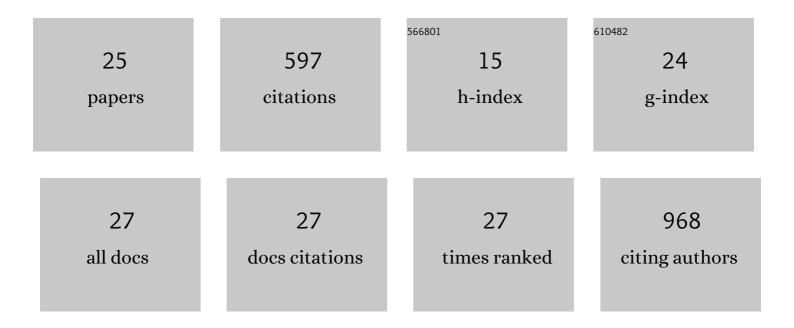
## Birte Möhlendick

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1961358/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Patients with Liver Cirrhosis Show High Immunogenicity upon COVID-19 Vaccination but Develop Premature Deterioration of Antibody Titers. Vaccines, 2022, 10, 377.	2.1	30
2	NAT2 polymorphisms as a cause of metamizole-induced agranulocytosis. Pharmacogenetics and Genomics, 2021, Publish Ahead of Print, 140-143.	0.7	2
3	Transmembrane serine protease 2 Polymorphisms and Susceptibility to Severe Acute Respiratory Syndrome Coronavirus Type 2 Infection: A German Case-Control Study. Frontiers in Genetics, 2021, 12, 667231.	1.1	43
4	ACE2 polymorphism and susceptibility for SARS-CoV-2 infection and severity of COVID-19. Pharmacogenetics and Genomics, 2021, 31, 165-171.	0.7	73
5	The influence of IFITM3 polymorphisms on susceptibility to SARS-CoV-2 infection and severity of COVID-19. Cytokine, 2021, 142, 155492.	1.4	37
6	Development Of De Novo Donor-Specific HLA Antibodies and ABMR in Renal Transplant Patients Depends on CYP3A5 Genotype. Transplantation, 2021, Publish Ahead of Print, .	0.5	3
7	BRAF <sup>V600E</sup> mutation: A promising target in colorectal neuroendocrine carcinoma. International Journal of Cancer, 2019, 144, 1379-1390.	2.3	33
8	The <i>GNAS</i> SNP c.393C>T (rs7121) as a marker for disease progression and survival in cancer. Pharmacogenomics, 2019, 20, 553-562.	0.6	11
9	Diagnostic pathology of early systemic cancer: <i>ERBB2</i> gene amplification in single disseminated cancer cells determines patient survival in operable esophageal cancer. International Journal of Cancer, 2018, 142, 833-843.	2.3	15
10	Establishment and Characterization of a Novel Cell Line Derived from a Small Cell Neuroendocrine Carcinoma of the Anal Canal. Neuroendocrinology, 2018, 107, 246-256.	1.2	4
11	A streamlined workflow for single-cells genome-wide copy-number profiling by low-pass sequencing of LM-PCR whole-genome amplification products. PLoS ONE, 2018, 13, e0193689.	1.1	20
12	Disseminated tumour cells with highly aberrant genomes are linked to poor prognosis in operable oesophageal adenocarcinoma. British Journal of Cancer, 2017, 117, 725-733.	2.9	17
13	Preclinical assesement of survivin and XIAP as prognostic biomarkers and therapeutic targets in gastroenteropancreatic neuroendocrine neoplasia. Oncotarget, 2017, 8, 8369-8382.	0.8	17
14	Isolation of circulating tumor cells from pancreatic cancer by automated filtration. Oncotarget, 2017, 8, 86143-86156.	0.8	24
15	Peritoneal sarcomatosis: site of origin for the establishment of an in vitro and in vivo cell line model to study therapeutic resistance in dedifferentiated liposarcoma. Tumor Biology, 2016, 37, 2341-2351.	0.8	4
16	Unscrambling the genomic chaos of osteosarcoma reveals extensive transcript fusion, recurrent rearrangements and frequent novel TP53 aberrations. Oncotarget, 2016, 7, 5273-5288.	0.8	60
17	Abstract 2394: Scalable, rapid and affordable low-pass whole genome sequencing method for single-cell copy-number profiling on LM-PCR based WGA products. , 2016, , .		0
18	Serum microRNA profiles as prognostic or predictive markers in the multimodality treatment of patients with gastric cancer. Oncology Letters, 2015, 10, 869-874.	0.8	8

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19	Serum micro <scp>RNA</scp> profiles as prognostic/predictive markers in the multimodality therapy of locally advanced adenocarcinomas of the gastroesophageal junction. International Journal of Cancer, 2015, 137, 230-237.	2.3	24
20	New Model for Gastroenteropancreatic Large-Cell Neuroendocrine Carcinoma: Establishment of Two Clinically Relevant Cell Lines. PLoS ONE, 2014, 9, e88713.	1.1	28
21	A New Workflow for Whole-Genome Sequencing of Single Human Cells. Human Mutation, 2014, 35, 1260-1270.	1.1	19
22	Analysis of Copyâ€Number Alterations in Single Cells Using Microarrayâ€Based Comparative Genomic Hybridization (aCGH). Current Protocols in Cell Biology, 2014, 65, 22.19.1-23.	2.3	10
23	Genomic High-Resolution Profiling of Single CKpos/CD45neg Flow-Sorting Purified Circulating Tumor Cells from Patients with Metastatic Breast Cancer. Clinical Chemistry, 2014, 60, 1290-1297.	1.5	74
24	A Robust Method to Analyze Copy Number Alterations of Less than 100 kb in Single Cells Using Oligonucleotide Array CGH. PLoS ONE, 2013, 8, e67031.	1.1	40
25	From primary to secondary cancer formation – models for disseminating cancer cells. Drug Discovery Today: Disease Models, 2011, 8, 75-79.	1.2	1