Torben LÃ¹/₄ders

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/84532/publications.pdf

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30 papers

1,284 citations

430874 18 h-index 28 g-index

31 all docs

31 docs citations

times ranked

31

2805 citing authors

#	Article	IF	Citations
1	Hipposin, a histone-derived antimicrobial peptide in Atlantic halibut (Hippoglossus hippoglossus L.). Biochimica Et Biophysica Acta - Proteins and Proteomics, 2003, 1646, 207-215.	2.3	164
2	Breast cancer quantitative proteome and proteogenomic landscape. Nature Communications, 2019, 10, 1600.	12.8	152
3	Glycan gene expression signatures in normal and malignant breast tissue; possible role in diagnosis and progression. Molecular Oncology, 2010, 4, 98-118.	4.6	147
4	Deregulation of cancer-related miRNAs is a common event in both benign and malignant human breast tumors. Carcinogenesis, 2014, 35, 76-85.	2.8	119
5	Proline Conformation-Dependent Antimicrobial Activity of a Proline-Rich Histone H1 N-Terminal Peptide Fragment Isolated from the Skin Mucus of Atlantic Salmon. Antimicrobial Agents and Chemotherapy, 2005, 49, 2399-2406.	3.2	87
6	Integrative clustering reveals a novel split in the luminal A subtype of breast cancer with impact on outcome. Breast Cancer Research, 2017, 19, 44.	5.0	85
7	Strong Synergy between a Eukaryotic Antimicrobial Peptide and Bacteriocins from Lactic Acid Bacteria. Applied and Environmental Microbiology, 2003, 69, 1797-1799.	3.1	74
8	Subtypeâ€specific microâ€RNA expression signatures in breast cancer progression. International Journal of Cancer, 2016, 139, 1117-1128.	5.1	53
9	Glycanâ€related gene expression signatures in breast cancer subtypes; relation to survival. Molecular Oncology, 2015, 9, 861-876.	4.6	47
10	Gene expression profiles of breast biopsies from healthy women identify a group with claudin-low features. BMC Medical Genomics, 2011, 4, 77.	1.5	38
11	Expression levels of uridine 5'-diphospho-glucuronosyltransferase genes in breast tissue from healthy women are associated with mammographic density. Breast Cancer Research, 2010, 12, R65.	5.0	37
12	Serum estradiol levels associated with specific gene expression patterns in normal breast tissue and in breast carcinomas. BMC Cancer, 2011, 11, 332.	2.6	35
13	Long Non-Coding RNAs Differentially Expressed between Normal versus Primary Breast Tumor Tissues Disclose Converse Changes to Breast Cancer-Related Protein-Coding Genes. PLoS ONE, 2014, 9, e106076.	2.5	35
14	Age, estrogen, and immune response in breast adenocarcinoma and adjacent normal tissue. Oncolmmunology, 2017, 6, e1356142.	4.6	34
15	Canine Mammary Tumours Are Affected by Frequent Copy Number Aberrations, including Amplification of MYC and Loss of PTEN. PLoS ONE, 2015, 10, e0126371.	2.5	28
16	Noninvasive profiling of serum cytokines in breast cancer patients and clinicopathological characteristics. Oncolmmunology, 2019, 8, e1537691.	4.6	27
17	Expression of BMI-1 and Mel-18 in breast tissue - a diagnostic marker in patients with breast cancer. BMC Cancer, 2010, 10, 686.	2.6	23
18	Globular adiponectin and its downstream target genes are up-regulated locally in human colorectal tumors: ex vivo and in vitro studies. Metabolism: Clinical and Experimental, 2014, 63, 672-681.	3.4	23

#	Article	IF	CITATIONS
19	Serum cytokine levels in breast cancer patients during neoadjuvant treatment with bevacizumab. Oncolmmunology, 2018, 7, e1457598.	4.6	18
20	Identification and structural analysis of the antimicrobial domain in hipposin, a 51-mer antimicrobial peptide isolated from Atlantic halibut. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2004, 1699, 221-227.	2.3	17
21	Loss of progesterone receptor is associated with distinct tyrosine kinase profiles in breast cancer. Breast Cancer Research and Treatment, 2020, 183, 585-598.	2.5	10
22	Molecular Profiles of Pre- and Postoperative Breast Cancer Tumours Reveal Differentially Expressed Genes. ISRN Oncology, 2012, 2012, 1-12.	2.1	6
23	mRNA expression of adipocytokines and glucocorticoid-related genes are associated with downregulation of E-cadherin mRNA in colorectal adenocarcinomas. International Journal of Colorectal Disease, 2012, 27, 1021-1027.	2.2	5
24	Comparable cancerâ€relevant mutation profiles in synchronous ductal carcinoma in situ and invasive breast cancer. Cancer Reports, 2020, 3, e1248.	1.4	5
25	ZBTB11 dysfunction: spectrum of brain abnormalities, biochemical signature and cellular consequences. Brain, 2022, 145, 2602-2616.	7.6	5
26	Mucosal Gene Transcript Signatures in Treatment Naïve Inflammatory Bowel Disease: A Comparative Analysis of Disease to Symptomatic and Healthy Controls in the European IBD-Character Cohort. Clinical and Experimental Gastroenterology, 2022, Volume 15, 5-25.	2.3	5
27	Gene Expression Profile Analysis of T1 and T2 Breast Cancer Reveals Different Activation Pathways. ISRN Oncology, 2013, 2013, 1-12.	2.1	3
28	Gene expression analysis supports tumor threshold over 2.0Âcm for T-category breast cancer. Eurasip Journal on Bioinformatics and Systems Biology, 2016, 2016, 6.	1.4	2
29	Circulating mitochondrial DNA (mtDNA) variants to predict metastatic progression of rectal cancer Journal of Clinical Oncology, 2020, 38, e16132-e16132.	1.6	0
30	Abstract OT2-19-01: Presurgical treatment with ribociclib and letrozole in patients with locally advanced breast cancer: The NEOLETRIB study. Cancer Research, 2022, 82, OT2-19-01-OT2-19-01.	0.9	O