

Andrew Collin Stewart

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EXPERTISE

Data Analysis	Decision Analytics	Predictive Modeling
Research, Reports and Forecasts	Dashboards & Visualization	Big Data Queries
Data Mining	Machine Learning	Business Intelligence (BI)

PROFESSIONAL EXPERIENCE

Twist Bioscience	2015-2017
Scientist (Bioinformatics)	
Veracyte	2014-2015
Scientist (Data Analysis Team)	
Smithsonian Institution	2013-2015
Data Science Specialist (Office of the CIO)	
Keygene	2009-2012
Bioinformatician (Applied Systems Biology)	
Navy Medical Research Center	2006-2009
Research Assistant (Genomics)	

EDUCATION EXPERIENCE

Johns Hopkins University, M.S. Computer Science	2015
University of Maryland College Park, B.S. Biology	2006

TECHNICAL SKILLS

Programming	R, Python, Rust, C/C++, Java
Algorithms	Lorem ipsum dolor sit amet, consectetur adipiscing elit. Quisque cursus blandit odio, sit amet efficitur purus lobortis id. Nulla malesuada ac neque eu scelerisque.
Statistics	Lorem ipsum dolor sit amet, consectetur adipiscing elit. Quisque cursus blandit odio, sit amet efficitur purus lobortis id. Nulla malesuada ac neque eu scelerisque.

PUBLICATIONS

Chen, Peter E., Christopher Cook, Andrew C. Stewart, Niranjana Nagarajan, Dan D. Sommer, Mihai Pop, Brendan Thomason, et al. 2010. "Genomic Characterization of the Yersinia Genus." *Genome Biology* 11 (January):R1. <https://doi.org/10.1186/gb-2010-11-1-r1>.

Chen, Peter E., Kristin M. Willner, Amy Butani, Shakia Dorsey, Matroner George, Andrew Stewart, Shannon M. Lentz, et al. 2010. "Rapid Identification of Genetic Modifications in *Bacillus Anthracis* Using Whole Genome Draft Sequences Generated by 454 Pyrosequencing." *PLOS ONE* 5 (8):e12397. <https://doi.org/10.1371/journal.pone.0012397>.

"Meetings - Paper Digital Library." n.d. Accessed December 6, 2017. <https://dl.sciencesocieties.org/publications/meetings/2011am/8617/67897>.

Mukhopadhyay, Sanghamitra, Arya Akmal, Andrew C. Stewart, Ru-ching Hsia, and Timothy D. Read. 2009. "Identification of *Bacillus Anthracis* Spore Component Antigens Conserved Across Diverse *Bacillus Cereus* Ssensu Lato Strains." *Molecular & Cellular Proteomics* 8 (6):1174–91. <https://doi.org/10.1074/mcp.M800403-MCP200>.

Pagan, Moraima, Richard T. Kloos, Chu-Fang Lin, Kevin J. Travers, Hajime Matsuzaki, Ed Y. Tom, Su Yeon Kim, et al. 2016. "The Diagnostic Application of RNA Sequencing in Patients with Thyroid Cancer: An Analysis of 851 Variants and 133 Fusions in 524 Genes." *BMC Bioinformatics* 17 (1):S6. <https://doi.org/10.1186/s12859-015-0849-9>.

Stewart, Andrew C., Brian Osborne, and Timothy D. Read. 2009. "DIYA: A Bacterial Annotation Pipeline for Any Genomics Lab." *Bioinformatics* 25 (7):962–63. <https://doi.org/10.1093/bioinformatics/btp097>.

Zwick, Michael E., Sandeep J. Joseph, Xavier Didelot, Peter E. Chen, Kimberly A. Bishop-Lilly, Andrew C. Stewart, Kristin Willner, et al. 2012. "Genomic Characterization of the *Bacillus Cereus* Ssensu Lato Species: Backdrop to the Evolution of *Bacillus Anthracis*." *Genome Research* 22 (8):1512–24. <https://doi.org/10.1101/gr.134437.111>.

Zwick, Michael E., Maureen P. Kiley, Andrew C. Stewart, Alfred Mateczun, and Timothy D. Read. 2008. "Genotyping of *Bacillus Cereus* Strains by Microarray-Based Resequencing." *PLOS ONE* 3 (7):e2513. <https://doi.org/10.1371/journal.pone.0002513>.