Tune Wulff

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6617071/publications.pdf

Version: 2024-02-01

414414 394421 1,179 46 19 32 citations h-index g-index papers 51 51 51 1748 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	A Consensus Genome-scale Reconstruction of Chinese Hamster Ovary Cell Metabolism. Cell Systems, 2016, 3, 434-443.e8.	6.2	205
2	Drug-Driven Phenotypic Convergence Supports Rational Treatment Strategies of Chronic Infections. Cell, 2018, 172, 121-134.e14.	28.9	131
3	Predictable tuning of protein expression in bacteria. Nature Methods, 2016, 13, 233-236.	19.0	116
4	Multiplex secretome engineering enhances recombinant protein production and purity. Nature Communications, 2020, $11,1908$.	12.8	63
5	Authentication of Fish Products by Large-Scale Comparison of Tandem Mass Spectra. Journal of Proteome Research, 2013, 12, 5253-5259.	3.7	46
6	Changes in Liver Proteome Expression of Senegalese Sole (Solea senegalensis) in Response to Repeated Handling Stress. Marine Biotechnology, 2012, 14, 714-729.	2.4	41
7	Model-guided dynamic control of essential metabolic nodes boosts acetyl-coenzyme A–dependent bioproduction in rewired Pseudomonas putida. Metabolic Engineering, 2021, 67, 373-386.	7.0	41
8	Nutritional mitigation of winter thermal stress in gilthead seabream: Associated metabolic pathways and potential indicators of nutritional state. Journal of Proteomics, 2016, 142, 1-14.	2.4	36
9	Reversibility of exercise-induced translocation of Na+-K+ pump subunits to the plasma membrane in rat skeletal muscle. Pflugers Archiv European Journal of Physiology, 2001, 443, 212-217.	2.8	34
10	Effects of Preslaughter Stress Levels on the Post-mortem Sarcoplasmic Proteomic Profile of Gilthead Seabream Muscle. Journal of Agricultural and Food Chemistry, 2012, 60, 9443-9453.	5.2	32
11	Long term anoxia in rainbow trout investigated by 2â€DE and MS/MS. Proteomics, 2008, 8, 1009-1018.	2.2	31
12	Dietary Tools To Modulate Glycogen Storage in Gilthead Seabream Muscle: Glycerol Supplementation. Journal of Agricultural and Food Chemistry, 2012, 60, 10613-10624.	5.2	31
13	Dietary Creatine Supplementation in Gilthead Seabream (Sparus aurata): Comparative Proteomics Analysis on Fish Allergens, Muscle Quality, and Liver. Frontiers in Physiology, 2018, 9, 1844.	2.8	31
14	Time-dependent changes in protein expression in rainbow trout muscle following hypoxia. Journal of Proteomics, 2012, 75, 2342-2351.	2.4	30
15	Influence of supplemental maslinic acid (olive-derived triterpene) on the post-mortem muscle properties and quality traits of gilthead seabream. Aquaculture, 2013, 396-399, 146-155.	3.5	30
16	Dietary Supplementation with Vitamin K Affects Transcriptome and Proteome of Senegalese Sole, Improving Larval Performance and Quality. Marine Biotechnology, 2014, 16, 522-537.	2.4	30
17	Reprogramming AA catabolism in CHO cells with CRISPR/Cas9 genome editing improves cell growth and reduces byproduct secretion. Metabolic Engineering, 2019, 56, 120-129.	7.0	22
18	Comparison of two anoxia models in rainbow trout cells by a 2â€DE and MS/MSâ€based proteome approach. Proteomics, 2008, 8, 2035-2044.	2,2	21

#	Article	IF	CITATIONS
19	Industrializing a Bacterial Strain for <scp>l</scp> -Serine Production through Translation Initiation Optimization. ACS Synthetic Biology, 2019, 8, 2347-2358.	3.8	21
20	Adaptation of hydroxymethylbutenyl diphosphate reductase enables volatile isoprenoid production. ELife, 2020, 9, .	6.0	19
21	Regulation of the mitogen-activated protein kinase p44 ERK activity during anoxia/recovery in rainbow trout hypodermal fibroblasts. Journal of Experimental Biology, 2006, 209, 1765-1776.	1.7	17
22	Fluctuations in glucose availability prevent global proteome changes and physiological transition during prolonged chemostat cultivations of Saccharomyces cerevisiae. Biotechnology and Bioengineering, 2020, 117, 2074-2088.	3.3	15
23	Triton X-114 cloud point extraction to subfractionate blood plasma proteins for two-dimensional gel electrophoresis. Analytical Biochemistry, 2015, 485, 11-17.	2.4	14
24	Proteome Analysis of Pyloric Ceca: A Methodology for Fish Feed Development?. Journal of Agricultural and Food Chemistry, 2012, 60, 8457-8464.	5.2	13
25	Identification and validation of novel small proteins in <i>Pseudomonas putida</i> . Environmental Microbiology Reports, 2016, 8, 966-974.	2.4	13
26	CRISPR interference of nucleotide biosynthesis improves production of a singleâ€domain antibody in <i>Escherichia coli</i> . Biotechnology and Bioengineering, 2020, 117, 3835-3848.	3.3	13
27	Effect of $\hat{l}\pm$ -lactalbumin and \hat{l}^2 -lactoglobulin on the oxidative stability of 10% fish oil-in-water emulsions depends on pH. Food Chemistry, 2013, 141, 574-581.	8.2	10
28	Muscle Protein Profiles Used for Prediction of Texture of Farmed Salmon (<i>Salmo salar</i> L.). Journal of Agricultural and Food Chemistry, 2017, 65, 3413-3421.	5.2	9
29	Directed Metabolic Pathway Evolution Enables Functional Pterin-Dependent Aromatic-Amino-Acid Hydroxylation in <i>Escherichia coli</i>). ACS Synthetic Biology, 2020, 9, 494-499.	3.8	9
30	Compartmentalized Proteomic Profiling Outlines the Crucial Role of the Classical Secretory Pathway during Recombinant Protein Production in Chinese Hamster Ovary Cells. ACS Omega, 2021, 6, 12439-12458.	3.5	9
31	Effect of <i>in vitro</i> digested cod liver oil of different quality on oxidative, proteomic and inflammatory responses in the yeast <i>Saccharomyces cerevisiae</i> and human monocyte-derived dendritic cells. Journal of the Science of Food and Agriculture, 2015, 95, 3096-3106.	3.5	8
32	Homologous Desensitisation of the Mouse Leukotriene B ₄ Receptor Involves Protein Kinase C-Mediated Phosphorylation of Serine 127. Cellular Physiology and Biochemistry, 2007, 20, 143-156.	1.6	5
33	KCNK5 is Functionally Down-Regulated Upon Long-Term Hypotonicity in Ehrlich Ascites Tumor Cells. Cellular Physiology and Biochemistry, 2013, 32, 1238-1246.	1.6	5
34	Emergence of Phenotypically Distinct Subpopulations Is a Factor in Adaptation of Recombinant Saccharomyces cerevisiae under Glucose-Limited Conditions. Applied and Environmental Microbiology, 2022, 88, e0230721.	3.1	5
35	Differentiating samples and experimental protocols by direct comparison of tandem mass spectra. Rapid Communications in Mass Spectrometry, 2016, 30, 731-738.	1.5	4
36	Enrichment of microsomes from Chinese hamster ovary cells by subcellular fractionation for its use in proteomic analysis. PLoS ONE, 2020, 15, e0237930.	2.5	4

#	Article	IF	CITATIONS
37	CHOmics: A web-based tool for multi-omics data analysis and interactive visualization in CHO cell lines. PLoS Computational Biology, 2020, 16, e1008498.	3.2	4
38	Bioactivity of Cod and Chicken Protein Hydrolysates before and after in vitro Gastrointestinal Digestion. Food Technology and Biotechnology, 2017, 55, 360-367.	2.1	3
39	Co-expression of mCysLT1 receptors and IK channels in Xenopus laevis oocytes elicits LTD4-stimulated IK current, independent of an increase in [Ca2+]i. Biochimica Et Biophysica Acta - Biomembranes, 2004, 1660, 75-79.	2.6	2
40	Tissue damage in organic rainbow trout muscle investigated by proteomics and bioinformatics. Proteomics, 2013, 13, 2180-2190.	2.2	0
41	Title is missing!. , 2020, 16, e1008498.		0
42	Title is missing!. , 2020, 16, e1008498.		0
43	Title is missing!. , 2020, 16, e1008498.		0
44	Title is missing!. , 2020, 16, e1008498.		0
45	Title is missing!. , 2020, 16, e1008498.		0
46	Title is missing!. , 2020, 16, e1008498.		0