

Dmitry Mozzherin

Curriculum Vitae

Illinois Natural History Survey
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🌐 <http://globalnames.org>



Education and Postdoctoral Training

- 1994–2001 **Postdoctoral Associate**, *State University of New York*, Stony Brook, NY, Department of Pharmacology.
- 1989–1993 **Graduate Student**, *The Engelhardt Institute of Molecular Biology*, Moscow, Russia, Lab of Chemical and Biochemical Analysis of Biopolymers and Cells.
Ph.D. in Biochemistry
- 1979–1985 **Undergraduate Student**, *Ural State University* — *Biology Major*, Yekaterinburg, Russia, Department of Physiology and Biochemistry of Plants.
Speciality: Biologist, Teacher of Biology and Chemistry

Professional and Academic Appointments

- 2015–present **Bioinformatician**, *University of Illinois*, Champaign, IL, Species File Group, Global Names Architecture.
- 2009–2015 **Research Associate**, *Marine Biological Laboratory*, Woods Hole, MA, Encyclopedia of Life, Global Names Architecture.
- 2008–2015 **Scientific Informatics Project Leader**, *Marine Biological Laboratory*, Woods Hole, MA, Encyclopedia of Life, Global Names Architecture.
- 2001–2008 **Research Instructor, Senior Programmer Analyst**, *State University of New York*, Stony Brook, NY, Department of Pharmacology, Department of Medical Informatics.
- 1993–1994 **Scientific Researcher**, *The Engelhardt Institute of Molecular Biology*, *Russian Academy of Sciences*, Moscow, Russia, Lab of Chemical and Biochemical Analysis of Biopolymers and Cells.
- 1985–1989 **Senior Engineer**, *Omutninsk Chemical Plant*, Kirov region, Russia.
Head of the group responsible for purification of *Avian Myeloblastosis Virus Reverse Transcriptase*

Experience and Skills

Science	Molecular Biologist, Biochemist, Biodiversity Informatician, Computer Scientist
Programming	Go, Ruby, Scala, Elm, Python, R, JavaScript, C, Assembly, Java, Perl, PHP, Cold Fusion, etc.
Management	NSF Primary Investigator, Agile/Scrum Master
DevOps	Linux System Administration, Ansible, Docker, Kubernetes, Chef
Markup	HTML, CSS, SASS, HAML, Markdown, \LaTeX
Design/Art	Blender, Photoshop, Gimp, Photography, Sculpture

Honors and Awards

2014	PI on NSF Award DBI-1356347	<i>University of Illinois</i>
2002	PI on NLM Award 1G07LM007762-01	<i>State University of New York at Stony Brook</i>
1996	Catacosinos Cancer Research Award	<i>State University of New York at Stony Brook</i>
1993	Outstanding young scientist	<i>The Engelhardt Institute of Molecular Biology</i>

Extra-Curricular Awards

2006	Winner of Best of Wild Life Photography	<i>Smithsonian National Museum of Natural History (as part of Long Island Wild Life Photography submission)</i>
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Interests

- Biodiversity Informatics
- Open Source Programming
- Sculpture
- Wild Life Photography
- Molecular Biology
- Machine Learning
- 3D Modelling
- Biking

Teaching

2018-2019	Mentoring students for Computer Science 492 Course	<i>University of Illinois</i>
2015-2019	Mentoring students working on Phylotastic project	<i>Phylotastic project</i>
2010, 2011, 2015	Computer Science Students Mentoring	<i>Google Summer of Code</i>
2011, 2012	Computer Science Students Mentoring	<i>UCOSP Capstone Project Canada</i>
2013,2014	Scientific Illustration Students Mentoring	<i>Rode Island School of Design, Marine Biological Laboratory</i>

Selected Open Source Software (principle of major contributor)

2020–present	BHL Names	<i>Linking taxonomic publications to BHL pages</i>
2020–present	Global Names Verifier (Go)	<i>Scientific name reconciliation/resolution</i>
2018–present	Global Names Parser (Go)	<i>Scientific name parsing</i>
2017–present	bhlindex	<i>Indexer of scientific names in Biodiversity Heritage Library</i>
2017–present	gnfinder	<i>General purpose scientific names detection</i>
2015–present	Global Names Parser (Scala)	<i>Scientific name parsing</i>
2015–present	Global Names Resolver	<i>Taxonomic resolution of scientific names</i>
2015–present	Global Names Crossmap	<i>Bulk resolution of scientific names from CSV file</i>
2012–present	Global Names Recognition and Discovery	<i>Scientific names finding service</i>
2012–present	NameSpotter	<i>A library for finding scientific names</i>
2012–present	DamerauLevenshtein	<i>A library to determine edit distance between strings</i>
2012–present	BHL Indexer	<i>A tool for finding names in Biodiversity Heritage Library</i>
2011–present	DwCA Hunter	<i>A convertor of biodiversity resources to DarwinCore Archive</i>
2010–present	DwC Archive	<i>A DarwinCore Archive reader/writer</i>
2009–present	Taxamatch for Ruby	<i>A fuzzy-matching library for scientific names</i>
2009–2013	Global Names Interactive Editor	<i>Classification Editor</i>

2008–present	Global Names Index	<i>Global index of scientific names</i>
2008–2015	Biodiversity	<i>Scientific names parser</i>
2008–2015	Encyclopedia of Life	<i>An online encyclopedia about known species</i>
2000–2004	BioMail	<i>Biomedical publications alert service</i>

Peer-reviewed Publications

1. Mozzherin, Dmitry, Myltsev, Alexander, Patterson, David: "gnparser": a powerful parser for scientific names based on Parsing Expression Grammar. *BMC Bioinformatics* (2017). doi:10.1186/s12859-017-1663-3
2. Patterson, D., Mozzherin, D., Shorthouse, D., Thessen, A.: Challenges with using names to link digital biodiversity information. *Biodiversity Data Journal* **4**, 8080 (2016). doi:10.3897/BDJ.4.e8080
3. Kripke, E., Mozzherin, D., Senft, S., Hanlon, R.: Visualizing Biological Complexity in Cephalopod Skin: A Synergy of Art and Science Technologies. *Leonardo* **48**(5), 486–487 (2015). doi:10.1162/LEON_a_01124
4. Fischer, A.H.L., Mozzherin, D., Eren, a.M., Lans, K.D., Wilson, N., Cosentino, C., Smith, J.: SeaBase: a multispecies transcriptomic resource and platform for gene network inference. *Integrative and comparative biology* **54**(2), 250–63 (2014). doi:10.1093/icb/icu065
5. Boyle, B., Hopkins, N., Lu, Z., Raygoza Garay, J.A., Mozzherin, D., Rees, T., Matasci, N., Narro, M.L., Piel, W.H., McKay, S.J., Lowry, S., Freeland, C., Peet, R.K., Enquist, B.J.: The taxonomic name resolution service: an online tool for automated standardization of plant names. *BMC bioinformatics* **14**(1), 16 (2013). doi:10.1186/1471-2105-14-16
6. Morris, R.a., Barve, V., Carausu, M., Chavan, V., Cuadra, J., Freeland, C., Hagedorn, G., Leary, P., Mozzherin, D., Olson, A., Riccardi, G., Teage, I., Whitbread, G.: Discovery and publishing of primary biodiversity data associated with multimedia resources: the audubon core strategies and approaches. *Biodiversity Informatics* (1), 185–197 (2013). doi:10.17161/bi.v8i2.4117
7. Thessen, A.E., Cui, H., Mozzherin, D.: Applications of natural language processing in biodiversity science. *Advances in bioinformatics* **2012**, 391574 (2012). doi:10.1155/2012/391574
8. Mozzherin, D.J., McConnell, M., Miller, H., Fisher, P.a.: Site-specific mutagenesis of *Drosophila* proliferating cell nuclear antigen enhances its effects on calf thymus DNA polymerase delta. *BMC biochemistry* **5**, 13 (2004). doi:10.1186/1471-2091-5-13
9. Fisher, P.A., Moutsakis, D.L., McConnell, M., Mozzherin, D.: A Single Amino Acid Change (E85K) in Human PCNA That Leads, Relative to Wild Type, to Enhanced DNA Synthesis by DNA Polymerase δ past Nucleotide Base Lesions (TLS) as Well as on Unmodified Templates. *Biochemistry* **43**(50), 15915–15921 (2004). doi:10.1021/bi048558x
10. Mozzherin, D.J., Tan, C.K., Downey, K.M., Fisher, P.A.: Architecture of the active DNA polymerase δ -proliferating cell nuclear antigen-template-primer complex. *Journal of Biological Chemistry* **274**(28), 19862–7 (1999). doi:10.1074/jbc.274.28.19862
11. Mozzherin, D.J., McConnell, M., Fisher, P.A.: *Drosophila* replication and repair proteins: proliferating cell nuclear antigen (PCNA). *Methods* **18**(3), 401–406 (1999). doi:10.1006/meth.1999.0798
12. Zaika, A., Mozzherin, D.J., Tan, C.K., Downey, K.M., Fisher, P.A.: A two-dimensional support for selective binding of polyhistidine-tagged proteins: identification of a proliferating cell nuclear antigen point mutant with altered function in vitro. *Anal Biochem* **268**(2), 193–200 (1999)

13. Mozzherin, D.J., Shibutani, S., Tan, C.K., Downey, K.M., Fisher, P.a.: Proliferating cell nuclear antigen promotes DNA synthesis past template lesions by mammalian DNA polymerase delta. *Proceedings of the National Academy of Sciences of the United States of America* **94**(12), 6126–31 (1997)
14. Mozzherin, D.J., McConnell, M., Jasko, M.V., Krayevsky, A.A., Tan, C.K., Downey, K.M., Fisher, P.A.: Proliferating cell nuclear antigen promotes misincorporation catalyzed by calf thymus DNA polymerase delta. *J Biol Chem* **271**(49), 31711–31717 (1996)
15. Mozzherin, D.J., Fisher, P.A.: Human DNA polymerase epsilon: enzymologic mechanism and gap-filling synthesis. *Biochemistry* **35**(11), 3572–3577 (1996)
16. McConnell, M., Miller, H., Mozzherin, D.J., Quamina, a., Tan, C.K., Downey, K.M., Fisher, P.a.: The mammalian DNA polymerase delta–proliferating cell nuclear antigen–template-primer complex: molecular characterization by direct binding. *Biochemistry* **35**(25), 8268–74 (1996). doi:10.1021/bi9530649
17. Jasko, M.V., Fedorov, I.I., Atrazhev, A.M., Mozzherin, D.Y., Novicov, N.A., Bochkarev, A.V., Gurskaya, G.V., Krayevsky, A.A.: Synthesis, Molecular and Crystal Structure of 3'-N-Alkylamino-3'-deoxythymidines and Some Biochemical Properties of Their Phosphorous Esters. *Nucleosides and Nucleotides* **14**(1-2), 23–37 (1995). doi:10.1080/15257779508014650
18. Jasko, M.V., Semizarov, D.G., Victorova, L.S., Mozzherin, D., Krayevsky, A.A., Kukhanova, M.K.: New modified substrates for discriminating between human DNA polymerases alpha and epsilon. *FEBS Lett* **357**(1), 23–26 (1995). doi:001457939401319V [pii]
19. Kukhanova, M., Liu, S.H., Mozzherin, D., Lin, T.S., Chu, C.K., Cheng, Y.C.: L- and D-enantiomers of 2',3'-dideoxycytidine 5'-triphosphate analogs as substrates for human DNA polymerases. Implications for the mechanism of toxicity. *J Biol Chem* **270**(39), 23055–9 (1995)
20. Mozzherin, D.J., Atrazhev, A.M., Kukhanova, M.K.: [A method of isolation and properties of DNA-dependent DNA-polymerase epsilon from human placenta]. *Molekuliarnaia Biologiya* **26**(5), 999–1010 (1992)
21. Viktorova, L.S., Rozovskaia, T.A., Mozzherin, D.J., Krayevsky, A.A.: [Acyclic analogs of 2',3'-dideoxy-2',3'-didehydronucleoside-5'-triphosphates–terminators of DNA synthesis, catalyzed by a broad set of DNA polymerases]. *Molekuliarnaia Biologiya* **27**(1), 143–152 (1992)
22. Victorova, L.S., Dyatkina, N.B., Mozzherin, D.J., Atrazhev, A.M., Kraevsky, A.A., Kukhanova, M.K.: Formation of phosphonoester bonds catalyzed by DNA polymerases. *Nucleic Acids Research* **20**(4), 783–789 (1992). doi:10.1093/nar/20.4.783
23. Jasko, M.V., Atrazhev, A.M., Mozzherin, D.J., Novikov, N.A., Fedorov, I.I., Kraevsky, A.A.: [Synthesis and various biochemical properties of alkylated derivatives of 2',3'-dideoxy-3'-aminothymidine]. *Bioorganicheskaya Khimia* **18**(2), 299–301 (1992)