## Torben Lüders

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	ZBTB11 dysfunction: spectrum of brain abnormalities, biochemical signature and cellular consequences. Brain, 2022, 145, 2602-2616.	7.6	5
2	Mucosal Gene Transcript Signatures in Treatment NaÃ <sup>-</sup> ve 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
3	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	0
4	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
5	Comparable cancerâ€relevant mutation profiles in synchronous ductal carcinoma in situ and invasive breast cancer. Cancer Reports, 2020, 3, e1248.	1.4	5
6	Circulating mitochondrial DNA (mtDNA) variants to predict metastatic progression of rectal cancer Journal of Clinical Oncology, 2020, 38, e16132-e16132.	1.6	0
7	Breast cancer quantitative proteome and proteogenomic landscape. Nature Communications, 2019, 10, 1600.	12.8	152
8	Noninvasive profiling of serum cytokines in breast cancer patients and clinicopathological characteristics. Oncolmmunology, 2019, 8, e1537691.	4.6	27
9	Serum cytokine levels in breast cancer patients during neoadjuvant treatment with bevacizumab. Oncolmmunology, 2018, 7, e1457598.	4.6	18
10	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
11	Age, estrogen, and immune response in breast adenocarcinoma and adjacent normal tissue. Oncolmmunology, 2017, 6, e1356142.	4.6	34
12	Subtypeâ€ <b>s</b> pecific microâ€RNA expression signatures in breast cancer progression. International Journal of Cancer, 2016, 139, 1117-1128.	5.1	53
13	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
14	Glycanâ€related gene expression signatures in breast cancer subtypes; relation to survival. Molecular Oncology, 2015, 9, 861-876.	4.6	47
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	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
17	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
18	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

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19	Gene Expression Profile Analysis of T1 and T2 Breast Cancer Reveals Different Activation Pathways. ISRN Oncology, 2013, 2013, 1-12.	2.1	3
20	Molecular Profiles of Pre- and Postoperative Breast Cancer Tumours Reveal Differentially Expressed Genes. ISRN Oncology, 2012, 2012, 1-12.	2.1	6
21	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
22	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
23	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
24	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
25	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
26	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
27	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
28	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
29	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
30	Strong Synergy between a Eukaryotic Antimicrobial Peptide and Bacteriocins from Lactic Acid Bacteria. Applied and Environmental Microbiology, 2003, 69, 1797-1799.	3.1	74