## Georg Hemmrich-Stanisak

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genomics and drug profiling of fatal TCF3-HLFâ	21.4	190
2	A comprehensive, cell specific microRNA catalogue of human peripheral blood. Nucleic Acids Research, 2017, 45, 9290-9301.	14.5	159
3	Sparse Modeling Reveals miRNA Signatures for Diagnostics of Inflammatory Bowel Disease. PLoS ONE, 2015, 10, e0140155.	2.5	31
4	Genome-wide analysis of 944 133 individuals provides insights into the etiology of haemorrhoidal disease. Gut, 2021, 70, 1538-1549.	12.1	21
5	Depletion of erythropoietic miR-486-5p and miR-451a improves detectability of rare microRNAs in peripheral blood-derived small RNA sequencing libraries. NAR Genomics and Bioinformatics, 2020, 2, lqaa008.	3.2	12
6	NGS-based methylation profiling differentiates TCF3-HLF and TCF3-PBX1 positive B-cell acute lymphoblastic leukemia. Epigenomics, 2018, 10, 133-147.	2.1	10
7	Integrating Culture-based Antibiotic Resistance Profiles with Whole-genome Sequencing Data for 11,087 Clinical Isolates. Genomics, Proteomics and Bioinformatics, 2019, 17, 169-182.	6.9	8
8	Comparing genome versus proteome-based identification of clinical bacterial isolates. Briefings in Bioinformatics, 2018, 19, bbw122.	6.5	7
9	miRNome Profiling and Functional Analysis Reveal Involvement of hsa-miR-1246 in Colon Adenoma-Carcinoma Transition by Targeting AXIN2 and CFTR. International Journal of Molecular Sciences, 2022, 23, 2107.	4.1	7
10	Detailed Transcriptional Landscape of Peripheral Blood Points to Increased Neutrophil Activation in Treatment-NaÃ <sup>-</sup> ve Inflammatory Bowel Disease. Journal of Crohn's and Colitis, 2022, 16, 1097-1109.	1.3	5
11	RNA based individualized drug selection in breast cancer patients without patient-matched normal tissue. Oncotarget, 2018, 9, 32362-32372.	1.8	1