Silvia Bresolin

List of Publications by Year in descending order

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51	3,046	21 h-index	46
papers	citations		g-index
53	53	53	6139
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Neonatal Manifestations of Chronic Granulomatous Disease: MAS/HLH and Necrotizing Pneumonia as Unusual Phenotypes and Review of the Literature. Journal of Clinical Immunology, 2022, 42, 299-311.	2.0	11
2	<i>NUP214–ABL1</i> fusion in childhood Tâ€ALL. Pediatric Blood and Cancer, 2022, 69, e29643.	0.8	4
3	Customized bioreactor enables the production of 3D diaphragmatic constructs influencing matrix remodeling and fibroblast overgrowth. Npj Regenerative Medicine, 2022, 7, 25.	2.5	5
4	Targeting mesenchymal stromal cells plasticity to reroute acute myeloid leukemia course. Blood, 2021, 138, 557-570.	0.6	26
5	Ruxolitinib as a Novel Therapeutic Option for Poor Prognosis T-LBL Pediatric Patients. Cancers, 2021, 13, 3724.	1.7	2
6	iWhale: a computational pipeline based on Docker and SCons for detection and annotation of somatic variants in cancer WES data. Briefings in Bioinformatics, 2021, 22, .	3.2	8
7	Case Report: Intestinal Nodular Lymphoid Hyperplasia as First Manifestation of Activated PI3Kδ Syndrome Due to a Novel PIK3CD Variant. Frontiers in Pediatrics, 2021, 9, 703056.	0.9	5
8	A novel germline variant in <scp><i>PIK3R1</i></scp> results in <scp>SHORT</scp> syndrome associated with <scp><i>TAL</i></scp> <i>/i><i>LMO</i> Tâ€cell acute lymphoblastic leukemia. American Journal of Hematology, 2020, 95, E335-E338.</i>	2.0	11
9	CircRNAs Dysregulated in Juvenile Myelomonocytic Leukemia: CircMCTP1 Stands Out. Frontiers in Cell and Developmental Biology, 2020, 8, 613540.	1.8	12
10	Posttranslational Regulation of the Exon Skipping Machinery Controls Aberrant Splicing in Leukemia. Cancer Discovery, 2020, 10, 1388-1409.	7.7	37
11	USP7 Cooperates with NOTCH1 to Drive the Oncogenic Transcriptional Program in T-Cell Leukemia. Clinical Cancer Research, 2019, 25, 222-239.	3. 2	66
12	Pre-Clinical Efficacy of the Novel Kinase Inhibitor Nintedanib on PAX5 Fusion Genes in Pediatric Ph-like B-Cell Precursor Acute Lymphoblastic Leukemia. Blood, 2019, 134, 745-745.	0.6	0
13	Somatic mutations activating Wiskott-Aldrich syndrome protein concomitant with RAS pathway mutations in juvenile myelomonocytic leukemia patients. Human Mutation, 2018, 39, 579-587.	1.1	16
14	The long non-coding RNA landscape in juvenile myelomonocytic leukemia. Haematologica, 2018, 103, e501-e504.	1.7	13
15	Differences in circulating microRNA signature in Prader–Willi syndrome and non-syndromic obesity. Endocrine Connections, 2018, 7, 1262-1274.	0.8	10
16	SRC/ABL inhibition disrupts CRLF2-driven signaling to induce cell death in B-cell acute lymphoblastic leukemia. Oncotarget, 2018, 9, 22872-22885.	0.8	11
17	Glucocorticoid resistance is reverted by LCK inhibition in pediatric T-cell acute lymphoblastic leukemia. Blood, 2017, 130, 2750-2761.	0.6	54
18	ZNF521 sustains the differentiation block in MLL-rearranged acute myeloid leukemia. Oncotarget, 2017, 8, 26129-26141.	0.8	21

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19	Induction of Expandable Tissue-Specific Stem/Progenitor Cells through Transient Expression of YAP/TAZ. Cell Stem Cell, 2016, 19, 725-737.	5.2	204
20	LIN28B overexpression defines a novel fetal-like subgroup of juvenile myelomonocytic leukemia. Blood, 2016, 127, 1163-1172.	0.6	48
21	Deciphering KRAS and NRAS mutated clone dynamics in MLL-AF4 paediatric leukaemia by ultra deep sequencing analysis. Scientific Reports, 2016, 6, 34449.	1.6	20
22	LIN28B is over-expressed in specific subtypes of pediatric leukemia and regulates lncRNA H19. Haematologica, 2016, 101, e240-e244.	1.7	18
23	GSK3 Deficiencies in Hematopoietic Stem Cells Initiate Pre-neoplastic State that Is Predictive of Clinical Outcomes of Human Acute Leukemia. Cancer Cell, 2016, 29, 61-74.	7.7	52
24	MicroRNA fingerprints in juvenile myelomonocytic leukemia (JMML) identified miR-150-5p as a tumor suppressor and potential target for treatment. Oncotarget, 2016, 7, 55395-55408.	0.8	30
25	Mutations of <i>SETBP1</i> and <i>JAK3</i> in juvenile myelomonocytic leukemia: a report from the Italian AIEOP study group. Oncotarget, 2016, 7, 28914-28919.	0.8	21
26	Refinement of IKZF1 status in pediatric Philadelphia-positive acute lymphoblastic leukemia. Leukemia, 2015, 29, 2107-2110.	3.3	18
27	Aerobic glycolysis tunes <scp>YAP</scp> / <scp>TAZ</scp> transcriptional activity. EMBO Journal, 2015, 34, 1349-1370.	3.5	306
28	Different outcomes of allogeneic hematopoietic stem cell transplant in a pair of twins affected by juvenile myelomonocytic leukemia. International Journal of Hematology, 2014, 99, 208-212.	0.7	7
29	YAP/TAZ Incorporation in the \hat{I}^2 -Catenin Destruction Complex Orchestrates the Wnt Response. Cell, 2014, 158, 157-170.	13.5	873
30	MLL-AF6 fusion oncogene sequesters AF6 into the nucleus to trigger RAS activation in myeloid leukemia. Blood, 2014, 124, 263-272.	0.6	41
31	Epigenetic Silencing of TFPI-2 in Canine Diffuse Large B-Cell Lymphoma. PLoS ONE, 2014, 9, e92707.	1.1	33
32	Array-Based Comparative Genomic Hybridization Analysis Reveals Chromosomal Copy Number Aberrations Associated with Clinical Outcome in Canine Diffuse Large B-Cell Lymphoma. PLoS ONE, 2014, 9, e111817.	1.1	25
33	Low $\langle i \rangle$ PKCα $\langle i \rangle$ expression within the MRD-HR stratum defines a new subgroup of childhood T-ALL with very poor outcome. Oncotarget, 2014, 5, 5234-5245.	0.8	20
34	Wnt activation promotes neuronal differentiation of Glioblastoma. Cell Death and Disease, 2013, 4, e500-e500.	2.7	89
35	Patient-derived induced pluripotent stem cells recapitulate hematopoietic abnormalities of juvenile myelomonocytic leukemia. Blood, 2013, 121, 4925-4929.	0.6	104
36	Validation of flow cytometric phospho-STAT5 as a diagnostic tool for juvenile myelomonocytic leukemia. Blood Cancer Journal, 2013, 3, e160-e160.	2.8	35

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37	MicroRNA-34b promoter hypermethylation induces CREB overexpression and contributes to myeloid transformation. Haematologica, 2013, 98, 602-610.	1.7	42
38	Mesenchymal stem cells from Shwachman–Diamond syndrome patients display normal functions and do not contribute to hematological defects. Blood Cancer Journal, 2012, 2, e94-e94.	2.8	17
39	Role of TAZ as Mediator of Wnt Signaling. Cell, 2012, 151, 1443-1456.	13.5	419
40	Gene expression signatures of pediatric myelodysplastic syndromes are associated with risk of evolution into acute myeloid leukemia. Leukemia, 2012, 26, 1717-1719.	3.3	6
41	DNA methyltransferase 3a hot-spot locus is not mutated in pediatric patients affected by acute myeloid or T-cell acute lymphoblastic leukemia: an Italian study. Haematologica, 2011, 96, 1886-1887.	1.7	11
42	Enforced expression of MLL-AF4 fusion in cord blood CD34+ cells enhances the hematopoietic repopulating cell function and clonogenic potential but is not sufficient to initiate leukemia. Blood, 2011, 117, 4746-4758.	0.6	84
43	MLL partner genes drive distinct gene expression profiles and genomic alterations in pediatric acute myeloid leukemia: an AIEOP study. Leukemia, 2011, 25, 560-563.	3.3	31
44	The Interlaboratory RObustness of Next-generation sequencing (IRON) study: a deep sequencing investigation of TET2, CBL and KRAS mutations by an international consortium involving 10 laboratories. Leukemia, 2011, 25, 1840-1848.	3.3	96
45	Phenotypical and Functional Characterization of Mesenchymal Stem Cells Derived From Patients Affected by Schwachman-Diamond Syndrome. Blood, 2011, 118, 1336-1336.	0.6	0
46	Lack of Protein Kinase C Alpha Is Associated with Poor Prognosis in Pediatric T-Lineage Acute Lymphoblastic Leukemia. Blood, 2011, 118, 744-744.	0.6	0
47	Evolution of Sub-Clones with KRAS Mutations In Pediatric Patients with MLL-AF4 Rearrangements. Blood, 2011, 118, 2454-2454.	0.6	6
48	Gene Expression–Based Classification As an Independent Predictor of Clinical Outcome in Juvenile Myelomonocytic Leukemia. Journal of Clinical Oncology, 2010, 28, 1919-1927.	0.8	74
49	The Interlaboratory RObustness of Next-Generation Sequencing (IRON) Study: Deep-Sequencing Investigating TET2, CBL, and KRAS Mutations In 4464 Amplicons by An International Group Involving 8 Laboratories Blood, 2010, 116, 1665-1665.	0.6	0
50	STAT5 Phosphorylation Status by Flow Cytometry Is a Rapid and Reliable Tool for Diagnosis and Follow-up of Juvenile Myelomonocytic Leukemia. Blood, 2010, 116, 2751-2751.	0.6	0
51	Gene Expression Profile Analysis of Pediatric MDS Patients Correlates with FAB Classification and Has Prognostic Relevance. Blood, 2008, 112, 2695-2695.	0.6	О