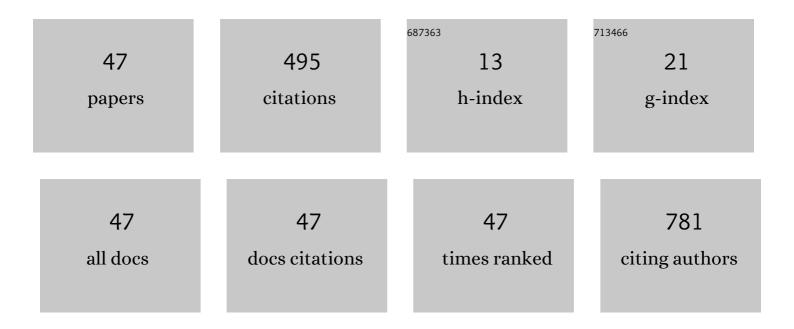
Woo-Jin Kim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2836180/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Genome sequence of pacific abalone (Haliotis discus hannai): the first draft genome in family Haliotidae. GigaScience, 2017, 6, 1-8.	6.4	84
2	Distinct immune tones are established by Lactococcus lactis BFE920 and Lactobacillus plantarum FGL0001 in the gut of olive flounder (Paralichthys olivaceus). Fish and Shellfish Immunology, 2016, 55, 434-443.	3.6	41
3	Twelve quick steps for genome assembly and annotation in the classroom. PLoS Computational Biology, 2020, 16, e1008325.	3.2	34
4	Antimicrobial Activity of Peptides Derived from Olive Flounder Lipopolysaccharide Binding Protein/Bactericidal Permeability-Increasing Protein (LBP/BPI). Marine Drugs, 2014, 12, 5240-5257.	4.6	25
5	Molecular characterization of a tandem-repeat galectin-9 (RuGlec9) from Korean rose bitterling (Rhodeus uyekii). Fish and Shellfish Immunology, 2012, 32, 939-944.	3.6	22
6	Optimizing Hybrid de Novo Transcriptome Assembly and Extending Genomic Resources for Giant Freshwater Prawns (Macrobrachium rosenbergii): The Identification of Genes and Markers Associated with Reproduction. International Journal of Molecular Sciences, 2016, 17, 690.	4.1	20
7	Development of Type I Genetic Markers from Expressed Sequence Tags in Highly Polymorphic Species. Marine Biotechnology, 2011, 13, 127-132.	2.4	19
8	Characterization of the flounder IL-6 promoter and its regulation by the p65 NF-κB subunit. Fish and Shellfish Immunology, 2010, 28, 961-964.	3.6	18
9	Mitochondrial DNA sequence analysis from multiple gene fragments reveals genetic heterogeneity of Crassostrea ariakensis in East Asia. Genes and Genomics, 2014, 36, 611-624.	1.4	18
10	A cDNA microarray analysis to identify genes involved in the acute-phase response pathway of the olive flounder after infection with Edwardsiella tarda. Veterinary Immunology and Immunopathology, 2014, 161, 49-56.	1.2	16
11	Cloning and characterization of hypusine-containing protein elF5A from the olive flounder Paralichthys olivaceus. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2009, 153, 281-287.	1.6	15
12	Microarray analysis of gene expression in olive flounder liver infected with viral haemorrhagic septicaemia virus (VHSV). Fish and Shellfish Immunology, 2016, 49, 66-78.	3.6	15
13	Molecular characterization and expression analysis of a peroxiredoxin 1 cDNA from Korean rose bitterling (Rhodeus uyekii). Molecular Biology Reports, 2014, 41, 2363-2370.	2.3	14
14	Gene expression profiles alteration after infection of virus, bacteria, and parasite in the Olive flounder (Paralichthys olivaceus). Scientific Reports, 2018, 8, 18065.	3.3	14
15	Molecular Characterization of Paralichthys olivaceus MAF1 and Its Potential Role as an Anti-Viral Hemorrhagic Septicaemia Virus Factor in Hirame Natural Embryo Cells. International Journal of Molecular Sciences, 2021, 22, 1353.	4.1	14
16	Effects of food availability on growth performance and immune-related gene expression of juvenile olive flounder (Paralichthys olivaceus). Fish and Shellfish Immunology, 2018, 80, 348-356.	3.6	12
17	Characterization, Expression Profile, and Promoter Analysis of the Rhodeus uyekii Vitellogenin Ao1 Gene. International Journal of Molecular Sciences, 2014, 15, 18804-18818.	4.1	11
18	Evolutionary conservation and expression of miR - 10a - 3p in olive flounder and rock bream. Gene, 2017, 628, 16-23.	2.2	11

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19	Characterization of the Long Terminal Repeat of the Endogenous Retrovirus-derived microRNAs in the Olive Flounder. Scientific Reports, 2019, 9, 14007.	3.3	11
20	Draft Genome Sequence of Kocuria rhizophila P7-4. Journal of Bacteriology, 2011, 193, 4286-4287.	2.2	10
21	Antimicrobial and Antitumor Activities of Novel Peptides Derived from the Lipopolysaccharide- and β-1,3-Glucan Binding Protein of the Pacific Abalone Haliotis discus hannai. Marine Drugs, 2016, 14, 227.	4.6	10
22	RNA-Seq-based transcriptome analysis of Korean rose bitterling (Rhodeus uyekii) exposed to synthetic estrogen 17-α-ethinylestradiol (EE2). Marine Genomics, 2015, 24, 233-236.	1.1	8
23	Transcriptome profiling of olive flounder responses under acute and chronic heat stress. Genes and Genomics, 2021, 43, 151-159.	1.4	8
24	Molecular and Functional Characterization of Thioredoxin 1from Korean Rose Bitterling (Rhodeus) Tj ETQq0 0 0	rgBT /Ove 4.1	rloçk 10 Tf 50
25	Characterization of sexual size dimorphism and sex-biased genes expression profile in the olive flounder. Molecular Biology Reports, 2020, 47, 8317-8324.	2.3	7
26	Genomic cloning and promoter analysis of the β-actin gene from Korean rose bitterling (Rhodeus) Tj ETQqO 0 0	rgBT/Ove 1.4	rloçk 10 Tf 50
27	A comparison of oneâ€slope straight brokenâ€line, twoâ€slope straight brokenâ€line, quadratic brokenâ€line and quadratic models to estimate an accurate optimum feeding rate for juvenile olive flounder () Tj ETQq1 1 0.7	'84 317 4 rgE	3T /Dverlock]
28	Development of 81 new polymorphic EST-derived microsatellite markers for the olive flounder, ParalichthysÂolivaceus. Conservation Genetics, 2009, 10, 1105-1111.	1.5	3
29	Development of 52 new polymorphic microsatellite markers for the olive flounder, <i>Paralichthys olivaceus</i> . Molecular Ecology Resources, 2009, 9, 839-842.	4.8	3
30	Molecular cloning and characterization of cathepsin F gene from olive flounder Paralichthys Olivaceus. Genes and Genomics, 2010, 32, 137-142.	1.4	3
31	PRIMER NOTE: Isolation and characterization of polymorphic microsatellite markers for the endangered Korean freshwater fish Hemibarbus mylodon. Molecular Ecology Notes, 2006, 7, 516-518.	1.7	2
32	Expression and promoter activity of endogenous retroviruses in the Olive flounder (Paralichthys) Tj ETQq0 0 0 rg	3BT/Qverlo 1.4	ock ₂ 10 Tf 50 2
33	EST-based Survey of Gene Expression in Seven Tissue Types from the Abalone Haliotis discus hannai. Journal of Fisheries Science and Technology, 2007, 10, 119-126.	0.2	2
34	Molecular cloning and characterization of PLCB1 (phospholipase C, beta 1) gene from the olive flounder, Paralichthys olivaceus. Genes and Genomics, 2011, 33, 701-709.	1.4	1
35	Isolation and inheritance of microsatellite loci for the oily bittering (Acheilognathus koreensis): applications for analysis of genetic diversity of wild populations. Animal Cells and Systems, 2012, 16, 321-328.	2.2	1
36	The GAL4 enhancer-trap line for analysis of definitive hematopoiesis in zebrafish. Animal Cells and Systems, 2015, 19, 96-100.	2.2	1

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37	Detection of LINE RT elements in the olive flounder (Paralichthys olivaceus) genome and expression analysis after infection with S. parauberis. Genes and Genomics, 2016, 38, 1105-1110.	1.4	1
38	Fish Myogenic Regulatory Protein LUC7L: Characterization and Expression Analysis in Korean Rose Bitterling (Rhodeus uyekii). Development & Reproduction, 2014, 18, 251-258.	0.5	1
39	The Regulatory Region of Muscle-Specific Alpha Actin 1 Drives Fluorescent Protein Expression in Olive Flounder Paralichthys olivaceus. Development & Reproduction, 2019, 23, 55-61.	0.4	1
40	Identification of differentially expressed genes in the developmental stages from olive flounderParalichthys olivaceususing an annealing control primer system. Animal Cells and Systems, 2010, 14, 25-30.	2.2	0
41	Molecular characterization and gene expression analysis of a metalloprotease from Pacific abalone <i>Haliotis discus hannai</i> . Animal Cells and Systems, 2014, 18, 267-274.	2.2	0
42	Molecular cloning and characterization of peroxisome proliferatorâ€activated receptorâ€Â¥Ã£ from olive flounder, Paralichthys olivaceus. FASEB Journal, 2008, 22, 198-198.	0.5	0
43	Molecular cloning and characterization of cathepsin F from olive flounder Paralichthys olivaceus. FASEB Journal, 2008, 22, 199-199.	0.5	0
44	Cloning of the autophagyâ€related gene Beclinâ€1 of the marine teleost Paralichthys olivaceus and its possible implication in viral replication. FASEB Journal, 2011, 25, 948.4.	0.5	0
45	GENE CLONING, PURIFICATION, AND CHARACTERIZATION OF A COLDâ€ADAPTED LIPASE FROM ACINETOBACTER SP. V28â€28. FASEB Journal, 2011, 25, 919.3.	0.5	0
46	Gene organizations of two types of the flounder warm temperature acclimationâ€related 65 kDa protein Wap65, and their gene expression profiles. FASEB Journal, 2013, 27, lb208.	0.5	0
47	Expression Analysis of the Caspase10 from Olive Flounder (Paralichthys olivaceus) against Viral Hemorrhagic Septicemia Virus (VHSV) Challenge. Development & Reproduction, 2020, 24, 187-196.	0.4	0