Hannes Ponstingl

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genetic associations at regulatory phenotypes improve fine-mapping of causal variants for 12 immune-mediated diseases. Nature Genetics, 2022, 54, 251-262.	21.4	23
2	Genetic perturbation of PU.1 binding and chromatin looping at neutrophil enhancers associates with autoimmune disease. Nature Communications, 2021, 12, 2298.	12.8	32
3	SETBP1 overexpression acts in the place of class-defining mutations to drive FLT3-ITD–mutant AML. Blood Advances, 2021, 5, 2412-2425.	5.2	10
4	The Polygenic and Monogenic Basis of Blood Traits and Diseases. Cell, 2020, 182, 1214-1231.e11.	28.9	388
5	PiggyBac mutagenesis and exome sequencing identify genetic driver landscapes and potential therapeutic targets of EGFR-mutant gliomas. Genome Biology, 2020, 21, 181.	8.8	18
6	ATM orchestrates the DNA-damage response to counter toxic non-homologous end-joining at broken replication forks. Nature Communications, 2019, 10, 87.	12.8	133
7	SRPK1 maintains acute myeloid leukemia through effects on isoform usage of epigenetic regulators including BRD4. Nature Communications, 2018, 9, 5378.	12.8	60
8	Genome-wide transposon screening and quantitative insertion site sequencing for cancer gene discovery in mice. Nature Protocols, 2017, 12, 289-309.	12.0	41
9	A single-copy Sleeping Beauty transposon mutagenesis screen identifies new PTEN-cooperating tumor suppressor genes. Nature Genetics, 2017, 49, 730-741.	21.4	53
10	Promoter-bound METTL3 maintains myeloid leukaemia by m6A-dependent translation control. Nature, 2017, 552, 126-131.	27.8	833
11	A CRISPR Dropout Screen Identifies Genetic Vulnerabilities and Therapeutic Targets in Acute Myeloid Leukemia. Cell Reports, 2016, 17, 1193-1205.	6.4	556
12	Leukemia-Associated Somatic Mutations Drive Distinct Patterns of Age-Related Clonal Hemopoiesis. Cell Reports, 2015, 10, 1239-1245.	6.4	443
13	A conditional piggyBac transposition system for genetic screening in mice identifies oncogenic networks in pancreatic cancer. Nature Genetics, 2015, 47, 47-56.	21.4	77
14	A Genetic Progression Model of BrafV600E-Induced Intestinal Tumorigenesis Reveals Targets for Therapeutic Intervention. Cancer Cell, 2013, 24, 15-29.	16.8	183