

# Michael Domaratzki

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1978468/publications.pdf>

Version: 2024-02-01

39  
papers

486  
citations

840776

11  
h-index

752698

20  
g-index

44  
all docs

44  
docs citations

44  
times ranked

660  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | MicroRNA-guided regulation of heat stress response in wheat. BMC Genomics, 2019, 20, 488.   | 2.8  | 78        |
| 2  | Continent-wide effects of urbanization on bird and mammal genetic diversity. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20192497.                              | 2.6  | 63        |
| 3  | Deep sequencing of wheat sRNA transcriptome reveals distinct temporal expression pattern of miRNAs in response to heat, light and UV. Scientific Reports, 2016, 6, 39373.               | 3.3  | 51        |
| 4  | State complexity of power. Theoretical Computer Science, 2009, 410, 2377-2392.  | 0.9  | 41        |
| 5  | Deletion along trajectories. Theoretical Computer Science, 2004, 320, 293-313.  | 0.9  | 36        |
| 6  | Competitive Fitness of Essential Gene Knockdowns Reveals a Broad-Spectrum Antibacterial Inhibitor of the Cell Division Protein FtsZ. Antimicrobial Agents and Chemotherapy, 2018, 62, . | 3.2  | 28        |
| 7  | Trajectory-based codes. Acta Informatica, 2004, 40, 491-527.  | 0.5  | 19        |
| 8  | Neptune: a bioinformatics tool for rapid discovery of genomic variation in bacterial populations. Nucleic Acids Research, 2017, 45, e159-e159.  | 14.5 | 16        |
| 9  | Simulating finite automata with context-free grammars. Information Processing Letters, 2002, 84, 339-344.   | 0.6  | 13        |
| 10 | Decidability of trajectory-based equations. Theoretical Computer Science, 2005, 345, 304-330.   | 0.9  | 13        |
| 11 | Transition complexity of language operations. Theoretical Computer Science, 2007, 387, 147-154.   | 0.9  | 12        |
| 12 | Minimality in template-guided recombination. Information and Computation, 2009, 207, 1209-1220.   | 0.7  | 12        |
| 13 | Semantic Shuffle on and Deletion Along Trajectories. Lecture Notes in Computer Science, 2004, , 163-174.  | 1.3  | 9         |
| 14 | Metabolic network prediction through pairwise rational kernels. BMC Bioinformatics, 2014, 15, 318.  | 2.6  | 8         |
| 15 | Lower bounds for the transition complexity of NFAs. Journal of Computer and System Sciences, 2008, 74, 1116-1130.   | 1.2  | 7         |
| 16 | INTRA-MOLECULAR TEMPLATE-GUIDED RECOMBINATION. International Journal of Foundations of Computer Science, 2007, 18, 1177-1186.   | 1.1  | 6         |
| 17 | Hairpin Structures Defined by DNA Trajectories. Lecture Notes in Computer Science, 2006, , 182-194.   | 1.3  | 6         |
| 18 | Assessing feature selection method performance with class imbalance data. Machine Learning With Applications, 2021, 6, 100170.  | 4.4  | 6         |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | NON-UNIQUENESS AND RADIUS OF CYCLIC UNARY NFAs. International Journal of Foundations of Computer Science, 2005, 16, 883-896.  | 1.1 | 5         |
| 20 | RESTRICTED SETS OF TRAJECTORIES AND DECIDABILITY OF SHUFFLE DECOMPOSITIONS. International Journal of Foundations of Computer Science, 2005, 16, 897-912.            | 1.1 | 5         |
| 21 | Codes defined by multiple sets of trajectories. Theoretical Computer Science, 2006, 366, 182-193.   | 0.9 | 5         |
| 22 | Bond-free DNA language classes. Natural Computing, 2007, 6, 371-402.  | 3.0 | 5         |
| 23 | Equivalence in template-guided recombination. Natural Computing, 2008, 7, 439-449.  | 3.0 | 5         |
| 24 | Hairpin Structures Defined by DNA Trajectories. Theory of Computing Systems, 2009, 44, 432-454.   | 1.1 | 5         |
| 25 | ABELIAN PRIMITIVE WORDS. International Journal of Foundations of Computer Science, 2012, 23, 1021-1033.   | 1.1 | 5         |
| 26 | IMPROVED BOUNDS ON THE NUMBER OF AUTOMATA ACCEPTING FINITE LANGUAGES. International Journal of Foundations of Computer Science, 2004, 15, 143-161.                  | 1.1 | 4         |
| 27 | Representing recursively enumerable languages by iterated deletion. Theoretical Computer Science, 2004, 314, 451-457.   | 0.9 | 4         |
| 28 | Somatic Copy Number Alteration-Based Prediction of Molecular Subtypes of Breast Cancer Using Deep Learning Model. Lecture Notes in Computer Science, 2017, , 57-63. | 1.3 | 4         |
| 29 | Composition and orbits of language operations: finiteness and upper bounds. International Journal of Computer Mathematics, 2013, 90, 1171-1196.                     | 1.8 | 3         |
| 30 | Abelian Primitive Words. Lecture Notes in Computer Science, 2011, , 204-215.  | 1.3 | 3         |
| 31 | Characterizing DNA Bond Shapes Using Trajectories. Lecture Notes in Computer Science, 2006, , 180-191.  | 1.3 | 2         |
| 32 | Decidability of Trajectory-Based Equations. Lecture Notes in Computer Science, 2004, , 723-734.   | 1.3 | 2         |
| 33 | Template-Guided Recombination: From Theory to Laboratory. Natural Computing Series, 2009, , 117-137.  | 2.2 | 2         |
| 34 | On codes defined by bio-operations. Theoretical Computer Science, 2007, 378, 3-16.  | 0.9 | 1         |
| 35 | Pairwise Rational Kernels Obtained by Automaton Operations. Lecture Notes in Computer Science, 2014, , 332-345.   | 1.3 | 1         |
| 36 | On Language Decompositions and Primality. Lecture Notes in Computer Science, 2011, , 63-75.   | 1.3 | 1         |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 37 | Algebraic properties of substitution on trajectories. Theoretical Computer Science, 2006, 369, 183-196.             | 0.9 | 0         |
| 38 | Identification of significantly mutated subnetworks in the breast cancer genome. Scientific Reports, 2021, 11, 642. | 3.3 | 0         |
| 39 | On Codes Defined by Bio-operations. Lecture Notes in Computer Science, 2004, , 127-138.                             | 1.3 | 0         |