## Lukasz B Huminiecki

List of Publications by Year in descending order

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| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Virtual Gene Concept and a Corresponding Pragmatic Research Program in Genetical Data Science.<br>Entropy, 2022, 24, 17.   | 2.2  | 0         |
| 2  | A Contemporary Message from Mendel's Logical Empiricism. BioEssays, 2020, 42, e2000120.  | 2.5  | 5         |
| 3  | Models of the Gene Must Inform Data-Mining Strategies in Genomics. Entropy, 2020, 22, 942.   | 2.2  | 5         |
| 4  | Etiology of atherosclerosis informs choice of animal models and tissues for initial functional genomic studies of resveratrol. Pharmacological Research, 2020, 156, 104598.  | 7.1  | 6         |
| 5  | Molecular neuroscience at its "high― bibliometric analysis of the most cited papers on<br>endocannabinoid system, cannabis and cannabinoids. Journal of Cannabis Research, 2019, 1, 4.                                       | 3.2  | 7         |
| 6  | Magic roundabout is an endothelial-specific ohnolog of ROBO1 which neo-functionalized to an essential new role in angiogenesis. PLoS ONE, 2019, 14, e0208952.  | 2.5  | 7         |
| 7  | Ethnopharmacological Applications Targeting Alcohol Abuse: Overview and Outlook. Frontiers in Pharmacology, 2019, 10, 1593.  | 3.5  | 10        |
| 8  | The functional genomic studies of resveratrol in respect to its anti-cancer effects. Biotechnology<br>Advances, 2018, 36, 1699-1708.   | 11.7 | 49        |
| 9  | Vascular smooth muscle cell proliferation as a therapeutic target. Part 2: Natural products inhibiting proliferation. Biotechnology Advances, 2018, 36, 1608-1621.   | 11.7 | 38        |
| 10 | Vascular smooth muscle cell proliferation as a therapeutic target. Part 1: molecular targets and pathways. Biotechnology Advances, 2018, 36, 1586-1607.  | 11.7 | 78        |
| 11 | Pecan nuts: A review of reported bioactivities and health effects. Trends in Food Science and Technology, 2018, 71, 246-257.   | 15.1 | 97        |
| 12 | Modelling of the breadth of expression from promoter architectures identifies pro-housekeeping transcription factors. PLoS ONE, 2018, 13, e0198961.  | 2.5  | 4         |
| 13 | Ethnopharmacological Approaches for Dementia Therapy and Significance of Natural Products and<br>Herbal Drugs. Frontiers in Aging Neuroscience, 2018, 10, 3.   | 3.4  | 93        |
| 14 | Nutrients Composition in Fit Snacks Made from Ostrich, Beef and Chicken Dried Meat. Molecules, 2018, 23, 1267.   | 3.8  | 16        |
| 15 | Can We Predict Gene Expression by Understanding Proximal Promoter Architecture?. Trends in Biotechnology, 2017, 35, 530-546.   | 9.3  | 18        |
| 16 | The functional genomic studies of curcumin. Seminars in Cancer Biology, 2017, 46, 107-118.   | 9.6  | 61        |
| 17 | Ethnopharmacological Approaches for Therapy of Jaundice: Part I. Frontiers in Pharmacology, 2017, 8, 518.  | 3.5  | 23        |
| 18 | Ethnopharmacological Approaches for Therapy of Jaundice: Part II. Highly Used Plant Species from<br>Acanthaceae, Euphorbiaceae, Asteraceae, Combretaceae, and Fabaceae Families. Frontiers in<br>Pharmacology, 2017, 8, 519. | 3.5  | 27        |

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|----|--|------|-----------|
| 19 | The Constrained Maximal Expression Level Owing to Haploidy Shapes Gene Content on the Mammalian<br>X Chromosome. PLoS Biology, 2015, 13, e1002315.   | 5.6  | 32        |
| 20 | A simple metric of promoter architecture robustly predicts expression breadth of human genes suggesting that most transcription factors are positive regulators. Genome Biology, 2014, 15, 413.                                | 8.8  | 20        |
| 21 | Differential roles of epigenetic changes and Foxp3 expression in regulatory T cell-specific transcriptional regulation. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 5289-5294. | 7.1  | 111       |
| 22 | A promoter-level mammalian expression atlas. Nature, 2014, 507, 462-470.   | 27.8 | 1,838     |
| 23 | Polyploidy and the Evolution of Complex Traits. International Journal of Evolutionary Biology, 2012, 2012, 1-12.   | 1.0  | 19        |
| 24 | 2R and remodeling of vertebrate signal transduction engine. BMC Biology, 2010, 8, 146.   | 3.8  | 77        |
| 25 | Emergence, development and diversification of the TGF-β signalling pathway within the animal kingdom.<br>BMC Evolutionary Biology, 2009, 9, 28.  | 3.2  | 137       |
| 26 | The transcriptional network that controls growth arrest and differentiation in a human myeloid leukemia cell line. Nature Genetics, 2009, 41, 553-562.   | 21.4 | 408       |
| 27 | Pseudo–Messenger RNA: Phantoms of the Transcriptome. PLoS Genetics, 2006, 2, e23.  | 3.5  | 58        |
| 28 | Clusters of Co-expressed Genes in Mammalian Genomes Are Conserved by Natural Selection.<br>Molecular Biology and Evolution, 2005, 22, 767-775.   | 8.9  | 154       |
| 29 | The Transcriptional Landscape of the Mammalian Genome. Science, 2005, 309, 1559-1563.  | 12.6 | 3,227     |
| 30 | Divergence of Spatial Gene Expression Profiles Following Species-Specific Gene Duplications in Human and Mouse. Genome Research, 2004, 14, 1870-1879.  | 5.5  | 139       |
| 31 | Congruence of tissue expression profiles from Gene Expression Atlas, SAGEmap and TissueInfo databases. BMC Genomics, 2003, 4, 31.  | 2.8  | 77        |
| 32 | EndoPDI, a Novel Protein-disulfide Isomerase-like Protein That Is Preferentially Expressed in<br>Endothelial Cells Acts as a Stress Survival Factor. Journal of Biological Chemistry, 2003, 278,<br>47079-47088.               | 3.4  | 149       |
| 33 | Magic Roundabout Is a New Member of the Roundabout Receptor Family That Is Endothelial Specific and Expressed at Sites of Active Angiogenesis. Genomics, 2002, 79, 547-552.  | 2.9  | 299       |
| 34 | The Ensembl genome database project. Nucleic Acids Research, 2002, 30, 38-41.  | 14.5 | 1,411     |
| 35 | Vascular endothelial growth factor transgenic mice exhibit reduced male fertility and placental rejection. Molecular Human Reproduction, 2001, 7, 255-264.   | 2.8  | 26        |
| 36 | In Silico Cloning of Novel Endothelial-Specific Genes. Genome Research, 2000, 10, 1796-1806.   | 5.5  | 130       |