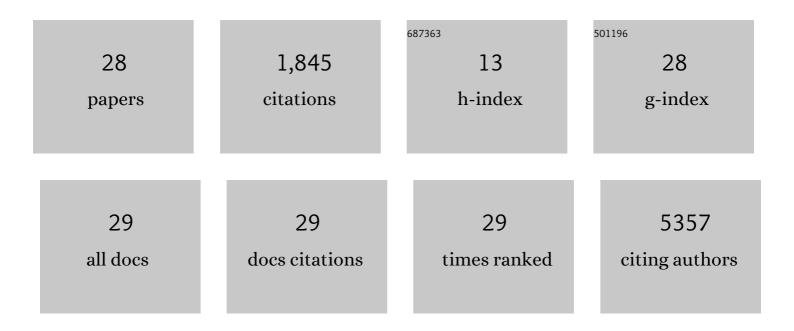
Michael Forster

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Detection of Cancer Mutations by Urine Liquid Biopsy as a Potential Tool in the Clinical Management of Bladder Cancer Patients. Cancers, 2022, 14, 969.	3.7	2
2	High-throughput method for the hybridisation-based targeted enrichment of long genomic fragments for PacBio third-generation sequencing. NAR Genomics and Bioinformatics, 2022, 4, .	3.2	7
3	Clinical responses to PD-1 inhibition and their molecular characterization in six patients with mismatch repair-deficient metastatic cancer of the digestive system. Journal of Cancer Research and Clinical Oncology, 2021, 147, 263-273.	2.5	5
4	Quantifying sequencing error and effective sequencing depth of liquid biopsy NGS with UMI error correction. BioTechniques, 2021, 70, 226-232.	1.8	5
5	Genome-wide analysis of 944 133 individuals provides insights into the etiology of haemorrhoidal disease. Gut, 2021, 70, 1538-1549.	12.1	21
6	Liquid Biopsy in Gastric Cancer: Analysis of Somatic Cancer Tissue Mutations in Plasma Cell-Free DNA for Predicting Disease State and Patient Survival. Clinical and Translational Gastroenterology, 2021, 12, e00403.	2.5	8
7	Genetic Variation in ABCC4 and CFTR and Acute Pancreatitis during Treatment of Pediatric Acute Lymphoblastic Leukemia. Journal of Clinical Medicine, 2021, 10, 4815.	2.4	2
8	Deficiency in X-linked inhibitor of apoptosis protein promotes susceptibility to microbial triggers of intestinal inflammation. Science Immunology, 2021, 6, eabf7473.	11.9	15
9	Rapid response of stage IV colorectal cancer with APC/TP53/KRAS mutations to FOLFIRI and Bevacizumab combination chemotherapy: a case report of use of liquid biopsy. BMC Medical Genetics, 2020, 21, 3.	2.1	5
10	Two Prevalent â^¼100-kb <i>GYPB</i> Deletions Causative of the GPB-Deficient Blood Group MNS Phenotype S–s–U– in Black Africans. Transfusion Medicine and Hemotherapy, 2020, 47, 326-336.	1.6	8
11	Longitudinal Multi-omics Analyses Identify Responses of Megakaryocytes, Erythroid Cells, and Plasmablasts as Hallmarks of Severe COVID-19. Immunity, 2020, 53, 1296-1314.e9.	14.3	278
12	Stage IV Colorectal Cancer Patients with High Risk Mutation Profiles Survived 16 Months Longer with Individualized Therapies. Cancers, 2020, 12, 393.	3.7	3
13	Phylogenetic network analysis of SARS-CoV-2 genomes. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 9241-9243.	7.1	840
14	Reply to Sánchez-Pacheco et al., Chookajorn, and Mavian et al.: Explaining phylogenetic network analysis of SARS-CoV-2 genomes. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 12524-12525.	7.1	19
15	Abstract 3617: Very low coverage whole genome sequencing improves clinically relevant copy number variation calling compared to targeted sequencing. , 2020, , .		0
16	Effects of Quantification Methods, Isolation Kits, Plasma Biobanking, and Hemolysis on Cell-Free DNA Analysis in Plasma. Biopreservation and Biobanking, 2019, 17, 553-561.	1.0	11
17	Leptin induces TNFα-dependent inflammation in acquired generalized lipodystrophy and combined Crohn's disease. Nature Communications, 2019, 10, 5629.	12.8	27
18	NGS-based methylation profiling differentiates TCF3-HLF and TCF3-PBX1 positive B-cell acute lymphoblastic leukemia. Epigenomics, 2018, 10, 133-147.	2.1	10

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19	Grow With the Challenge – Microbial Effects on Epithelial Proliferation, Carcinogenesis, and Cancer Therapy. Frontiers in Microbiology, 2018, 9, 2020.	3.5	26
20	RNA based individualized drug selection in breast cancer patients without patient-matched normal tissue. Oncotarget, 2018, 9, 32362-32372.	1.8	1
21	Quantifying cell free DNA in urine: comparison between commercial kits, impact of gender and inter-individual variation. BioTechniques, 2018, 64, 225-230.	1.8	31
22	Metastatic triple-negative breast cancer patient with <i>TP53</i> tumor mutation experienced 11 months progression-free survival on bortezomib monotherapy without adverse events after ending standard treatments with grade 3 adverse events. Journal of Physical Education and Sports Management, 2017, 3, a001677.	1.2	14
23	Identification and characterization of two functional variants in the human longevity gene FOXO3. Nature Communications, 2017, 8, 2063.	12.8	69
24	Development of a high-resolution NGS-based HLA-typing and analysis pipeline. Nucleic Acids Research, 2015, 43, e70-e70.	14.5	77
25	Genomics and drug profiling of fatal TCF3-HLFâ^ positive acute lymphoblastic leukemia identifies recurrent mutation patterns and therapeutic options. Nature Genetics, 2015, 47, 1020-1029.	21.4	190
26	Vy-PER: eliminating false positive detection of virus integration events in next generation sequencing data. Scientific Reports, 2015, 5, 11534.	3.3	42
27	Early-onset Crohn's disease and autoimmunity associated with a variant in CTLA-4. Gut, 2015, 64, 1889-1897.	12.1	106
28	From next-generation sequencing alignments to accurate comparison and validation of single-nucleotide variants: the pibase software. Nucleic Acids Research, 2013, 41, e16-e16.	14.5	21