

# Kevin Bonham, Ph.D.

Senior Research Scientist

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## Education

2008 - 2014 | **Harvard - Cambridge, MA**  
*PhD in Immunology*  
Thesis: Cellular and Biochemical Events in Toll-like Receptor Signaling

2002 - 2006 | **University of California, San Diego**  
*BS Cum laude in Biochemistry and Cell Biology*

## Research Positions

Jan 2019 - Present | **Wellesley College – Wellesley, MA**  
*Senior Research Scientist*  
Promoted from Research Scientist to Senior RS in March, 2021.  
Projects:

- ECHO / Khula - Longitudinal cohorts of child brain development and the microbiome
- PASC - Subsets of Post-acute sequelae of COVID-19 (Long-COVID) with machine learning
- Microbiome.jl - software for microbial community data analysis
- GaPLAC - Gaussian Process modeling software tool for microbiome analysis

May 2017 - Dec 2018 | **Broad Institute and Harvard T.H. Chan School of Public Health – Boston, MA**  
*Postdoctoral Fellow - Huttenhower Lab*  
Projects:

- Human microbiome associations with inflammatory arthritis
- Human Microbiome Project phase 2 (HMP-II)
- Computational infrastructure for juvenile diabetes research consortium

May 2014 - Apr 2017 | **Harvard University and UCSD – Cambridge, MA**  
*Postdoctoral Fellow - Dutton Lab*  
Project: Horizontal gene transfer (HGT) in cheese-associated bacteria

Sep 2008 - Apr 2014 | **Boston Children's Hospital – Boston, MA**  
*Graduate Research Assistant - Kagan Lab*  
Project: Cellular localization and function in endosomal Toll-like receptor (TLR) signaling

May 2006 - Aug 2008 | **Scripps Research Institute – La Jolla, CA**  
*Lab Manager, Research Technician - Mowen Lab*  
Project: Small molecule inhibitors of enzymes (PRMTs) in T-cell signaling

## Honors and Awards

2014 | Jeffrey Modell Immunology Prize

2003 - 2006 | Provost's Honor

2002 - 2006 | UCSD Millenium Scholarship

## Research Funding

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### Current

2021 - present | **Co-Investigator on Wellcome LEAP 1kD**  
*A multi-scale approach to characterizing developing executive function*

### Past

2018-2019 | **Sloan Foundation - JuliaLang Diversity and Inclusion Award**  
*Increasing Representation of Women in Computational Biology*

2009 - 2013 | **NSF Graduate Research Fellowship** *Integration of Distinct Signaling Pathways: Toll-like Receptors and Cytokine-Activated Macrophages*

## Major Research Interests

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1. Effects of microbial metabolism on neurocognitive development
2. Dynamics of microbial communities
3. Computational and statistical methods for longitudinal microbiome research

### Narrative report

The development of the human gut microbiome and neurocognitive development are intimately linked. They are also dynamic, complex, and multi-causal, necessitating advanced computational and statistical methods. My current focus is on understanding the role of the human microbiome in neurocognitive development, and on developing software tools that facilitate investigation of longitudinal microbiome data and other complex biological data in humans.

## Publications

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\* indicates co-first authorship.

† indicates corresponding authorship.

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| In review<br>or in prep | <ol style="list-style-type: none"><li>1. <b>Bonham, K. S.</b>, Bottino, G. Z., McCann, S. H., Beauchemin, J., Weisse, E., O'Muircheartaigh, J., Huttenhower, C., D'Sa, V., Bruchhage, M., Deoni, S. C. L. &amp; Klepac-Ceraj, V. Gut microbes and their genes in infant children predict neurocognitive development. <i>In preparation</i>.</li><li>2. * Laue, H. E., <b>Bonham, K. S.</b>, Coker, M. O., Moroishi, Y., Pathmasiri, W., McRitchie, S., Sumner, S., Hoen, A. G., Karagas, M. R., Klepac-Ceraj, V. &amp; Madan, J. C. Prospective Association of the Infant Gut Microbiome with Autism-Related Behaviors in the ECHO Consortium. <i>In Review - Scientific Reports</i>.</li><li>3. Schoenborn, A. A., Yannarell, S. M., MacVicar, C. T., Barriga-Medina, N. N., Markillie, M., Mitchell, H., <b>Bonham, K. S.</b>, Leon-Reyes, A., Riveros-Iregui, D., Klepac-Ceraj, V. &amp; Shank, E. A. Microclimate is a strong predictor of the native and invasive plant-associated soil microbiota on San Cristóbal Island, Galápagos archipelago. <i>In revision - Environmental Microbiology</i>.</li><li>4. Thompson, K. N., <b>Bonham, K. S.</b>, <i>et al.</i> Alterations in the gut microbiome in inflammatory arthritis implicate key taxa and metabolic pathways across arthritis phenotypes. <i>In revision - Science Translational Medicine</i>.</li><li>5. * Woodruff, M. C., <b>Bonham, K. S.</b>, <i>et al.</i> Inflammation and autoreactivity define a discrete subset of patients with post-acute sequelae of COVID-19, or long-COVID. <i>In revision - Nature; medRxiv - doi: 10.1101/2021.09.21.21263845</i>.</li></ol> |
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- 2021 | 6. † **Bonham, K. S.**, Kayisire, A., Luo, A. & Klepac-Ceraj, V. Microbiome.jl and BiobakeryUtils.jl - Julia packages for working with microbial community data. *J. Open Source Softw.* **6**, 3876 (Nov. 2021).
7. Gauthier, A. E., Chandler, C. E., *et al.* Deep-sea microbes as tools to refine the rules of innate immune pattern recognition. en. *Sci Immunol* **6** (Mar. 2021).
8. Lewis, C. R., **Bonham, K. S.**, McCann, S. H., Volpe, A. R., D'Sa, V., Naymik, M., De Both, M. D., Huentelman, M. J., Lemery-Chalfant, K., Highlander, S. K., Deoni, S. C. L. & Klepac-Ceraj, V. Family SES Is Associated with the Gut Microbiome in Infants and Children. en. *Microorganisms* **9**, 1608 (July 2021).
9. Peterson, D., **Bonham, K. S.**, Rowland, S., Pattanayak, C. W., RESONANCE Consortium & Klepac-Ceraj, V. Comparative Analysis of 16S rRNA Gene and Metagenome Sequencing in Pediatric Gut Microbiomes. en. *Front. Microbiol.* **12**, 670336 (July 2021).
10. \* Tso, L., **Bonham, K. S.**, Fishbein, A., Rowland, S. & Klepac-Ceraj, V. Targeted High-Resolution Taxonomic Identification of *Bifidobacterium longum* subsp. *infantis* Using Human Milk Oligosaccharide Metabolizing Genes. en. *Nutrients* **13**, 2833 (Aug. 2021).
- 2019 | 16. Lloyd-Price, J., Arze, C., *et al.* Multi-omics of the gut microbial ecosystem in inflammatory bowel diseases. en. *Nature* **569**, 655–662 (May 2019).
19. Tett, A., Huang, K. D., *et al.* The Prevotella copri Complex Comprises Four Distinct Clades Underrepresented in Westernized Populations. en. *Cell Host Microbe* **26**, 666–679.e7 (Nov. 2019).
- 2017 | 22. † **Bonham, K. S.** & Stefan, M. I. Women are underrepresented in computational biology: An analysis of the scholarly literature in biology, computer science and computational biology. en. *PLoS Comput. Biol.* **13**, e1005134 (Oct. 2017).
24. **Bonham, K. S.**, Wolfe, B. E. & Dutton, R. J. Extensive horizontal gene transfer in cheese-associated bacteria. en. *Elife* **6** (June 2017).
- 2015 | 26. Brubaker, S. W., **Bonham, K. S.**, Zanoni, I. & Kagan, J. C. Innate immune pattern recognition: a cell biological perspective. en. *Annu. Rev. Immunol.* **33**, 257–290 (Jan. 2015).
- 2014 | 27. **Bonham, K. S.** & Kagan, J. C. Endosomes as platforms for NOD-like receptor signaling. en. *Cell Host Microbe* **15**, 523–525 (May 2014).
28. **Bonham, K. S.**, Orzalli, M. H., Hayashi, K., Wolf, A. I., Glanemann, C., Weninger, W., Iwasaki, A., Knipe, D. M. & Kagan, J. C. A promiscuous lipid-binding protein diversifies the subcellular sites of toll-like receptor signal transduction. *Cell* **156**, 705–716 (2014).
- 2010 | 30. **Bonham, K. S.**, Hemmers, S., Lim, Y.-H., Hill, D. M., Finn, M. G. & Mowen, K. A. Effects of a novel arginine methyltransferase inhibitor on T-helper cell cytokine production. *FEBS J.* **277**, 2096–2108 (2010).
31. Fathman, J. W., Gurish, M. F., Hemmers, S., **Bonham, K. S.**, Friend, D. S., Grusby, M. J., Glimcher, L. H. & Mowen, K. A. NIP45 controls the magnitude of the type 2 T helper cell response. en. *Proc. Natl. Acad. Sci. U. S. A.* **107**, 3663–3668 (Feb. 2010).

## Presentations

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### Invited Talks

- 2022 | **Boston Bacterial Meeting - Cambridge, MA**  
*Microbiomes and Microbial Ecosystems*  
Panel discussion
- 2019 | **Wellesley Science Center Faculty Seminar - Wellesley, MA**  
*The role of human gut microbial communities in the neurocognitive development of children*  
**JuliaCon - Baltimore, MD**  
*Raising Diversity and Inclusion among Julia users*  
with Anna Harris and Elwin van t' Wout
- 2018 | **HSPH Biostatistics Retreat - Boston, MA**  
*Strain-resolved microbial profiling in inflammatory arthritis*
- 2016 | **Bowdoin College Biology Department Seminar - Brunswick, ME**  
*Extensive horizontal gene transfer in cheese-associated bacteria*

### Workshops Taught

- 2018 | **Juvenile Diabetes Research Foundation Microbiome Initiative - Cambridge, MA**  
*The bioBakery for human microbiome epidemiology*  
**Wageningen University - Wageningen, Netherlands**  
*Creating Effective Graphics for Scientific Presentations*  
**SETAC, North Atlantic Chapter - Durham, NH**  
*Creating Effective Graphics for Scientific Presentations*  
**Physalia Microbiome Analysis, Berlin DE**  
*Taxonomic profiling with MetaPhlan*  
*Functional profiling with HUMAnN*  
*Targeted functional profiling with ShortBRED*  
*Searching for horizontal gene transfer with WAAFLE*
- 2017 | **ACM Conference on Bioinformatics, Computational Biology, and Health Informatics - Boston, MA**  
*Workshop on Algorithms in Bioinformatics - HUMAnN2*  
**PEGS Summit - Boston, MA**  
*Immunology for Drug Discovery Scientists*
- 2016 | **PEGS Summit - Boston, MA**  
*Immunology for Drug Discovery Scientists*

### Conference Posters

Since 2014

- 2020 | 11. **Bonham, K. S.**, Bruchhage, M., Rowland, S., Volpe, A., Dyer, K., D'Sa, V., Huttenhower, C., Deoni, S. & Klepac-Ceraj, V. *ASM Microbe - Gut microbes and their genes are associated with brain development and cognitive function in healthy children*. June 2020.
12. Peterson, D., Rowland, S., **Bonham, K. S.** & Klepac-Ceraj, V. *Boston Bacterial Meeting - Comparing early childhood gut microbiomes obtained from 16S rRNA gene and metagenome sequencing*. July 2020.
13. Tso, L., **Bonham, K. S.**, Rowland, S. & Klepac-Ceraj, V. *ASM Microbe - Baby steps: Characterizing Bifidobacterium longum subsp. infantis and its presence in American infants*. June 2020.
14. Tso, L., **Bonham, K. S.**, Rowland, S. & Klepac-Ceraj, V. *Boston Bacterial Meeting - Baby steps: Characterizing Bifidobacterium longum subsp. infantis and its presence in American infants*. July 2020.
- 2019 | 15. **Bonham, K. S.**, Rowland, S., Bruchhage, M., D'Sa, V., Huttenhower, C., Deoni, S. & Klepac-Ceraj, V. *Boston Bacterial Meeting - The relationship of the gut microbiome, environmental exposure and neurocognitive development in infants and children* May 2019.
17. Peterson, D., Rowland, S., Tso, L., **Bonham, K. S.**, Bruchhage, M., D'Sa, V., Huttenhower, C. & Klepac-Ceraj, V. *MIT-Harvard Microbiome Symposium - The relationship of the gut microbiome, environmental exposures, and neurocognitive development in infants and children* Mar. 2019.
18. Rowland, S., **Bonham, K. S.**, Bruchhage, M., D'Sa, V., Huttenhower, C., Deoni, S. & Klepac-Ceraj, V. *ASM Microbe - The early childhood gut microbiome, environmental exposures, and neurocognitive development*. June 2019.
- 2018 | 20. **Bonham, K. S.**, Peterson, D., Tso, L., Rowland, S., Deoni, S., Huttenhower, C. & Klepac-Ceraj, V. *Lake Arrowhead Microbial Genomics -The role of the gut microbiome in early childhood cognitive development* Sept. 2018.
21. **Bonham, K. S.**, Franzosa, E. A., Sayoldin, B., Ilott, N. E., Fehlner-Peach, H., Bullers, S., Littman, D. R., Young, S. P., Raza, K., Powrie, F. & Huttenhower, C. *Keystone: Microbiome, Host Resistance and Disease - Strain-resolved microbial and metabolomic profiling in inflammatory arthritis* Jan. 2018.
- 2017 | 23. **Bonham, K. S.**, Wolfe, B. E. & Dutton, R. J. *American Society of Microbiology, Mechanisms of Interbacterial Competition and Cooperation - Extensive horizontal transfer in cheese-associated bacteria* Mar. 2017.
- 2015 | 25. **Bonham, K. S.**, Wolfe, B. E. & Dutton, R. J. *ASM Microbe - Extensive horizontal transfer in cheese-associated bacteria* May 2015.
- 2014 | 29. **Bonham, K. S.**, Wolfe, B. E. & Dutton, R. J. *Boston Bacterial Meeting - Identifying horizontal transfer in cheese-associated bacteria* May 2014.

# Teaching

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## Positions

May 2016 - Apr 2017	<b>Harvard Medical School – Boston, MA</b> <i>Course Lead - Harvard Medical School Online</i> Course: Biochemistry Fundamentals
May 2014 - Apr 2016	<b>Harvard Medical School – Boston, MA</b> <i>Instructor in Microbiology and Immunobiology, Curriculum Fellow</i> Role: Founding instructor for HMS Masters of Medical Science in Immunology. Designed and taught 2 courses: <ul style="list-style-type: none"><li>• Research Methods in Experimental Immunology</li><li>• Understanding Immunology Literature</li></ul>
Spring 2015	<b>Harvard Extension School – Cambridge, MA</b> <i>Instructor</i> Course: Viruses: Molecular machines existing on the boundaries of life
Spring 2012, 2014	<b>Emmerson College – Boston, MA</b> <i>Adjunct Professor</i> Course: Plagues and Pandemics

## Graduate Courses

2018	<b>Harvard T.H. Chan School of Public Health - Boston, MA</b> <i>BST273 - Introduction to programming</i> Co-taught with Eric Franzosa.
2014-2016	<b>Harvard Medical School - Boston, MA</b> <i>IMM701 - Research Methods in Experimental Immunology</i>
2014-2016	<b>Harvard Medical School - Boston, MA</b> <i>IMM703 - Understanding Immunology Literature</i>

## Undergraduate Courses

2022	<b>Wellesley College - Wellesley, MA</b> <i>BISC314 - Environmental Microbiology Lab</i>
2021	<b>Wellesley College - Wellesley, MA</b> <i>BISC195 - Essential skills for computational biology</i>
2016-2017	<b>Harvard Medical School Online - Boston, MA</b> <i>Biochemistry essentials</i>
2015	<b>Harvard Extension School - Cambridge, MA</b> <i>BIOS E-157 - Viruses: A molecular arms race</i>
2013, 2014	<b>Emmerson College - Boston, MA</b> <i>SC214 - Plagues and Pandemics</i>

# Open Source Software

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## Package Author

2020-present	<b>GaPLAC - bioBakery</b> <a href="http://github.com/biobakery/GaPLAC">http://github.com/biobakery/GaPLAC</a> <ul style="list-style-type: none"><li>• Gaussian Process (GP) command line tool</li><li>• Ideal for longitudinal data, especially when sampled at irregular intervals</li><li>• Use GLM model-like syntax for specifying formula</li></ul>
2020-present	<b>Airtable.jl</b> <a href="http://github.com/kescobo/Airtable.jl">http://github.com/kescobo/Airtable.jl</a> <ul style="list-style-type: none"><li>• Interact with airtable.com REST API</li><li>• Rate-limiting control over request frequency to avoid limits</li><li>• FETCH, POST, and PATCH functionality using julia types</li></ul>
2016-present	<b>Microbiome.jl - BioJulia</b> <a href="http://github.com/BioJulia/Microbiome.jl">http://github.com/BioJulia/Microbiome.jl</a> <ul style="list-style-type: none"><li>• Data structures for biosamples, sample features (eg taxa)</li><li>• Attaching metadata to biosamples</li><li>• Data structures for taxonomic and community profiles</li><li>• Interfaces with statistical packages (eg <code>Distances.jl</code> and <code>Hclust.jl</code>)</li></ul> <b>BiobakeryUtils.jl - BioJulia</b> <a href="http://github.com/BioJulia/BiobakeryUtils.jl">http://github.com/BioJulia/BiobakeryUtils.jl</a> <ul style="list-style-type: none"><li>• Utilities for I/O of file types used with bioBakery tools</li><li>• Plotting utilities</li></ul>
2014-2016	<b>Kvasier - Dutton Lab</b> <a href="https://github.com/DuttonLab/kvasir">https://github.com/DuttonLab/kvasir</a> <ul style="list-style-type: none"><li>• Python-based command line tool for HGT discovery</li><li>• Stores genomic information, BLAST hits in MongoDB</li><li>• API for reading genomes, performing search, and generating tables</li></ul>

## Package Maintainer

2017-present	<b>Co-founder, BioJulia</b> <a href="http://github.com/BioJulia/">http://github.com/BioJulia/</a> <b>YAML.jl - JuliaData</b> <a href="http://github.com/JuliaData/YAML.jl">http://github.com/JuliaData/YAML.jl</a>
2018-present	<b>ClusterManagers.jl - JuliaParallel</b> <a href="https://github.com/JuliaParallel/ClusterManagers.jl">https://github.com/JuliaParallel/ClusterManagers.jl</a>
2020-present	<b>PowerAnalysis.jl</b> <a href="https://github.com/johnmyleswhite/PowerAnalysis.jl">https://github.com/johnmyleswhite/PowerAnalysis.jl</a>

## Package Contributor

2021	<b>Franklin.jl</b> - <a href="https://github.com/tlienart/Franklin.jl">https://github.com/tlienart/Franklin.jl</a> <b>LoggingExtras.jl</b> - <a href="https://github.com/JuliaLogging/LoggingExtras.jl">https://github.com/JuliaLogging/LoggingExtras.jl</a> <b>EcoBase.jl</b> - <a href="https://github.com/EcoJulia/EcoBase.jl">https://github.com/EcoJulia/EcoBase.jl</a>
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2020	<b>Julia</b> - <a href="https://github.com/JuliaLang/julia">https://github.com/JuliaLang/julia</a> <b>DataFrames.jl</b> - <a href="https://github.com/JuliaData/DataFrames.jl">https://github.com/JuliaData/DataFrames.jl</a> <b>CSV.jl</b> - <a href="https://github.com/JuliaData/CSV.jl">https://github.com/JuliaData/CSV.jl</a> <b>Documenter.jl</b> - <a href="https://github.com/JuliaDocs/Documenter.jl">https://github.com/JuliaDocs/Documenter.jl</a> <b>Literate.jl</b> - <a href="https://github.com/fredrikekre/Literate.jl">https://github.com/fredrikekre/Literate.jl</a> <b>LightGraphs.jl</b> - <a href="https://github.com/JuliaGraphs/LightGraphs.jl">https://github.com/JuliaGraphs/LightGraphs.jl</a> <b>SQLite.jl</b> - <a href="https://github.com/JuliaDatabases/SQLite.jl">https://github.com/JuliaDatabases/SQLite.jl</a>
2019	<b>StatsPlots.jl</b> - <a href="https://github.com/JuliaPlots/StatsPlots.jl">https://github.com/JuliaPlots/StatsPlots.jl</a> <b>Documenter.jl</b> - <a href="https://github.com/JuliaDocs/Documenter.jl">https://github.com/JuliaDocs/Documenter.jl</a> <b>DataDeps.jl</b> - <a href="https://github.com/oxinabox/DataDeps.jl">https://github.com/oxinabox/DataDeps.jl</a> <b>Clustering.jl</b> - <a href="https://github.com/JuliaStats/Clustering.jl">https://github.com/JuliaStats/Clustering.jl</a> <b>language-weave</b> - <a href="https://github.com/JunoLab/language-weave">https://github.com/JunoLab/language-weave</a>
2018	<b>Colors.jl</b> - <a href="https://github.com/JuliaGraphics/Colors.jl">https://github.com/JuliaGraphics/Colors.jl</a> <b>Julia</b> - <a href="https://github.com/JuliaLang/julia">https://github.com/JuliaLang/julia</a> <b>DataFrames.jl</b> - <a href="https://github.com/JuliaData/DataFrames.jl">https://github.com/JuliaData/DataFrames.jl</a> <b>SpatialEcology.jl</b> - <a href="https://github.com/EcoJulia/SpatialEcology.jl">https://github.com/EcoJulia/SpatialEcology.jl</a>
2017	<b>Distances.jl</b> - <a href="https://github.com/JuliaStats/Distances.jl">https://github.com/JuliaStats/Distances.jl</a> <b>BioSequences.jl</b> - <a href="https://github.com/BioJulia/BioSequences.jl">https://github.com/BioJulia/BioSequences.jl</a> <b>Bio.jl</b> - <a href="https://github.com/BioJulia/Bio.jl">https://github.com/BioJulia/Bio.jl</a>

## Outreach

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### Online Publications

2014-2019	<b>Co-founder of “Emmunity.org”</b> <b>Co-Host of the podcast Audiommunity</b>
2014-Present	<b>Creator: Adobe Illustrator for Scientists tutorial videos (youtube)</b>
2013-2016	<b>Blogger: “Food Matters” Scientific American Blogs</b> <i>Notable posts (links included):</i> <ul style="list-style-type: none"> <li>• What’s in your poo?</li> <li>• Time is the enemy, unless it’s colonic transfer time</li> <li>• Antibiotics and Obesity—an Unexpected Casualty in the War on Microbes</li> <li>• My new fermentation obsession</li> <li>• Probiotics, the immune system, and mouse balls</li> </ul>



2009-2013 | **Founder: “We, Beasties,” ScienceBlogs.com**

*Notable posts (links included):*

- Snow, cold, influenza and colds - Temperature and Infectious Disease
- Ebola Outbreak in Uganda - Both More and Less Frightening Than You Think
- The future of science publishing
- Autoimmunity to spunk
- A Bitter Sweet Nobel - Beutler, Janeway, and the Dawn of Innate Immunity

## Other

2016 | **Panel Moderator: Boston Fermentation Festival**

2015 | **Presenter: Boston Science Museum Health Science Fair**

2009-2013 | **Lecturer: Harvard Science in the News (SITN).**

- Autoimmunity and Disease: When the Body Attacks Itself (2009)
- Our Microbial Organ: The Good and Bad Bugs of The Human Gut (2010)
- How to Spot a Virus: The Origins of an Immune Response (2011)
- Avian flu and scientific censorship: When should scientists keep their mouths shut? (2012)
- Living Factories: Engineering Cells to Manufacture Molecules (2013)

2010-2012 | **Co-founder: Harvard Policy PATH**

2011 | **Student Advocate: ASBMB “Hill Day”**