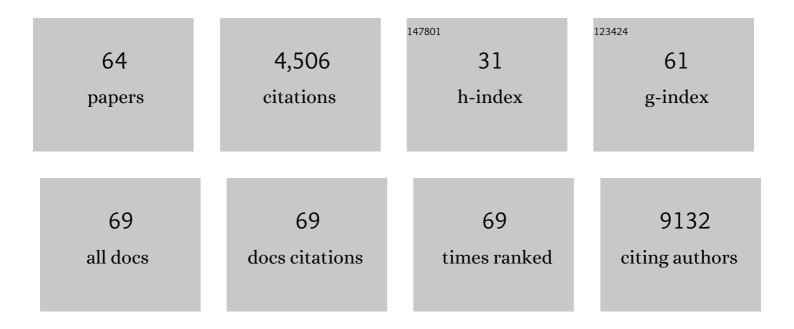
Hubert Rehrauer

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A MicroRNA Next-Generation-Sequencing Discovery Assay (miND) for Genome-Scale Analysis and Absolute Quantitation of Circulating MicroRNA Biomarkers. International Journal of Molecular Sciences, 2022, 23, 1226.	4.1	16
2	Phenotype of Mrps5-Associated Phylogenetic Polymorphisms Is Intimately Linked to Mitoribosomal Misreading. International Journal of Molecular Sciences, 2022, 23, 4384.	4.1	1
3	Effects of ACTH-Induced Long-Term Hypercortisolism on the Transcriptome of Canine Visceral Adipose Tissue. Veterinary Sciences, 2022, 9, 250.	1.7	1
4	Monocyte-derived alveolar macrophages autonomously determine severe outcome of respiratory viral infection. Science Immunology, 2022, 7, .	11.9	39
5	In vivo cytidine base editing of hepatocytes without detectable off-target mutations in RNA and DNA. Nature Biomedical Engineering, 2021, 5, 179-189.	22.5	62
6	Mitochondrial Mistranslation in Brain Provokes a Metabolic Response Which Mitigates the Age-Associated Decline in Mitochondrial Gene Expression. International Journal of Molecular Sciences, 2021, 22, 2746.	4.1	5
7	Endogenous retrovirus expression activates type-l interferon signaling in an experimental mouse model of mesothelioma development. Cancer Letters, 2021, 507, 26-38.	7.2	18
8	CD40 Agonist Monoclonal Antibody-Mediated Hepatitis in TNF-Receptor 1 Gene Knockout Mice. Biomedicines, 2021, 9, 863.	3.2	0
9	Double-Stranded RNA Structural Elements Holding the Key to Translational Regulation in Cancer: The Case of Editing in RNA-Binding Motif Protein 8A. Cells, 2021, 10, 3543.	4.1	5
10	Genetic deletion of microRNA biogenesis in muscle cells reveals a hierarchical non-clustered network that controls focal adhesion signaling during muscle regeneration. Molecular Metabolism, 2020, 36, 100967.	6.5	10
11	Ribosomal mistranslation leads to silencing of the unfolded protein response and increased mitochondrial biogenesis. Communications Biology, 2019, 2, 381.	4.4	33
12	Effects of a male meiotic driver on male and female transcriptomes in the house mouse. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20191927.	2.6	12
13	Gene expression profiling of the canine placenta during normal and antigestagen-induced luteolysis. General and Comparative Endocrinology, 2019, 282, 113194.	1.8	16
14	Long fragments achieve lower base quality in Illumina paired-end sequencing. Scientific Reports, 2019, 9, 2856.	3.3	66
15	An experimental genetically attenuated live vaccine to prevent transmission of Toxoplasma gondii by cats. Scientific Reports, 2019, 9, 1474.	3.3	112
16	Global Transcriptomic Analysis of the Canine corpus luteum (CL) During the First Half of Diestrus and Changes Induced by in vivo Inhibition of Prostaglandin Synthase 2 (PTGS2/COX2). Frontiers in Endocrinology, 2019, 10, 715.	3.5	6
17	How asbestos drives the tissue towards tumors: YAP activation, macrophage and mesothelial precursor recruitment, RNA editing, and somatic mutations. Oncogene, 2018, 37, 2645-2659.	5.9	53
18	Mutant <scp>MRPS</scp> 5 affects mitoribosomal accuracy and confers stressâ€related behavioral alterations. EMBO Reports, 2018, 19, .	4.5	26

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19	Non-Coding Transcript Heterogeneity in Mesothelioma: Insights from Asbestos-Exposed Mice. International Journal of Molecular Sciences, 2018, 19, 1163.	4.1	8
20	Epigenetically-driven anatomical diversity of synovial fibroblasts guides joint-specific fibroblast functions. Nature Communications, 2017, 8, 14852.	12.8	126
21	RNA-Seq Data Analysis: From Raw Data Quality Control to Differential Expression Analysis. Methods in Molecular Biology, 2017, 1669, 295-307.	0.9	29
22	Transcriptome analysis reveals differences in mechanisms regulating cessation of luteal function in pregnant and non-pregnant dogs. BMC Genomics, 2017, 18, 757.	2.8	26
23	SUSHI: an exquisite recipe for fully documented, reproducible and reusable NGS data analysis. BMC Bioinformatics, 2016, 17, 228.	2.6	90
24	Diurnal changes in the histone H3 signature H3K9ac H3K27ac H3S28p are associated with diurnal gene expression in <i>Arabidopsis</i> . Plant, Cell and Environment, 2016, 39, 2557-2569.	5.7	31
25	Field Trial and Molecular Characterization of RNAi-Transgenic Tomato Plants That Exhibit Resistance to Tomato Yellow Leaf Curl Geminivirus. Molecular Plant-Microbe Interactions, 2016, 29, 197-209.	2.6	60
26	MicroRNA MiR-199a-5p Regulates Smooth Muscle Cell Proliferation and Morphology by Targeting WNT2 Signaling Pathway. Journal of Biological Chemistry, 2015, 290, 7067-7086.	3.4	59
27	The Phosphoinositide 3-Kinase p110α Isoform Regulates Leukemia Inhibitory Factor Receptor Expression via c-Myc and miR-125b to Promote Cell Proliferation in Medulloblastoma. PLoS ONE, 2015, 10, e0123958.	2.5	24
28	Hypoxia attenuates the proinflammatory response in colon cancer cells by regulating lκB. Oncotarget, 2015, 6, 20288-20301.	1.8	23
29	Circadian behavior is light-reprogrammed by plastic DNA methylation. Nature Neuroscience, 2014, 17, 377-382.	14.8	218
30	Induction of the nuclear receptor PPAR-γ by the cytokine GM-CSF is critical for the differentiation of fetal monocytes into alveolar macrophages. Nature Immunology, 2014, 15, 1026-1037.	14.5	443
31	Targeting the Phosphoinositide 3-Kinase p110-α Isoform Impairs Cell Proliferation, Survival, and Tumor Growth in Small Cell Lung Cancer. Clinical Cancer Research, 2013, 19, 96-105.	7.0	30
32	An <scp>ER</scp> â€directed transcriptional response to unfolded protein stress in the absence of conserved sensorâ€ŧransducer proteins in <i><scp>G</scp>iardia lamblia</i> . Molecular Microbiology, 2013, 88, 754-771.	2.5	11
33	Role of Hedgehog Signaling in Malignant Pleural Mesothelioma. Clinical Cancer Research, 2012, 18, 4646-4656.	7.0	60
34	RIP-chip-SRM—a new combinatorial large-scale approach identifies a set of translationally regulated bantam/miR-58 targets in <i>C. elegans</i> . Genome Research, 2012, 22, 1360-1371.	5.5	18
35	Synthetic transactivation screening reveals ETV4 as broad coactivator of hypoxia-inducible factor signaling. Nucleic Acids Research, 2012, 40, 1928-1943.	14.5	32
36	Genomeâ€wide transcription analysis of <i>Escherichia coli</i> in response to extremely lowâ€frequency magnetic fields. Bioelectromagnetics, 2012, 33, 488-496.	1.6	18

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37	Induction of senescence markers after neo-adjuvant chemotherapy of malignant pleural mesothelioma and association with clinical outcome: An exploratory analysis. European Journal of Cancer, 2011, 47, 326-332.	2.8	58
38	Chronic exposure to cigarette smoke condensate in vitro induces epithelial to mesenchymal transition-like changes in human bronchial epithelial cells, BEAS-2B. Toxicology in Vitro, 2011, 25, 446-453.	2.4	77
39	CD133 Positive Embryonal Rhabdomyosarcoma Stem-Like Cell Population Is Enriched in Rhabdospheres. PLoS ONE, 2011, 6, e19506.	2.5	111
40	Myc-mediated repression of microRNA-34a promotes high-grade transformation of B-cell lymphoma by dysregulation of FoxP1. Blood, 2011, 117, 6227-6236.	1.4	165
41	Epigenetic Silencing of MicroRNA-203 Dysregulates ABL1 Expression and Drives <i>Helicobacter</i> -Associated Gastric Lymphomagenesis. Cancer Research, 2011, 71, 3616-3624.	0.9	101
42	Pleural mesothelioma side populations have a precursor phenotype. Carcinogenesis, 2011, 32, 1324-1332.	2.8	38
43	Application of anifHmicroarray to assess the impact of environmental factors on free-living diazotrophs in a glacier forefield. Canadian Journal of Microbiology, 2011, 57, 105-114.	1.7	4
44	Interaction of HIF and USF Signaling Pathways in Human Genes Flanked by Hypoxia-Response Elements and E-box Palindromes. Molecular Cancer Research, 2011, 9, 1520-1536.	3.4	21
45	Rhizobial Adaptation to Hosts, a New Facet in the Legume Root-Nodule Symbiosis. Molecular Plant-Microbe Interactions, 2010, 23, 784-790.	2.6	68
46	Lysyl oxidase expression is an independent marker of prognosis and a predictor of lymph node metastasis in oral and oropharyngeal squamous cell carcinoma (OSCC). International Journal of Cancer, 2010, 126, 2653-2662.	5.1	56
47	Epigenetic mechanisms regulate stage differentiation in the minimized protozoan <i>Giardia lamblia</i> . Molecular Microbiology, 2010, 76, 48-67.	2.5	85
48	The Transcriptional Response to Encystation Stimuli in Giardia lamblia Is Restricted to a Small Set of Genes. Eukaryotic Cell, 2010, 9, 1566-1576.	3.4	73
49	Absence of Poly(ADP-Ribose) Polymerase 1 Delays the Onset of <i>Salmonella enterica</i> Serovar Typhimurium-Induced Gut Inflammation. Infection and Immunity, 2010, 78, 3420-3431.	2.2	29
50	The <i>Streptococcus mutans</i> Serine/Threonine Kinase, PknB, Regulates Competence Development, Bacteriocin Production, and Cell Wall Metabolism. Infection and Immunity, 2010, 78, 2209-2220.	2.2	81
51	AGRONOMICS1: A New Resource for Arabidopsis Transcriptome Profiling Â. Plant Physiology, 2010, 152, 487-499.	4.8	61
52	A global network of coexisting microbes from environmental and whole-genome sequence data. Genome Research, 2010, 20, 947-959.	5.5	425
53	Ablation of Dicer from Murine Schwann Cells Increases Their Proliferation while Blocking Myelination. PLoS ONE, 2010, 5, e12450.	2.5	69
54	The CD4+ T Cell-Mediated IFN-Î ³ Response to <i>Helicobacter</i> Infection Is Essential for Clearance and Determines Gastric Cancer Risk. Journal of Immunology, 2009, 182, 7085-7101.	0.8	151

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55	Deterministic protein inference for shotgun proteomics data provides new insights into Arabidopsis pollen development and function. Genome Research, 2009, 19, 1786-1800.	5.5	151
56	16S rRNA geneâ€based phylogenetic microarray for simultaneous identification of members of the genus <i>Burkholderia</i> . Environmental Microbiology, 2009, 11, 779-800.	3.8	22
57	Development and experimental validation of a <i>nifH</i> oligonucleotide microarray to study diazotrophic communities in a glacier forefield. Environmental Microbiology, 2009, 11, 2179-2189.	3.8	16
58	Malignant Catarrhal Fever of Cattle Is Associated with Low Abundance of IL-2 Transcript and a Predominantly Latent Profile of Ovine Herpesvirus 2 Gene Expression. PLoS ONE, 2009, 4, e6265.	2.5	25
59	Genome-wide gene expression profiling reveals renal genes regulated during metabolic acidosis. Physiological Genomics, 2008, 32, 322-334.	2.3	77
60	Functional relevance of novel p300-mediated lysine 314 and 315 acetylation of RelA/p65. Nucleic Acids Research, 2008, 36, 1665-1680.	14.5	91
61	Transcriptome Profile of Human Colorectal Adenomas. Molecular Cancer Research, 2007, 5, 1263-1275.	3.4	428
62	Sirolimus ameliorates the enhanced expression of metalloproteinases in a rat model of autosomal dominant polycystic kidney disease. Nephrology Dialysis Transplantation, 2007, 23, 880-889.	0.7	27
63	Genome-Wide Transcript Analysis of <i>Bradyrhizobium japonicum</i> Bacteroids in Soybean Root Nodules. Molecular Plant-Microbe Interactions, 2007, 20, 1353-1363.	2.6	187
64	Gene Expression Profiles of Different Breast Cancer Cells Compared with their Responsiveness to Fermented Mistletoe (Viscum album L.) Extracts Iscador from Oak (Quercus), Pine (Pinus), White Fir (Abies) and Apple Tree (Malus) in vitro. Arzneimittelforschung, 2006, 56, 483-496.	0.4	8