

JAMES (JAMIE) T. MORTON

(513) · 907 · 9853 ◊ jamietmorton@gmail.com ◊ <https://github.com/mortonjt>

EDUCATION

Graduate 2015 - Present	University of California, San Diego, CA PhD student in Computer Science Expected Graduation Date: August 2018
Graduate 2014 - 2015	University of Colorado, Boulder, CO PhD student in Computer Science Integrative Quantitative Biology Program
Undergraduate 2010 - 2014	Miami University, Oxford, OH B.S. Computer Science (Cum Laude) B.S. Electrical Engineering (Cum Laude) B.S. Mathematics and Statistics B.S. Engineering Physics
Study Abroad Spring 2012	Hong Kong University of Science and Technology, Hong Kong

HONORS

- **NSF Graduate Fellow**, 2014 - present
- **Integrated Quantitative Biology Fellowship**, University of Colorado Boulder, 2014 - 2015
- **National Barry Goldwater Scholar**, 2013
- **Benjamin Harrison Scholar**, Miami University, 2010-2014
- **First place**, Institute of Navigation (ION) Autonomous Snowplow Competition, 2014
- **NSF REU**, Cold Spring Harbor Laboratories, Summer 2012
- **Provost Academic Achievement Award**, Miami University, 2012
- **Ohio Space Grant Scholar Award**, NASA, 2012 - 2014
- **Dean's List**, Miami University, 2010-13
- **R.L. Edwards Scholarship**, Department of Physics, Miami University, 2011, 2013
- **Mary Jeannette and Clifford Harvey Scholarship**, Department of Mathematics, Miami U., 2013
- **Mary Jean and Joseph R. Priest Scholarship**, Department of Physics, Miami University, 2012
- **President List**, Miami University, 2010-11
- **Nestle Scholar**, Computer Sci. and Software Eng. Dept, Miami University, 2011
- **Faculty Prize**, Department of Mathematics, Miami University, 2011
- **Joseph A. Culler Award**, Department of Physics, Miami University, 2010, 2011
- **NSF Travel Grant**, Coupling, Energetics, & Dynamics of Atmospheric Regions workshop, 2010
- **Wright Scholar**, Air Force Research Laboratory, Wright Patterson Air Force Base, 2009

RESEARCH INTERESTS

High dimensional statistics, compositional data analysis, multi-omics data fusion and machine learning techniques with applications to microbial ecology with special interests in functional genomics, microbial metabolism and microbial niche differentiation.

PUBLICATIONS

1. **Morton, J. T.**, Sanders, J., Quinn, R. A., McDonald, D., Gonzalez, A., Vázquez-Baeza, Y., Navas-Molina, J. A., Song, S. J., Metcalf, J. L., and Hyde, E. R. (2017a). Balance trees reveal microbial niche differentiation. *mSystems*, 2(1):e00162–16
2. **Morton, J. T.**, Toran, L., Edlund, A., Metcalf, J. L., Lauber, C., and Knight, R. (2017e). Uncovering the horseshoe effect in microbial analyses. *mSystems*, 2(1):e00166–16

3. Amir, A., McDonald, D., Navas-Molina, J. A., Debelius, J., **Morton, J. T.**, Hyde, E., Robbins-Pianka, A., and Knight, R. (2017a). Correcting for microbial blooms in fecal samples during room-temperature shipping. *mSystems*, 2(2):e00199–16
4. Amir, A., McDonald, D., Navas-Molina, J. A., Kopylova, E., **Morton, J. T.**, Zhenjiang, Z. X., Kightley, E. P., Thompson, L. R., Hyde, E. R., Gonzalez, A., and Knight, R. (2017b). Deblur rapidly resolves single-nucleotide community sequence patterns. *mSystems*, 2(2)
5. Vázquez-Baeza, Y., Gonzalez, A., Smarr, L., McDonald, D., **Morton, J. T.**, Navas-Molina, J. A., and Knight, R. (2017). Bringing the dynamic microbiome to life with animations. *Cell Host & Microbe*, 21(1):7–10
6. Vrbancac, A., Debelius, J. W., Jiang, L., **Morton, J. T.**, Dorrestein, P., and Knight, R. (2017a). An elegans (t) screen for drug-microbe interactions. *Cell Host & Microbe*, 21(5):555–556
7. Hill-Burns, E. M., Debelius, J. W., **Morton, J. T.**, Wissemann, W. T., Lewis, M. R., Wallen, Z. D., Peddada, S. D., Factor, S. A., Molho, E., Zabetian, C. P., Knight, R., and Payami, H. (2017). Parkinson’s disease and parkinson’s disease medications have distinct signatures of the gut microbiome. *Movement Disorders*, 32(5):739–749
8. Hemmings, S. M., Malan-Muller, S., van den Heuvel, L. L., Demmitt, B. A., Stanislawski, M. A., Smith, D. G., Bohr, A. D., Stamper, C. E., Hyde, E. R., **Morton, J. T.**, Marotz, C. A., Siebler, P. H., Braspenning, Maarten Irand Van Crieke, W., Irand Hoisington, A. J., Brenner, L. A., Postolache, T. T., McQueen, M. B., Krauter, K. S., Knight, R., Seedat, S., and Lowry, C. A. (2017). The microbiome in posttraumatic stress disorder and trauma-exposed controls: An exploratory study. *Psychosomatic Medicine*
9. Reber, S. O., Siebler, P. H., Donner, N. C., **Morton, J. T.**, Smith, D. G., Kopelman, J. M., Lowe, K. R., Wheeler, K. J., Fox, J. H., Jr., J. E. H., Greenwood, B. N., Jansch, C., Lechner, A., Schmidt, D., Uschold-Schmidt, N., Fchsl, A. M., Langgartner, D., Walker, F. R., Hale, M. W., Perez, G. L., Treuren, W. V., Gonzlez, A., Halweg-Edwards, A. L., Fleshner, M., Raison, C. L., Rook, G. A., Peddada, S. D., Knight, R., , and Lowry, C. A. (2016a). Immunization with a heat-killed preparation of the environmental bacterium mycobacterium vaccae promotes stress resilience in mice. *Proceedings of the National Academy of Sciences*, page 201600324
10. Gilbert, J. A., Quinn, R. A., Debelius, J., Xu, Z. Z., **Morton, J. T.**, Garg, N., Jansson, J. K., Dorrestein, P. C., and Knight, R. (2016). Microbiome-wide association studies link dynamic microbial consortia to disease. *Nature*, 535(7610):94–103
11. Nellore, A., Collado-Torres, L., Jaffe, A. E., Alquicira-Hernández, J., Wilks, C., Pritt, J., **Morton, J. T.**, Leek, J. T., and Langmead, B. (2016). Rail-rna: Scalable analysis of rna-seq splicing and coverage. *Bioinformatics*, page btw575
12. Petras, D., Nothias, L.-F., Quinn, R. A., Alexandrov, T., Bandeira, N., Bouslimani, A., Castro-Falcn, G., Chen, L., Dang, T., Floros, D. J., Hook, V., Garg, N., Hoffner, N., Jiang, Y., Kapon, C. A., Koester, I., Knight, R., Leber, C. A., Ling, T.-J., Luzzatto-Knaan, T., McCall, L.-I., McGrath, A. P., Meehan, M. J., Merritt, J. K., Mills, R. H., **Morton, J. T.**, Podvin, S., Protsyuk, I., Purdy, T., Satterfield, K., Searles, S., Shah, S., Shires, S., Steffen, D., White, M., Todoric, J., Tuttle, R., Wojnicz, A., Sapp, V., Vargas, F., Yang, J., Zhang, C., and Dorrestein, P. C. (2016). Mass spectrometry-based visualization of molecules associated with human habitats. *Analytical Chemistry*, 88(22):10775–10784. PMID: 27732780
13. Barberán, A., Dunn, R. R., Reich, B. J., Pacifici, K., Laber, E. B., Menninger, H. L., **Morton, J. T.**, Henley, J. B., Leff, J. W., Miller, S. L., and Fierer, N. (2015). The ecology of microscopic life in household dust. *Proceedings of the Royal Society of London B: Biological Sciences*, 282(1814)

14. **Morton, J. T.**, Freed, S. D., Lee, S. W., and Friedberg, I. (2015a). A large scale prediction of bacteriocin gene blocks suggests a wide functional spectrum for bacteriocins. *BMC bioinformatics*, 16(1):381
15. **Morton, J. T.**, Abrudan, P., Figueroa, N., Liang, C., and Karro, J. E. (2014a). Scope++: Sequence classification of homopolymer emissions. *Genomics*, 104(3):157–162

PUBLICATIONS IN REVIEW

1. Washburne, A., **Morton, J. T.**, Sanders, J., McDonald, D., Zhu, Q., Oliverio, AM, and Knight, R. (2017). Phylogenetic analysis of microbiome data: making sense of microbiomes in light of evolution. *Nature Microbiology*
2. McCall, L.-I., **Morton, J. T.**, Bernatchez, J. A., de Siqueira-Neto, J. L., Knight, R., C., D. P., and McKerrow, J. H. (2017). Molecular cartography of experimental cardiac infection by the parasite trypanosoma cruzi. *Analytical Chemistry*
3. Jiang, L., Amir, A., **Morton, J. T.**, Heller, R., Arias-Castro, E., and Knight, R. (2017). Discrete false discovery rate improves identification of differentially abundant microbes. *mSystems*
4. Thompson, L. R., Sanders, J. G., McDonald, D., Ladau, J., Locey, K. J., Navas-Molina, J. A., Prill, R. J., Gibbons, S. M., Gonzalez, A., Amir, A., Tripathi, A., Song, S. J., Vazquez-Baeza, Y., Kopylova, E., **Morton, J. T.**, Mirarab, S., Haroon, M. F., Kosciolk, T., Xu, Z. Z., Bokulich, N. A., Humphrey, G. C., Ackermann, G., Owens, S. M., Janssen, S., BrislawnK, C. J., Lefler, J., Hampton-Marcell, J., Zhu, Q., Kanbar, J., Berg-Lyons, D., Fierer, N., Shade, A., Pollard, K. S., Goodwin, K. D., JanssonK, J. K., Gilbert, J. A., Knight, R., and the Earth Microbiome Project Consortium (2017). The earth microbiome project. *Nature*
5. Vazquez-Baeza, Y., Callewaert, C., Debelius, J., Hyde, E., Marotz, C., **Morton, J. T.**, Swafford, A., Vrbanc, A., Dorrestein, P. C., and Knight, R. (2017). Impacts of the human gut microbiome on therapeutics. *Annual Reviews*
6. Metcalf, J. L., Song, S. J., **Morton, J. T.**, Weiss, S., Seguin-Orlando, A., Joly, F., Feh, C., Taberlet, P., Coissac, E., Amir, A., Willerslev, E., Knight, R., McKenzie, V., and Orlando, L. (2017). Domestication and captivity shape the horse gut microbiome. *Scientific Reports*
7. Vrbanc, A., Taylor, B. C., Aksenov, A., Callewaert, C., Debelius, J., Gonzalez, A., McCall, L.-I., McDonald, D., Melnik, A. V., **Morton, J. T.**, Navas, J., Quinn, R., Sanders, J. G., Swafford, A. D., Thompson, L. R., Tripathi, A., Vazquez-Baeza, Y., Xu, Z. Z., Zaneveld, J., Zhu, Q., Caporaso, J. G., C., D. P., and Knight, R. (2017b). Best practices for analyzing microbiomes. *Nature Reviews Microbiology*
8. Debelius, J. W., McDonald, D., Hyde, E., Ackermann, G., Aksenov, A., Chen, Y., Dorrestein, P. C., Dunn, R. R., Fahimpour, A. K., Gaffney, J., Gilbert, J. A., Gogul, G., Gonzalez, A., Green, J. L., Hugenholtz, P., Humphrey, G., Huttenhower, C., Jackson, M. A., Kelley, S. T., Knights, D., JoshuaLadau, Leach, J., Melnik, A., Metcalf, J. L., Montassier, E., **Morton, J. T.**, Navas-Molina, J., Peddada, S., Pollard, K. S., Rahnavard, G., Robbins-Pianka, A., Sangwan, N., Shorenstein, J., Spector, T., Thackray, V. G., Thompson, L. R., Vazquez-Baeza, Y., Wischmeyer, P., Wolfe, E., Consortium, T. A. G., and Knight, R. (2017). American gut: an open platform for citizen-science microbiome research. *Science*
9. Quinn, R. A., Comstock, W., Zhang, T., **Morton, J. T.**, Silva, R. d., Tran, A., Aksenov, A., Nothias-Scaglia1, L.-F., Wangpraseurt, D., Melnik, A. V., Ackerman, G., Conrad, D., Klapper, I., Knight, R., and Dorrestein, P. C. (2017). Niche partitioning of a polymicrobial infection driven by chemical gradients. *Nature*

PUBLICATIONS IN PREPARATION

1. Kapono, C. A., **Morton, J. T.**, Bouslimani, A., Orlinsky, K., Melnik, A. V., Knaan, T. L., Garg, N., Vázquez-Baeza, Y., Alexandrov, T., Protsyuk, I., Smarr, L., Knight, R., and Dorrestein, P. C. (2017). 3d chemical tracking of a human habitat and its human and microbial occupants. *Scientific Reports*
2. Martino, C., **Morton, J. T.**, Knight, R., and Zengler, K. (2017). Extrapolations across the vast unobserved microbial space with matrix completion. *mSystems*
3. Lavrinienko, A., Mappes, T., Tukalenko, E., Mousseau, T. A., Moller, A. P., Knight, R., **Morton, J. T.**, Thompson, L. R., and Watts, P. C. (2017). Environmental radiation alters the gut microbiome of the bank vole *Myodes glareolus*. *Nature*

PRESENTATIONS

- T., M. J. ((2017)). The microbiome and sex differences. Sex and the Kidneys: Sex Differences in Renal Disease Workshop in Bethesda, MD
- **Morton, J. T.**, Sanders, J., Quinn, R. A., McDonald, D., Gonzalez, A., Vázquez-Baeza, Y., Navas-Molina, J. A., Song, S. J., Metcalf, J. L., and Hyde, E. R. ((2017)b). Balances reveal microbial niche differentiation. CODAwork in Abbadia San Salvatore, Siena
- **Morton, J. T.**, Sanders, J., Quinn, R. A., McDonald, D., Gonzalez, A., Vázquez-Baeza, Y., Navas-Molina, J. A., Song, S. J., Metcalf, J. L., and Hyde, E. R. ((2017)d). Probabilities to balances: An alternative approach. Information Theory and Applications Workshop in San Diego, CA
- **Morton, J. T.**, Sanders, J., Quinn, R. A., McDonald, D., Gonzalez, A., Vázquez-Baeza, Y., Navas-Molina, J. A., Song, S. J., Metcalf, J. L., and Hyde, E. R. ((2017)c). From probabilities to balances: An alternative approach. Random Processes and Time Series: Theory and Applications in San Diego, CA
- Reber, S. O., Siebler, P. H., Donner, N. C., **Morton, J. T.**, Smith, D. G., Kopelman, J. M., Lowe, K. R., Wheeler, K. J., Fox, J. H., Jr., J. E. H., Greenwood, B. N., Jansch, C., Lechner, A., Schmidt, D., Uschold-Schmidt, N., Fchsl, A. M., Langgartner, D., Walker, F. R., Hale, M. W., Perez, G. L., Treuren, W. V., Gonzlez, A., Halweg-Edwards, A. L., Fleshner, M., Raison, C. L., Rook, G. A., Peddada, S. D., Knight, R., , and Lowry, C. A. ((2016)b). An immunization strategy for prevention of post-traumatic stress disorder (ptsd) promotes stress resilience in mice. University California San Diego Pediatrics Symposium in San Diego, CA
- **Morton, J. T.**, M., L., and R., K. ((2015)b). Uncovering the unknown: A new approach in analyzing microbiome data. NSF Data Science Workshop in University of Washington, Seattle, WA
- **Morton, J. T.**, Freed, S., Lee, S., and Friedberg, I. ((2014)b). Prediction of bacteriocin associated operons. Rocky Mountain Bioinformatics Conference
- **Morton, J. T.**, Freed, S. L., and S. Friedberg, I. (2014c). Discovering the next antibiotic. Ohio Space Grant Consortium in Cleveland OH
- **Morton, J. T.**, Abrudan, P., Karro, J., and Liang, C. (2013). Sequence classification of homopolymer emissions (scope). Great Lakes Bioinformatics Conference, Pittsburgh, PA
- **Morton, J. T.**, J. Karro, C. L. (2011). A novel approach for identifying poly(a) tails in raw cdna sequence data using general hidden markov models. Genome Informatics in Cold Spring Harbor, NY

PROFESSIONAL MEMBERSHIPS

- International Society of Computational Biology Student member, 2014-Present
- Association for Computing Machinery Student member, 2011-Present
- Sigma Pi Sigma, Tau Beta Pi, Eta Kappa Nu, 2014
- National Society of Collegiate Scholars, 2012 - 2013
- Institute of Electrical and Electronics Engineers Student member, 2011-2016
- IEEE Miami Student Chapter Treasurer, 2011 - 2012

PROFESSIONAL SERVICES

- Poster Reviewer for ISMB 2015 2014-2016

WORKSHOPS

- Teaching Assistant. Qiime2 workshop. Las Vegas, NV June 21-23, 2017
- Teaching Assistant. STAMPS, Woodshole, MA, August 2-13, 2016

TEACHING

- Instructor. Qiime2 workshop. UBC Vancouver August 23-25, 2017
- Instructor. Qiime2 workshop. UBC Kelowna August 21-22, 2017

Student Mentoring

- Jue Wang, undergraduate student, Summer 2017
- Cameron Martino, undergraduate student, Spring 2017
- Liam Toran, masters student, Summer 2016
- Kayla Orlinsky, undergraduate student, Spring 2016

SKILLS

Foreign Language Skills

- Chinese – Fluent in Mandarin and competent in written Chinese

Highly skilled in the following languages and computational platforms

- Python • C/C++ • Java • Javascript • L^AT_EX • ROS • Hadoop
- Matlab • R • Unix • SQL • OpenCL • CUDA • git

OPEN SOURCE CONTRIBUTIONS

- Gneiss (Core Maintainer)
- Sci-kit Bio (Developer)
- Emperor (Developer)
- Qiime2 (Contributor)
- Deblur (Contributor)
- Micronota (Developer)
- BOA: Bacteriocin Operon Associator (Lead Developer)
- SCOPE++: Sequence Classification Of homoPolymer Emissions (Lead Developer)
- Rail-RNA (Contributor)
- Statsmodels (Contributor)
- Scipy (Contributor)
- Biopython (Contributor)

PERSONAL INTERESTS

- Music: Piano, Cello, Guitar, and Voice (base).
- Sports: Surfing, Scuba Diving, Ice Hockey, Skiing, Hiking, Biking, Kung Fu
- Others: Traveling, Cooking