

# Georg Hemmrich-Stanisak

## List of Publications by Year in descending order

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Version: 2024-02-01

11  
papers

453  
citations

1307594

7  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

1354  
citing authors

#	ARTICLE	IF	CITATIONS
1	Detailed Transcriptional Landscape of Peripheral Blood Points to Increased Neutrophil Activation in Treatment-Naïve Inflammatory Bowel Disease. <i>Journal of Crohn's and Colitis</i> , 2022, 16, 1097-1109.	1.3	5
2	miRNome Profiling and Functional Analysis Reveal Involvement of hsa-miR-1246 in Colon Adenoma-Carcinoma Transition by Targeting AXIN2 and CFTR. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2107.	4.1	7
3	Genome-wide analysis of 944 133 individuals provides insights into the etiology of haemorrhoidal disease. <i>Gut</i> , 2021, 70, 1538-1549.	12.1	21
4	Depletion of erythropoietic miR-486-5p and miR-451a improves detectability of rare microRNAs in peripheral blood-derived small RNA sequencing libraries. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa008.	3.2	12
5	Integrating Culture-based Antibiotic Resistance Profiles with Whole-genome Sequencing Data for 11,087 Clinical Isolates. <i>Genomics, Proteomics and Bioinformatics</i> , 2019, 17, 169-182.	6.9	8
6	Comparing genome versus proteome-based identification of clinical bacterial isolates. <i>Briefings in Bioinformatics</i> , 2018, 19, bbw122.	6.5	7
7	NGS-based methylation profiling differentiates TCF3-HLF and TCF3-PBX1 positive B-cell acute lymphoblastic leukemia. <i>Epigenomics</i> , 2018, 10, 133-147.	2.1	10
8	RNA based individualized drug selection in breast cancer patients without patient-matched normal tissue. <i>Oncotarget</i> , 2018, 9, 32362-32372.	1.8	1
9	A comprehensive, cell specific microRNA catalogue of human peripheral blood. <i>Nucleic Acids Research</i> , 2017, 45, 9290-9301.	14.5	159
10	Genomics and drug profiling of fatal TCF3-HLF <sup>+</sup> positive acute lymphoblastic leukemia identifies recurrent mutation patterns and therapeutic options. <i>Nature Genetics</i> , 2015, 47, 1020-1029.	21.4	190
11	Sparse Modeling Reveals miRNA Signatures for Diagnostics of Inflammatory Bowel Disease. <i>PLoS ONE</i> , 2015, 10, e0140155.	2.5	31