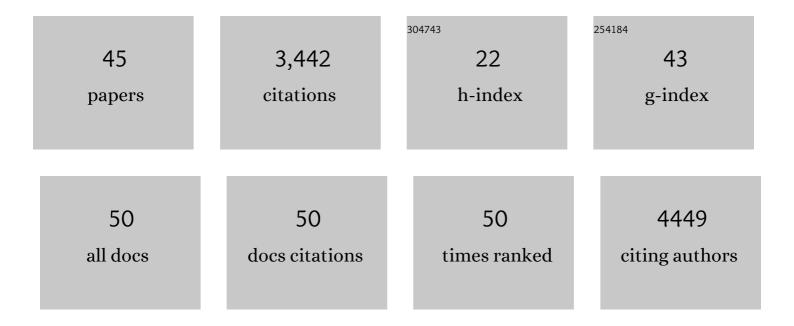
Gabriel V Markov

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6494925/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Genome sequence of the metazoan plant-parasitic nematode Meloidogyne incognita. Nature Biotechnology, 2008, 26, 909-915. | 17.5 | 1,012 |
| 2 | The Ectocarpus genome and the independent evolution of multicellularity in brown algae. Nature, 2010, 465, 617-621. | 27.8 | 774 |
| 3 | Genome structure and metabolic features in the red seaweed <i>Chondrus crispus</i> shed light on evolution of the Archaeplastida. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5247-5252. | 7.1 | 307 |
| 4 | Independent elaboration of steroid hormone signaling pathways in metazoans. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 11913-11918. | 7.1 | 163 |
| 5 | Origin and evolution of the ligand-binding ability of nuclear receptors. Molecular and Cellular Endocrinology, 2011, 334, 21-30. | 3.2 | 90 |
| 6 | Traceability, reproducibility and wiki-exploration for "Ã-la-carte―reconstructions of genome-scale metabolic models. PLoS Computational Biology, 2018, 14, e1006146. | 3.2 | 89 |
| 7 | Plastid genomes of two brown algae, Ectocarpus siliculosus and Fucus vesiculosus: further insights on the evolution of red-algal derived plastids. BMC Evolutionary Biology, 2009, 9, 253. | 3.2 | 77 |
| 8 | The amphioxus genome enlightens the evolution of the thyroid hormone signaling pathway. Development Genes and Evolution, 2008, 218, 667-680. | 0.9 | 59 |
| 9 | The Nuclear Hormone Receptor NHR-40 Acts Downstream of the Sulfatase EUD-1 as Part of a Developmental Plasticity Switch in Pristionchus. Current Biology, 2016, 26, 2174-2179. | 3.9 | 56 |
| 10 | Origin of an ancient hormone/receptor couple revealed by resurrection of an ancestral estrogen. Science Advances, 2017, 3, e1601778. | 10.3 | 49 |
| 11 | Ventx Factors Function as Nanog-Like Guardians of Developmental Potential in Xenopus. PLoS ONE, 2012, 7, e36855. | 2.5 | 48 |
| 12 | Natural Variation in Dauer Pheromone Production and Sensing Supports Intraspecific Competition in Nematodes. Current Biology, 2014, 24, 1536-1541. | 3.9 | 47 |
| 13 | The evolution of the ligand/receptor couple: A long road from comparative endocrinology to comparative genomics. Molecular and Cellular Endocrinology, 2008, 293, 5-16. | 3.2 | 43 |
| 14 | Genome and metabolic network of ââ,¬Å"Candidatus Phaeomarinobacter ectocarpiââ,¬Â•Ec32, a new candidate genus of Alphaproteobacteria frequently associated with brown algae. Frontiers in Genetics, 2014, 5, 241. | 2.3 | 43 |
| 15 | Chondrus crispus – A Present and Historical Model Organism for Red Seaweeds. Advances in Botanical Research, 2014, 71, 53-89. | 1.1 | 37 |
| 16 | Ancient gene duplications have shaped developmental stage-specific expression in Pristionchus pacificus. BMC Evolutionary Biology, 2015, 15, 185. | 3.2 | 36 |
| 17 | Draft Genome of the Scarab Beetle <i>Oryctes borbonicus</i> on La Réunion Island. Genome Biology and Evolution, 2016, 8, 2093-2105. | 2.5 | 35 |
| 18 | Functional Conservation and Divergence ofdaf-22Paralogs inPristionchus pacificusDauer Development. Molecular Biology and Evolution, 2016, 33, 2506-2514. | 8.9 | 34 |

GABRIEL V MARKOV

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 19 | Evolution of Nuclear Receptors and Ligand Signaling. Current Topics in Developmental Biology, 2017, 125, 1-38. | 2.2 | 34 |
| 20 | Linking Genomic and Metabolomic Natural Variation Uncovers Nematode Pheromone Biosynthesis. Cell Chemical Biology, 2018, 25, 787-796.e12. | 5.2 | 31 |
| 21 | Biological rhythms in the deep-sea hydrothermal mussel Bathymodiolus azoricus. Nature Communications, 2020, 11, 3454. | 12.8 | 30 |
| 22 | The Role of DAF-21/Hsp90 in Mouth-Form Plasticity in Pristionchus pacificus. Molecular Biology and Evolution, 2017, 34, 1644-1653. | 8.9 | 28 |
| 23 | The genome of Ectocarpus subulatus – A highly stress-tolerant brown alga. Marine Genomics, 2020, 52, 100740. | 1.1 | 26 |
| 24 | Metabolic Complementarity Between a Brown Alga and Associated Cultivable Bacteria Provide Indications of Beneficial Interactions. Frontiers in Marine Science, 2020, 7, . | 2.5 | 25 |
| 25 | Evolution of Nuclear Retinoic Acid Receptor Alpha (RARÂ) Phosphorylation Sites. Serine Gain Provides Fine-Tuned Regulation. Molecular Biology and Evolution, 2011, 28, 2125-2137. | 8.9 | 23 |
| 26 | The Same or Not the Same: Lineage-Specific Gene Expansions and Homology Relationships in Multigene Families in Nematodes. Journal of Molecular Evolution, 2015, 80, 18-36. | 1.8 | 23 |
| 27 | On the Origin and Evolutionary History of NANOG. PLoS ONE, 2014, 9, e85104. | 2.5 | 21 |
| 28 | qPCR-based relative quantification of the brown algal endophyte Laminarionema elsbetiae in Saccharina latissima: variation and dynamics of host—endophyte interactions. Journal of Applied Phycology, 2018, 30, 2901-2911. | 2.8 | 19 |
| 29 | Genome–Scale Metabolic Networks Shed Light on the Carotenoid Biosynthesis Pathway in the Brown Algae Saccharina japonica and Cladosiphon okamuranus. Antioxidants, 2019, 8, 564. | 5.1 | 19 |
| 30 | The Ectocarpus Genome and Brown Algal Genomics. Advances in Botanical Research, 2012, 64, 141-184. | 1.1 | 18 |
| 31 | Diversity and evolution of cytochromes P450 in stramenopiles. Planta, 2019, 249, 647-661. | 3.2 | 18 |
| 32 | NR3E receptors in cnidarians: A new family of steroid receptor relatives extends the possible mechanisms for ligand binding. Journal of Steroid Biochemistry and Molecular Biology, 2018, 184, 11-19. | 2.5 | 17 |
| 33 | The "street light syndromeâ€, or how protein taxonomy can bias experimental manipulations. BioEssays, 2008, 30, 349-357. | 2.5 | 16 |
| 34 | Herbivore-induced chemical and molecular responses of the kelps Laminaria digitata and Lessonia spicata. PLoS ONE, 2017, 12, e0173315. | 2.5 | 16 |
| 35 | Inferring Biochemical Reactions and Metabolite Structures to Understand Metabolic Pathway Drift. IScience, 2020, 23, 100849. | 4.1 | 15 |
| 36 | In Silico Survey of the Mitochondrial Protein Uptake and Maturation Systems in the Brown Alga Ectocarpus siliculosus. PLoS ONE, 2011, 6, e19540. | 2.5 | 10 |

GABRIEL V MARKOV

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 37 | Hormonally active phytochemicals from macroalgae: A largely untapped source of ligands to deorphanize nuclear receptors in emerging marine animal models. General and Comparative Endocrinology, 2018, 265, 41-45. | 1.8 | 8 |
| 38 | A structural signature motif enlightens the origin and diversification of nuclear receptors. PLoS Genetics, 2021, 17, e1009492. | 3.5 | 8 |
| 39 | Small molecules as products of evolution. Current Biology, 2022, 32, R100-R105. | 3.9 | 6 |
| 40 | The Evolution of Novelty in Conserved Gene Families. International Journal of Evolutionary Biology, 2012, 2012, 1-8. | 1.0 | 5 |
| 41 | Semi-Quantitative Targeted Gas Chromatography-Mass Spectrometry Profiling Supports a Late Side-Chain Reductase Cycloartenol-to-Cholesterol Biosynthesis Pathway in Brown Algae. Frontiers in Plant Science, 2021, 12, 648426. | 3.6 | 5 |
| 42 | Different Early Responses of Laminariales to an Endophytic Infection Provide Insights About Kelp Host Specificity. Frontiers in Marine Science, 2021, 8, . | 2.5 | 5 |
| 43 | What does Evolution Teach us about Nuclear Receptors?. , 2010, , 15-29. | | 4 |
| 44 | Independent Evolution of the MYB Family in Brown Algae. Frontiers in Genetics, 2021, 12, 811993. | 2.3 | 3 |
| 45 | Evolution of Hormonal Mechanisms. , 2019, , 16-22. | | Ο |