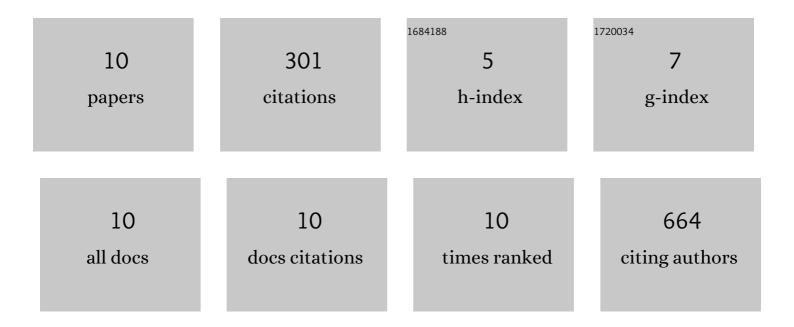
Pim G N J Mutsaers

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9786122/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Life-prolonging treatment restrictions and outcomes in patients with cancer and COVID-19: an update from the Dutch Oncology COVID-19 Consortium. European Journal of Cancer, 2022, 160, 261-272.	2.8	7
2	Diminished amplification of SARS-CoV-2 ORF1ab in a commercial dual-target qRT-PCR diagnostic assay. Journal of Virological Methods, 2022, 300, 114397.	2.1	5
3	Biological and Clinical Implications of Gene-Expression Profiling in Diffuse Large B-Cell Lymphoma: A Proposal for a Targeted BLYM-777 Consortium Panel as Part of a Multilayered Analytical Approach. Cancers, 2022, 14, 1857.	3.7	6
4	V-Domain Ig Suppressor of T Cell Activation (VISTA) Expression Is an Independent Prognostic Factor in Multiple Myeloma. Cancers, 2021, 13, 2219.	3.7	7
5	Poor outcome of patients with COVID-19 after CAR T-cell therapy for B-cell malignancies: results of a multicenter study on behalf of the European Society for Blood and Marrow Transplantation (EBMT) Infectious Diseases Working Party and the European Hematology Association (EHA) Lymphoma Group. Leukemia. 2021. 35. 3585-3588.	7.2	72
6	Subacute neurological deficits and respiratory insufficiency due to intrathecal methotrexate Journal of Clinical and Translational Research, 2021, 7, 809-810.	0.3	1
7	Dutch Oncology COVID-19 consortium: Outcome of COVID-19 in patients with cancer in a nationwide cohort study. European Journal of Cancer, 2020, 141, 171-184.	2.8	65
8	COVIDâ€19â€associated immune thrombocytopenia. British Journal of Haematology, 2020, 190, e61-e64.	2.5	138
9	Transcriptomics in Multiple Myeloma Demonstrates an Association between Survival and Expression of T Cell Co-Signaling Ligands in Bone Marrow Derived Myeloma Plasma Cells. Blood, 2018, 132, 241-241.	1.4	0
10	Familial AML With Germline CEBPA Mutations: Extended Clinical Outcomes and Analysis Of Secondary Mutations Using Whole Exome Sequencing. Blood, 2013, 122, 740-740.	1.4	0