## Eric M J Bindels

List of Publications by Year in descending order

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FRIC M I RINDELS

#	Article	IF	CITATIONS
1	SARS-CoV-2 infects the human kidney and drives fibrosis in kidney organoids. Cell Stem Cell, 2022, 29, 217-231.e8.	11.1	146
2	Kidney Organoids Are Capable of Forming Tumors, but Not Teratomas. Stem Cells, 2022, 40, 577-591.	3.2	3
3	A pipeline for copy number profiling of single circulating tumour cells to assess intrapatient tumour heterogeneity. Molecular Oncology, 2022, 16, 2981-3000.	4.6	6
4	Mapping the cardiac vascular niche in heart failure. Nature Communications, 2022, 13, .	12.8	31
5	Heterogeneous bone-marrow stromal progenitors drive myelofibrosis via a druggable alarmin axis. Cell Stem Cell, 2021, 28, 637-652.e8.	11.1	92
6	Extracellular Vesicles Derived From Adult and Fetal Bone Marrow Mesenchymal Stromal Cells Differentially Promote ex vivo Expansion of Hematopoietic Stem and Progenitor Cells. Frontiers in Bioengineering and Biotechnology, 2021, 9, 640419.	4.1	10
7	Selective Requirement of MYB for Oncogenic Hyperactivation of a Translocated Enhancer in Leukemia. Cancer Discovery, 2021, 11, 2868-2883.	9.4	25
8	Essential role for Cata2 in modulating lineage output from hematopoietic stem cells in zebrafish. Blood Advances, 2021, 5, 2687-2700.	5.2	21
9	The leukemic oncogene EVI1 hijacks a MYC super-enhancer by CTCF-facilitated loops. Nature Communications, 2021, 12, 5679.	12.8	31
10	Detection of Aneuploidy in Cerebrospinal Fluid from Patients with Breast Cancer Can Improve Diagnosis of Leptomeningeal Metastases. Clinical Cancer Research, 2021, 27, 2798-2806.	7.0	14
11	Causal integration of multiâ€omics data with prior knowledge to generate mechanistic hypotheses. Molecular Systems Biology, 2021, 17, e9730.	7.2	78
12	Decoding myofibroblast origins in human kidney fibrosis. Nature, 2021, 589, 281-286.	27.8	380
13	Malignant Transformation Involving CXXC4 Mutations Identified in a Leukemic Progression Model of Severe Congenital Neutropenia. Cell Reports Medicine, 2020, 1, 100074.	6.5	11
14	Identification of osteolineage cellâ€derived extracellular vesicle cargo implicated in hematopoietic support. FASEB Journal, 2020, 34, 5435-5452.	0.5	10
15	Activation of NF-κB driven inflammatory programs in mesenchymal elements attenuates hematopoiesis in low-risk myelodysplastic syndromes. Leukemia, 2019, 33, 536-541.	7.2	31
16	PML Plays a Key Role in Severe Congenital Neutropenia with Mutant ELANE Causing Neutrophil Elastase Protein Misfolding. Blood, 2019, 134, 213-213.	1.4	1
17	Allele-Specific Expression of GATA2 in AML with CEBPA Biallelic Mutations. Blood, 2019, 134, 1235-1235.	1.4	0
18	Temporal autoregulation during human PU.1 locus SubTAD formation. Blood, 2018, 132, 2643-2655.	1.4	12

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19	Complex 3q26/EVI1 Rearrangements Genocopy Inv(3)/t(3;3) Acute Myeloid Leukemias By Enhancer Hijacking, EVI1 Overexpression, Absent MDS1-EVI1 and Low GATA2 Expression. Blood, 2018, 132, 2766-2766.	1.4	ο
20	Mesenchymal Inflammation Drives Genotoxic Stress in Hematopoietic Stem Cells and Predicts Disease Evolution in Human Pre-leukemia. Cell Stem Cell, 2016, 19, 613-627.	11.1	277
21	Inflammatory Niche Signalling Drives Genotoxic Stress in Hematopoietic Stem Cells and Predicts Leukemic Evolution in Human Leukemia Predisposition Syndromes. Blood, 2016, 128, 428-428.	1.4	Ο
22	Mutational spectrum of myeloid malignancies with inv(3)/t(3;3) reveals a predominant involvement of RAS/RTK signaling pathways. Blood, 2015, 125, 133-139.	1.4	86
23	A Single Oncogenic Enhancer Rearrangement Causes Concomitant EVI1 and GATA2 Deregulation in Leukemia. Cell, 2014, 157, 369-381.	28.9	571
24	EVI1 is critical for the pathogenesis of a subset of MLL-AF9–rearranged AMLs. Blood, 2012, 119, 5838-5849.	1.4	76