

John Didion | PhD

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Education

- 2008.08 to 2014.05 **Doctor of Philosophy, Bioinformatics and Computational Biology**
University of North Carolina, Chapel Hill, NC
- 1996.09 to 2001.06 **Bachelor of Science, Computer Science**
Northwestern University, Evanston, IL

Research Experience

- 2014.09 to Present **National Human Genome Research Institute** Bethesda, MD
Postdoctoral Fellow, Laboratory of Dr. Francis S Collins
Investigating the role of epigenetics in common complex disease, especially type 2 diabetes.
- 2009.05 to 2014.08 **University of North Carolina at Chapel Hill** Chapel Hill, NC
Research Assistant, Laboratory of Dr. Fernando Pardo-Manuel de Villena
Characterized the genetic determinants of two meiotic drive systems in the mouse. First, conducted a GWAS of wild mice to identify genes associated with the accumulation of Robertsonian translocations. Second, mapped a novel meiotic drive locus, R2d2, and multiple modifier loci responsible for extreme TRD in the Collaborative Cross. Also developed CLASP, a software tool for validation of cell lines used in research.
- 2007.08 to 2008.08 **Institute for Systems Biology** Seattle, WA
Computational Biology Software Engineer, Laboratory of Dr. Ruedi Aebersold
Developed informatics software to support proteomics research. Created TIQAM, a work flow management system for MRM (Multiple Reaction Monitoring) experiments.

Industry Experience

- 2004.09 to 2007.08 **Muze, Inc.** Seattle, WA
Software Development Engineer
- 2003.09 to 2004.09 **Encyclopaedia Britannica, Online Services** Chicago, IL
Software Developer
- 2001.05 to 2003.08 **ThoughtWorks, LLC** Chicago, IL
Software Developer/Consultant

Teaching Experience

- 2013.09 to 2014.05 **University of North Carolina at Chapel Hill**
Undergraduate senior research project, Mentor
- 2013.09 to 2013.12 **University of North Carolina at Chapel Hill**
Genetic Analysis 2, Teaching Assistant

Publications

Journal Articles.....

- [12] Chandler RL, Damrauer JS, Raab JR, Schisler JC, Wilkerson MD, **Didion JP**, et al. "Coexistent ARID1A-PIK3CA mutations promote ovarian clear-cell tumorigenesis through pro-tumorigenic inflammatory cytokine signaling," *Nature Communications*, vol. 6, p. 6118, 2015. DOI: 10.1038/ncomms7118.
- [11] Crowley JJ, ...13 others..., **Didion JP**, et al. "Analyses of allele-specific gene expression in highly divergent mouse crosses identifies pervasive allelic imbalance," *Nature Genetics*, online, 2015. DOI: 10.1038/ng.3222.
- [10] **Didion JP**, Morgan AP, Clayshulte AMF, et al. "A Multi-Megabase Copy Number Gain Causes Maternal Transmission Ratio Distortion on Mouse Chromosome 2," *PLoS Genetics*, vol. 11, no. 2, e1004850, 2015. DOI: 10.1371/journal.pgen.1004850.
- [9] **Didion JP**, Buus RJ, Naghashfar Z, et al. "SNP array profiling of mouse cell lines identifies their strains of origin and reveals cross-contamination and widespread aneuploidy," *BMC Genomics*, vol. 15, p. 847, 2014. DOI: 10.1186/1471-2164-15-847.
- [8] **Didion JP** and Pardo-Manuel de Villena F, "Deconstructing *Mus gemischus*: advances in understanding ancestry, structure, and variation in the genome of the laboratory mouse," *Mammalian Genome*, vol. 24, no. 1-2, pp. 1-20, 2013. DOI: 10.1007/s00335-012-9441-z.
- [7] Calaway JD, Lenarcic AB, **Didion JP**, et al. "Genetic architecture of skewed X inactivation in the laboratory mouse," *PLoS Genetics*, vol. 9, no. 10, e1003853, 2013. DOI: 10.1371/journal.pgen.1003853.
- [6] Collaborative Cross Consortium, "The genome architecture of the Collaborative Cross mouse genetic reference population," *Genetics*, vol. 190, no. 2, pp. 389-401, 2012. DOI: 10.1534/genetics.111.132639.
- [5] **Didion JP**, Yang H, Sheppard K, et al. "Discovery of novel variants in genotyping arrays improves genotype retention and reduces ascertainment bias," *BMC Genomics*, vol. 13, p. 34, 2012. DOI: 10.1186/1471-2164-13-34.
- [4] Aylor DL, ...11 others..., **Didion JP**, et al. "Genetic analysis of complex traits in the emerging Collaborative Cross," *Genome Research*, vol. 21, no. 8, pp. 1213-1222, 2011. DOI: 10.1101/gr.111310.110.
- [3] Yang H, Wang JR, **Didion JP**, et al. "Subspecific origin and haplotype diversity in the laboratory mouse," *Nature Genetics*, vol. 43, no. 7, pp. 648-655, 2011. DOI: 10.1038/ng.847.
- [2] Eisener-Dorman AF and **Didion JP** and Santos C and Calaway JD, "The 23rd International Mammalian Genome Conference meeting report," *Mammalian Genome*, vol. 21, no. 5-6, pp. 217-223, 2010. DOI: 10.1007/s00335-010-9265-7.
- [1] Lange V, Malmström JA, **Didion JP**, et al. "Targeted quantitative analysis of *Streptococcus pyogenes* virulence factors by multiple reaction monitoring," *Molecular & Cellular Proteomics*, vol. 7, no. 8, pp. 1489-1500, 2008. DOI: 10.1074/mcp.M800032-MCP200.

Oral Presentations

2013.09	27th Intl. Mammalian Genome Conference	Salamanca, Spain
2013.05	12th Annual Meeting of the Complex Traits Community	Madison, WI
2012.09	16th EBM Conference	Marseille, France
2012.09	UNC Genetics Department Retreat	Myrtle Beach, SC
2010.10	24th Intl. Mammalian Genome Conference	Heraklion, Greece
2010.05	9th Annual Meeting of the Complex Traits Community	Chicago, IL
2009.09	23rd Intl. Mammalian Genome Conference	San Diego, CA

Poster Presentations

2014.10	28th Intl. Mammalian Genome Conference	Bar Harbor, ME
2011.09	UNC Genetics Department Retreat	Myrtle Beach, SC
2011.06	Genetics Society of America Mouse Genetics	Washington D.C.
2011.06	National Centers for Systems Biology Annual Meeting	Duke University
2010.09	UNC Genetics Department Retreat	Myrtle Beach, SC
2009.09	UNC Genetics Department Retreat	Asheville, NC

Funding

2014.12	NIH Intramural Sequencing Center Pilot Grant <i>Epigenomic regulation of glucose response in a human pancreatic islet β cell line</i> Institutional award with funding for sequencing services
2014.12 to 2015.04	Department of Health and Human Services Ignite <i>LabGenius to The Smart Lab Notebook for Scientists</i> Incubator program with USD 5,000 funding
2009.09 to 2010.08	Bioinformatics and Computational Biology Training Grant <i>NIH GM067553-04</i> Institution-awarded fellowship
2009.09	International Mammalian Genome Society Fellowship <i>Travel grant</i>

Honors & Awards

- 2014.12 **Dean's Distinguished Dissertation Award**
University of North Carolina at Chapel Hill
Nominee from the Bioinformatics and Computational Biology program
- 2013.09 **Verne Chapman Young Scientist Award**
International Mammalian Genome Society
Best talk at the International Mammalian Genome Conference
- 2013.05 **Chicago Prize**
Complex Traits Consortium
Best graduate student talk at the Complex Traits Consortium meeting
- 2010.10 **Genome Research Award for Outstanding Poster**
International Mammalian Genome Society
Outstanding poster at the International Mammalian Genome Conference

Professional Memberships & Service

- 2013.09 to 2015.09 **International Mammalian Genome Society**
Secretariat member (honorary)
- 2013.08 **UNC Genetics Department Retreat**
Abstract Review Committee