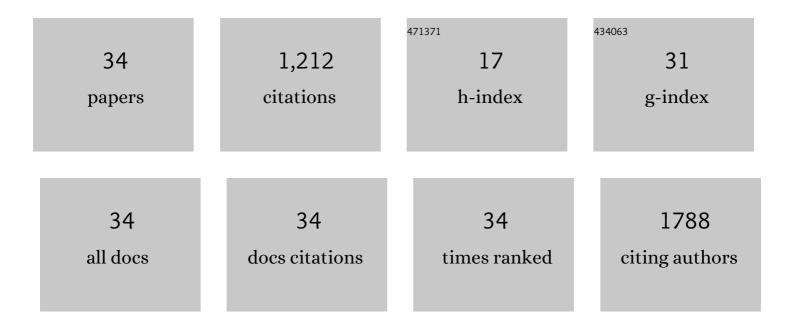
## M Kortüm

List of Publications by Year in descending order

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M KOPTÃ1/M

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | <sup>68</sup> Ga-Pentixafor PET/CT for Detection of Chemokine Receptor CXCR4 Expression in<br>Myeloproliferative Neoplasms. Journal of Nuclear Medicine, 2022, 63, 96-99.   | 2.8  | 13        |
| 2  | Transient regulatory T-cell targeting triggers immune control of multiple myeloma and prevents disease progression. Leukemia, 2022, 36, 790-800.  | 3.3  | 22        |
| 3  | Salvage therapy with "Daraâ€KDTâ€P(A)CE―in heavily pretreated, high―isk, proliferative,<br>relapsed/refractory multiple myeloma. Hematological Oncology, 2022, 40, 202-211.   | 0.8  | 9         |
| 4  | Minimal residual disease and imagingâ€guided consolidation strategies in newly diagnosed and relapsed refractory multiple myeloma. British Journal of Haematology, 2022, 198, 515-522.                                      | 1.2  | 7         |
| 5  | COVID-19 infection in patients with multiple myeloma: a German-Chinese experience from Würzburg and Wuhan. Annals of Hematology, 2021, 100, 843-846.  | 0.8  | 1         |
| 6  | Homozygous BCMA gene deletion in response to anti-BCMA CAR T cells in a patient with multiple myeloma. Nature Medicine, 2021, 27, 616-619.  | 15.2 | 140       |
| 7  | Actin cytoskeleton deregulation confers midostaurin resistance in FLT3-mutant acute myeloid<br>leukemia. Communications Biology, 2021, 4, 799.  | 2.0  | 16        |
| 8  | Cereblon enhancer methylation and IMiD resistance in multiple myeloma. Blood, 2021, 138, 1721-1726.   | 0.6  | 25        |
| 9  | Single- and double-hit events in genes encoding immune targets before and after T cell–engaging antibody therapy in MM. Blood Advances, 2021, 5, 3794-3798.   | 2.5  | 30        |
| 10 | <i>CIC</i> Mutation as a Molecular Mechanism of Acquired Resistance to Combined BRAF-MEK<br>Inhibition in Extramedullary Multiple Myeloma with Central Nervous System Involvement. Oncologist,<br>2020, 25, 112-118.        | 1.9  | 39        |
| 11 | 18F-FDG, 11C-Methionine, and 68Ga-Pentixafor PET/CT in Patients with Smoldering Multiple Myeloma:<br>Imaging Pattern and Clinical Features. Cancers, 2020, 12, 2333.  | 1.7  | 16        |
| 12 | The Link between Cytogenetics/Genomics and Imaging Patterns of Relapse and Progression in Patients<br>with Relapsed/Refractory Multiple Myeloma: A Pilot Study Utilizing 18F-FDG PET/CT. Cancers, 2020, 12,<br>2399.        | 1.7  | 4         |
| 13 | Toxicities of Chimeric Antigen Receptor T Cell Therapy in Multiple Myeloma: An Overview of Experience<br>From Clinical Trials, Pathophysiology, and Management Strategies. Frontiers in Immunology, 2020, 11,<br>620312.    | 2.2  | 21        |
| 14 | IKZF1/3 and CRL4 <sup>CRBN</sup> E3 ubiquitin ligase mutations and resistance to immunomodulatory drugs in multiple myeloma. Haematologica, 2020, 105, e237-e241.   | 1.7  | 41        |
| 15 | Carfilzomib Based Treatment Strategies in the Management of Relapsed/Refractory Multiple Myeloma<br>with Extramedullary Disease. Cancers, 2020, 12, 1035.   | 1.7  | 28        |
| 16 | 18F-FDG and 11C-Methionine PET/CT in Newly Diagnosed Multiple Myeloma Patients: Comparison of<br>Volume-Based PET Biomarkers. Cancers, 2020, 12, 1042.  | 1.7  | 24        |
| 17 | Different MAF translocations confer similar prognosis in newly diagnosed multiple myeloma patients.<br>Leukemia and Lymphoma, 2020, 61, 1885-1893.  | 0.6  | 3         |
| 18 | Sequential CD38 monoclonal antibody retreatment leads to deep remission in a patient with<br>relapsed/refractory multiple myeloma. International Journal of Immunopathology and Pharmacology,<br>2020, 34, 205873842098025. | 1.0  | 1         |

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| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Spectrum and functional validation of PSMB5 mutations in multiple myeloma. Leukemia, 2019, 33, 447-456.   | 3.3 | 93        |
| 20 | Response to daratumumab in rituximab-resistant EBV-associated PTLD following allogenic stem cell<br>transplantation from an EBV seronegative donor. Leukemia and Lymphoma, 2019, 60, 3573-3576.             | 0.6 | 10        |
| 21 | The Impact of Tumor Heterogeneity on Diagnostics and Novel Therapeutic Strategies in Multiple<br>Myeloma. International Journal of Molecular Sciences, 2019, 20, 1248.                                      | 1.8 | 54        |
| 22 | Assessment of TP53 lesions for p53 system functionality and drug resistance in multiple myeloma using an isogenic cell line model. Scientific Reports, 2019, 9, 18062.                                      | 1.6 | 14        |
| 23 | Protocol for M3P: A Comprehensive and Clinical Oriented Targeted Sequencing Panel for Routine<br>Molecular Analysis in Multiple Myeloma. Methods in Molecular Biology, 2018, 1792, 117-128.                 | 0.4 | 6         |
| 24 | CRISPR Genome-Wide Screening Identifies Dependence on the Proteasome Subunit PSMC6 for<br>Bortezomib Sensitivity in Multiple Myeloma. Molecular Cancer Therapeutics, 2017, 16, 2862-2870.                   | 1.9 | 54        |
| 25 | [ <sup>68</sup> Ga]Pentixafor-PET/CT for imaging of chemokine receptor CXCR4 expression in multiple<br>myeloma - Comparison to [ <sup>18</sup> F]FDG and laboratory values. Theranostics, 2017, 7, 205-212. | 4.6 | 138       |
| 26 | <sup>11</sup> C-Methionine-PET in Multiple Myeloma: A Combined Study from Two Different<br>Institutions. Theranostics, 2017, 7, 2956-2964.  | 4.6 | 63        |
| 27 | CXCR4-directed endoradiotherapy induces high response rates in extramedullary relapsed Multiple<br>Myeloma. Theranostics, 2017, 7, 1589-1597.   | 4.6 | 102       |
| 28 | Targeted sequencing of refractory myeloma reveals a high incidence of mutations in CRBN and Ras pathway genes. Blood, 2016, 128, 1226-1233.   | 0.6 | 185       |
| 29 | Proteasome inhibitors block Ikaros degradation by Lenalidomide in Multiple Myeloma. Haematologica, 2015, 100, e315-7.   | 1.7 | 20        |
| 30 | SnapShot: Multiple Myeloma. Cancer Cell, 2015, 28, 678-678.e1.  | 7.7 | 31        |
| 31 | Identification of FAM46C As a Multiple Myeloma Repressor. Blood, 2015, 126, 836-836.  | 0.6 | 2         |
| 32 | M3P Sequencing Panel Identifies TP53 Mutational Status As a Prognostic Factor in Chemotherapy-Naive<br>Multiple Myeloma. Blood, 2015, 126, 2984-2984.   | 0.6 | 0         |
| 33 | Mutations in Driver Genes and Changes in Clonal Dynamics Are Associated with Shorter Time to Treatment in MBL Cases. Blood, 2015, 126, 5264-5264.   | 0.6 | 0         |
| 34 | CXCR4 expression of multiple myeloma as a dynamic process: influence of therapeutic agents. Leukemia and Lymphoma, 0, , 1-10.   | 0.6 | 0         |