applied to autoimmune and neuropsychiatric diseases](#)  
 Buhm Han, Jennie G Pouget, **Kamil Slowikowski**, Eli Stahl, Cue Hyunkyu Lee, Dorothee Diogo, Xinli Hu, Yu Rang Park, Eunji Kim, Peter K Gregersen, Solbritt Rantapää Dahlqvist, Jane Worthington, Javier Martin, Steve Eyre, Lars Klareskog, Tom Huizinga, Wei-Min Chen, Suna Onengut-Gumuscu, Stephen S Rich, Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium, Naomi R Wray, Soumya Raychaudhuri  
*Nature Genetics*, 2016.
4. [Disentangling the Effects of Colocalizing Genomic Annotations to Functionally Prioritize Non-coding Variants within Complex-Trait Loci](#)  
 Gosia Trynka, Harm-Jan Westra, **Kamil Slowikowski**, Xinli Hu, Han Xu, Barbara E. Stranger, Robert J. Klein, Buhm Han, Soumya Raychaudhuri  
*The American Journal of Human Genetics*, 2015.
3. [Regulation of gene expression in autoimmune disease loci and the genetic basis of proliferation in CD4+ effector memory T cells](#)  
 Xinli Hu, Hyun Kim, Towfique Raj, Patrick J. Brennan, Gosia Trynka, Nikola Teslovich, **Kamil Slowikowski**, Wei-Min Chen, Suna Onengut, Clare Baecher-Allan, Philip L. De Jager, Stephen S. Rich, Barbara E. Stranger, Michael B. Brenner, Soumya Raychaudhuri  
*PLoS Genetics*, 2014.
2. [Common genetic variants modulate pathogen-sensing responses in human dendritic cells](#)  
 Mark N. Lee\*, Chun Ye\*, Alexandra-Chloé Villani, Towfique Raj, Weibo Li, Thomas M. Eisenhaure, Selina H. Imboywa, Portia I. Chipendo, E. Ann Ran, **Kamil Slowikowski**, Lucas D. Ward, Khadir Raddassi, Cristin McCabe, Michelle H. Lee, Irene Y. Frohlich, David A. Hafler, Manolis Kellis, Soumya Raychaudhuri, Feng Zhang, Barbara

E. Stranger, Christophe O. Benoist, Philip L. De Jager, Aviv Regev, Nir Hacohen  
*Science*, 2014.

1. [Computational and experimental analyses of retrotransposon-associated minisatellite DNAs in the soybean genome](#)  
Lauren S. Mogil\*, **Kamil Slowikowski\***, Howard M. Laten  
*BMC Bioinformatics* 2012.

#### PREPRINTS

2. [Using genetics to differentiate patients with similar symptoms: application to inflammatory arthritis in the rheumatology outpatient clinic](#)  
Rachel Knevel, Saskia le Cessie, Chikashi C. Terao, **Kamil Slowikowski**, Jing Cui, Tom W.J. Huizinga, Karen H. Costenbader, Katherine P. Liao, Elizabeth W. Karlson, Soumya Raychaudhuri  
*bioRxiv*, 2019.
1. [UBiT2: a client-side web-application for gene expression data analysis](#)  
Jean Fan, David Fan, **Kamil Slowikowski**, Nils Gehlenborg, Peter Kharchenko  
*bioRxiv*, 2017. [🔗 Try the app](#)

Updated March 10, 2020