

Associa€prof Eivind A B Undheim

List of Publications by Year in descending order

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Version: 2024-02-01

86
papers

3,826
citations

117625

34
h-index

144013

57
g-index

90
all docs

90
docs citations

90
times ranked

3204
citing authors

#	ARTICLE	IF	CITATIONS
1	Selective spider toxins reveal a role for the Nav1.1 channel in mechanical pain. <i>Nature</i> , 2016, 534, 494-499.	27.8	239
2	Discovery of a selective Na ^v 1.7 inhibitor from centipede venom with analgesic efficacy exceeding morphine in rodent pain models. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 17534-17539.	7.1	164
3	Production of Recombinant Disulfide-Rich Venom Peptides for Structural and Functional Analysis via Expression in the Periplasm of <i>E. coli</i> . <i>PLoS ONE</i> , 2013, 8, e63865.	2.5	140
4	The Diversity of Venom: The Importance of Behavior and Venom System Morphology in Understanding Its Ecology and Evolution. <i>Toxins</i> , 2019, 11, 666.	3.4	135
5	Intraspecific venom variation in the medically significant Southern Pacific Rattlesnake (<i>Crotalus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 99, 68-83.	2.4	114
6	Three-Fingered RAVERs: Rapid Accumulation of Variations in Exposed Residues of Snake Venom Toxins. <i>Toxins</i> , 2013, 5, 2172-2208.	3.4	111
7	On the venom system of centipedes (Chilopoda), a neglected group of venomous animals. <i>Toxicon</i> , 2011, 57, 512-524.	1.6	110
8	Selective Na ^v 1.1 activation rescues Dravet syndrome mice from seizures and premature death. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E8077-E8085.	7.1	105
9	Clawing through Evolution: Toxin Diversification and Convergence in the Ancient Lineage Chilopoda (Centipedes). <i>Molecular Biology and Evolution</i> , 2014, 31, 2124-2148.	8.9	100
10	Convergent evolution of pain-inducing defensive venom components in spitting cobras. <i>Science</i> , 2021, 371, 386-390.	12.6	96
11	Centipede Venom: Recent Discoveries and Current State of Knowledge. <i>Toxins</i> , 2015, 7, 679-704.	3.4	84
12	Differential Evolution and Neofunctionalization of Snake Venom Metalloprotease Domains. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 651-663.	3.8	83
13	Spider venomomics: implications for drug discovery. <i>Future Medicinal Chemistry</i> , 2014, 6, 1699-1714.	2.3	81
14	Venom peptides as therapeutics: advances, challenges and the future of venom-peptide discovery. <i>Expert Review of Proteomics</i> , 2017, 14, 931-939.	3.0	81
15	Evolution Stings: The Origin and Diversification of Scorpion Toxin Peptide Scaffolds. <i>Toxins</i> , 2013, 5, 2456-2487.	3.4	79
16	Toxin structures as evolutionary tools: Using conserved 3D folds to study the evolution of rapidly evolving peptides. <i>BioEssays</i> , 2016, 38, 539-548.	2.5	76
17	A comprehensive portrait of the venom of the giant red bull ant, <i>Myrmecia gulosa</i> , reveals a hyperdiverse hymenopteran toxin gene family. <i>Science Advances</i> , 2018, 4, eaau4640.	10.3	69
18	A Proteomics and Transcriptomics Investigation of the Venom from the Barychelid Spider <i>Trittame loki</i> (Brush-Foot Trapdoor). <i>Toxins</i> , 2013, 5, 2488-2503.	3.4	68

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19	The assassin bug <i>Pristhesancus plagipennis</i> produces two distinct venoms in separate gland lumens. <i>Nature Communications</i> , 2018, 9, 755.	12.8	67
20	Weaponization of a Hormone: Convergent Recruitment of Hyperglycemic Hormone into the Venom of Arthropod Predators. <i>Structure</i> , 2015, 23, 1283-1292.	3.3	66
21	Revisiting venom of the sea anemone <i>Stichodactyla haddoni</i> : Omics techniques reveal the complete toxin arsenal of a well-studied sea anemone genus. <i>Journal of Proteomics</i> , 2017, 166, 83-92.	2.4	64
22	Molecular Phylogeny and Evolution of the Proteins Encoded by Coleoid (Cuttlefish, Octopus, and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.8	62
23	Dracula's children: Molecular evolution of vampire bat venom. <i>Journal of Proteomics</i> , 2013, 89, 95-111.	2.4	61
24	Structural venomomics reveals evolution of a complex venom by duplication and diversification of an ancient peptide-encoding gene. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 11399-11408.	7.1	59
25	Production and packaging of a biological arsenal: Evolution of centipede venoms under morphological constraint. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 4026-4031.	7.1	56
26	Molecular Evolution of Vertebrate Neurotrophins: Co-Option of the Highly Conserved Nerve Growth Factor Gene into the Advanced Snake Venom Arsenal. <i>PLoS ONE</i> , 2013, 8, e81827.	2.5	56
27	Venom proteomic characterization and relative antivenom neutralization of two medically important Pakistani elapid snakes (<i>Bungarus sindanus</i> and <i>Naja naja</i>). <i>Journal of Proteomics</i> , 2013, 89, 15-23.	2.4	55
28	Venom Down Under: Dynamic Evolution of Australian Elapid Snake Toxins. <i>Toxins</i> , 2013, 5, 2621-2655.	3.4	55
29	Sea Anemone Toxins: A Structural Overview. <i>Marine Drugs</i> , 2019, 17, 325.	4.6	54
30	Melt With This Kiss: Paralyzing and Liquefying Venom of The Assassin Bug <i>Pristhesancus plagipennis</i> (Hemiptera: Reduviidae). <i>Molecular and Cellular Proteomics</i> , 2017, 16, 552-566.	3.8	53
31	Squeezers and Leaf-cutters: Differential Diversification and Degeneration of the Venom System in Toxiciferan Reptiles. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 1881-1899.	3.8	52
32	Diversification of a single ancestral gene into a successful toxin superfamily in highly venomous Australian funnel-web spiders. <i>BMC Genomics</i> , 2014, 15, 177.	2.8	49
33	A process of convergent amplification and tissue-specific expression dominates the evolution of toxin-like genes in sea anemones. <i>Molecular Ecology</i> , 2019, 28, 2272-2289.	3.9	48
34	Solenodon genome reveals convergent evolution of venom in eulipotyphlan mammals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 25745-25755.	7.1	42
35	True Lies: Using Proteomics to Assess the Accuracy of Transcriptome-Based Venomomics in Centipedes Uncovers False Positives and Reveals Startling Intraspecific Variation in <i>Scolopendra subspinipes</i> . <i>Toxins</i> , 2018, 10, 96.	3.4	40
36	Multifunctional warheads: Diversification of the toxin arsenal of centipedes via novel multidomain transcripts. <i>Journal of Proteomics</i> , 2014, 102, 1-10.	2.4	36

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37	Mapping Enzyme Activity on Tissue by Functional Mass Spectrometry Imaging. <i>Angewandte Chemie - International Edition</i> , 2020, 59, 3855-3858.	13.8	35
38	PHAB toxins: a unique family of predatory sea anemone toxins evolving via intra-gene concerted evolution defines a new peptide fold. <i>Cellular and Molecular Life Sciences</i> , 2018, 75, 4511-4524.	5.4	34
39	A Dipteran's Novel Sucker Punch: Evolution of Arthropod Atypical Venom with a Neurotoxic Component in Robber Flies (Asilidae, Diptera). <i>Toxins</i> , 2018, 10, 29.	3.4	33
40	Australian funnel-web spiders evolved human-lethal δ -hexatoxins for defense against vertebrate predators. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 24920-24928.	7.1	32
41	Proteomic comparison of <i>Hypnale hypnale</i> (Hump-Nosed Pit-Viper) and <i>Calloselasma rhodostoma</i> (Malayan Pit-Viper) venoms. <i>Journal of Proteomics</i> , 2013, 91, 338-343.	2.4	30
42	Venom on ice: First insights into Antarctic octopus venoms. <i>Toxicon</i> , 2010, 56, 897-913.	1.6	28
43	Centipede venoms as a source of drug leads. <i>Expert Opinion on Drug Discovery</i> , 2016, 11, 1139-1149.	5.0	28
44	Characterising Functional Venom Profiles of Anthozoans and Medusozoans within Their Ecological Context. <i>Marine Drugs</i> , 2020, 18, 202.	4.6	28
45	Genetic identification of Southern Ocean octopod samples using mtCOI. <i>Comptes Rendus - Biologies</i> , 2010, 333, 395-404.	0.2	27
46	Three Peptide Modulators of the Human Voltage-Gated Sodium Channel 1.7, an Important Analgesic Target, from the Venom of an Australian Tarantula. <i>Toxins</i> , 2015, 7, 2494-2513.	3.4	27
47	Venomomics of Remipede Crustaceans Reveals Novel Peptide Diversity and Illuminates the Venom's Biological Role. <i>Toxins</i> , 2017, 9, 234.	3.4	27
48	A complicated complex: Ion channels, voltage sensing, cell membranes and peptide inhibitors. <i>Neuroscience Letters</i> , 2018, 679, 35-47.	2.1	27
49	Structure-Activity Relationship of Chlorotoxin-Like Peptides. <i>Toxins</i> , 2016, 8, 36.	3.4	26
50	Deep venomomics of the <i>Pseudonaja</i> genus reveals inter- and intra-specific variation. <i>Journal of Proteomics</i> , 2016, 133, 20-32.	2.4	26
51	Deadly Proteomes: A Practical Guide to Proteotranscriptomics of Animal Venoms. <i>Proteomics</i> , 2020, 20, e1900324.	2.2	26
52	Toxins from scratch? Diverse, multimodal gene origins in the predatory robber fly <i>Dasypogon diadema</i> indicate a dynamic venom evolution in dipteran insects. <i>GigaScience</i> , 2019, 8, .	6.4	25
53	Parallel Evolution of Complex Centipede Venoms Revealed by Comparative Proteotranscriptomic Analyses. <i>Molecular Biology and Evolution</i> , 2019, 36, 2748-2763.	8.9	24
54	Modulation of Ion Channels by Cysteine-Rich Peptides. <i>Advances in Pharmacology</i> , 2017, 79, 199-223.	2.0	22

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55	Extreme venom variation in Middle Eastern vipers: A proteomics comparison of <i>Eristicophis macmahonii</i> , <i>Pseudocerastes fieldi</i> and <i>Pseudocerastes persicus</i> . <i>Journal of Proteomics</i> , 2015, 116, 106-113.	2.4	21
56	Mutual enlightenment: A toolbox of concepts and methods for integrating evolutionary and clinical toxinology via snake venomomics and the contextual stance. <i>Toxicon: X</i> , 2021, 9-10, 100070.	2.9	21
57	Venoms for all occasions: The functional toxin profiles of different anatomical regions in sea anemones are related to their ecological function. <i>Molecular Ecology</i> , 2022, 31, 866-883.	3.9	21
58	The Use of Imaging Mass Spectrometry to Study Peptide Toxin Distribution in Australian Sea Anemones. <i>Australian Journal of Chemistry</i> , 2017, 70, 1235.	0.9	20
59	Structural and Molecular Diversification of the Anguimorpha Lizard Mandibular Venom Gland System in the Arboreal Species <i>Abronia graminea</i> . <i>Journal of Molecular Evolution</i> , 2012, 75, 168-183.	1.8	19
60	A selective NaV1.1 activator with potential for treatment of Dravet syndrome epilepsy. <i>Biochemical Pharmacology</i> , 2020, 181, 113991.	4.4	19
61	Functional characterization on invertebrate and vertebrate tissues of tachykinin peptides from octopus venoms. <i>Peptides</i> , 2013, 47, 71-76.	2.4	18
62	An Integrated Proteomic and Transcriptomic Analysis Reveals the Venom Complexity of the Bullet Ant <i>Paraponera clavata</i> . <i>Toxins</i> , 2020, 12, 324.	3.4	18
63	Comparative analyses of glycerotoxin expression unveil a novel structural organization of the bloodworm venom system. <i>BMC Evolutionary Biology</i> , 2017, 17, 64.	3.2	17
64	Can we resolve the taxonomic bias in spider venom research?. <i>Toxicon: X</i> , 2019, 1, 100005.	2.9	17
65	A Centipede Toxin Family Defines an Ancient Class of CS β Defensins. <i>Structure</i> , 2019, 27, 315-326.e7.	3.3	17
66	Missiles of Mass Disruption: Composition and Glandular Origin of Venom Used as a Projectile Defensive Weapon by the Assassin Bug <i>Platymeris rhadamanthus</i> . <i>Toxins</i> , 2019, 11, 673.	3.4	16
67	Secreted Cysteine-Rich Repeat Proteins (SCREPs): A Novel Multi-Domain Architecture. <i>Frontiers in Pharmacology</i> , 2018, 9, 1333.	3.5	15
68	Phylogenetic analyses suggest centipede venom arsenals were repeatedly stocked by horizontal gene transfer. <i>Nature Communications</i> , 2021, 12, 818.	12.8	15
69	A peptide toxin in ant venom mimics vertebrate EGF-like hormones to cause long-lasting hypersensitivity in mammals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	15
70	CHAPTER 1. Seeing the Woods for the Trees: Understanding Venom Evolution as a Guide for Biodiscovery. <i>RSC Drug Discovery Series</i> , 2015, , 1-36.	0.3	13
71	Vintage venoms: Proteomic and pharmacological stability of snake venoms stored for up to eight decades. <i>Journal of Proteomics</i> , 2014, 105, 285-294.	2.4	12
72	Tentacle Morphological Variation Coincides with Differential Expression of Toxins in Sea Anemones. <i>Toxins</i> , 2021, 13, 452.	3.4	12

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73	Multipurpose peptides: The venoms of Amazonian stinging ants contain anthelmintic ponericins with diverse predatory and defensive activities. <i>Biochemical Pharmacology</i> , 2021, 192, 114693.	4.4	10
74	A Versatile and Robust Serine Protease Inhibitor Scaffold from <i>Actinia tenebrosa</i> . <i>Marine Drugs</i> , 2019, 17, 701.	4.6	9
75	Mapping Enzyme Activity on Tissue by Functional Mass Spectrometry Imaging. <i>Angewandte Chemie</i> , 2020, 132, 3883-3886.	2.0	8
76	Evolution, Expression Patterns, and Distribution of Novel Ribbon Worm Predatory and Defensive Toxins. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	8
77	CHAPTER 3. Venoms-Based Drug Discovery: Proteomic and Transcriptomic Approaches. <i>RSC Drug Discovery Series</i> , 2015, , 80-96.	0.3	7
78	A pain-causing and paralytic ant venom glycopeptide. <i>IScience</i> , 2021, 24, 103175.	4.1	7
79	Natural history of the slaty grey snake (<i>Stegonotus cucullatus</i>) (Serpentes:Colubridae) from tropical north Queensland, Australia. <i>Australian Journal of Zoology</i> , 2009, 57, 119.	1.0	7
80	Multitarget nociceptor sensitization by a promiscuous peptide from the venom of the King Baboon spider. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	7
81	Editorial: Animal Toxins as Comprehensive Pharmacological Tools to Identify Diverse Ion Channels. <i>Frontiers in Pharmacology</i> , 2019, 10, 423.	3.5	6
82	AsKC11, a Kunitz Peptide from <i>Anemonia sulcata</i> , Is a Novel Activator of G Protein-Coupled Inward-Rectifier Potassium Channels. <i>Marine Drugs</i> , 2022, 20, 140.	4.6	6
83	Physiological constraints dictate toxin spatial heterogeneity in snake venom glands. <i>BMC Biology</i> , 2022, 20, .	3.8	4
84	Differential Evolution and Neofunctionalization of Snake Venom Metalloprotease Domains. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 1488.	3.8	1
85	Biochemical Modulation of Venom by Spiders is Achieved Via Compartmentalized Toxin Production and Storage. <i>SSRN Electronic Journal</i> , 0, , .	0.4	1
86	A Centipede Toxin Family Defines a New Ancient Class of CSSS Defensins. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0