Wilhelm Dirks

List of Publications by Year in descending order

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WILLELM DIDKS

#	Article	IF	CITATIONS
1	Cell line authentication: a necessity for reproducible biomedical research. EMBO Journal, 2022, 41, .	7.8	19
2	Modulators of histone demethylase JMJD1C selectively target leukemic stem cells. FEBS Open Bio, 2021, 11, 265-277.	2.3	5
3	Cross contamination meets misclassification: Awakening of <scp>CHP</scp> â€100 from sleeping beauty sleep—A reviewed model for Ewing's sarcoma. International Journal of Cancer, 2021, 148, 2608-2613.	5.1	2
4	First report on establishment and characterization of a carcinosarcoma tumour cell line model of the bladder. Scientific Reports, 2021, 11, 6030.	3.3	5
5	Ethical Challenges Using Human Tumor Cell Lines in Cancer Research. Recent Results in Cancer Research, 2021, 218, 39-46.	1.8	0
6	Small molecular modulators of JMJD1C preferentially inhibit growth of leukemia cells. International Journal of Cancer, 2020, 146, 400-412.	5.1	29
7	DNMT3A R882H mutation in acute myeloid leukemia cell line SET-2. Leukemia Research, 2020, 88, 106270.	0.8	1
8	Cell Lines as Biological Models: Practical Steps for More Reliable Research. Chemical Research in Toxicology, 2019, 32, 1733-1736.	3.3	10
9	Differential Requirements for the RAD51 Paralogs in Genome Repair and Maintenance in Human Cells. PLoS Genetics, 2019, 15, e1008355.	3.5	39
10	Intact-Cell MALDI-ToF Mass Spectrometry for the Authentication of Drug-Adapted Cancer Cell Lines. Cells, 2019, 8, 1194.	4.1	3
11	Epstein-Barr virus (EBV) activates NKL homeobox gene HLX in DLBCL. PLoS ONE, 2019, 14, e0216898.	2.5	17
12	The LL-100 panel: 100 cell lines for blood cancer studies. Scientific Reports, 2019, 9, 8218.	3.3	74
13	High level EGFR amplification in a newly established glioblastoma cell line 170-MG-BA. Neoplasma, 2019, 66, 109-117.	1.6	2
14	KDM3B shows tumor-suppressive activity and transcriptionally regulates <i>HOXA1</i> through retinoic acid response elements in acute myeloid leukemia. Leukemia and Lymphoma, 2018, 59, 204-213.	1.3	25
15	Authentication of M14 melanoma cell line proves misidentification of MDAâ€MBâ€435 breast cancer cell line. International Journal of Cancer, 2018, 142, 561-572.	5.1	37
16	RBFOX2 and alternative splicing in B-cell lymphoma. Blood Cancer Journal, 2018, 8, 77.	6.2	11
17	Fetal bovine serum (FBS): Past – present – future. ALTEX: Alternatives To Animal Experimentation, 2018, 35, 99-118.	1.5	231
18	Peripheral T-cell lymphoma cell line T8ML-1 highlights conspicuous targeting of PVRL2 by t(14;19)(q11.2;q13.3). Haematologica, 2017, 102, e356-e359.	3.5	3

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19	False and mycoplasmaâ€contaminated leukemia–lymphoma cell lines: time for a reappraisal. International Journal of Cancer, 2017, 140, 1209-1214.	5.1	40
20	Differential cytotoxicity induced by the Titanium(IV)Salan complex Tc52 in G2-phase independent of DNA damage. BMC Cancer, 2016, 16, 469.	2.6	11
21	Identification of flubendazole as potential anti-neuroblastoma compound in a large cell line screen. Scientific Reports, 2015, 5, 8202.	3.3	68
22	Spatio-temporal regulation of the human licensing factor Cdc6 in replication and mitosis. Cell Cycle, 2015, 14, 1704-1715.	2.6	16
23	BCL6 - regulated by AhR/ARNT and wild-type MEF2B - drives expression of germinal center markers MYBL1 and LMO2. Haematologica, 2015, 100, 801-809.	3.5	13
24	Genomic Landscape of Primary Mediastinal B-Cell Lymphoma Cell Lines. PLoS ONE, 2015, 10, e0139663.	2.5	18
25	Localization of MLH3 at the Centrosomes. International Journal of Molecular Sciences, 2014, 15, 13932-13937.	4.1	4
26	2.5 Quality Control Essentials in Human Cell Culture: Cell Line Cross-contamination and Microbiological Infections. , 2014, , 102-114.		0
27	Association between acquired resistance to PLX4032 (vemurafenib) and ATP-binding cassette transporter expression. BMC Research Notes, 2014, 7, 710.	1.4	13
28	Cell line cross-contamination: WSU-CLL is a known derivative of REH and is unsuitable as a model for chronic lymphocytic leukaemia. Leukemia Research, 2014, 38, 999-1001.	0.8	7
29	Aurora Kinases as Targets in Drug-Resistant Neuroblastoma Cells. PLoS ONE, 2014, 9, e108758.	2.5	39
30	Highâ€ŧhroughput SNPâ€based authentication of human cell lines. International Journal of Cancer, 2013, 132, 308-314.	5.1	172
31	Match criteria for human cell line authentication: Where do we draw the line?. International Journal of Cancer, 2013, 132, 2510-2519.	5.1	148
32	Frameshift-derived neoantigens constitute immunotherapeutic targets for patients with microsatellite-instable haematological malignancies. European Journal of Cancer, 2013, 49, 2587-2595.	2.8	28
33	STR DNA Typing of Human Cell Lines: Detection of Intra- and Interspecies Cross-Contamination. Methods in Molecular Biology, 2013, 946, 27-38.	0.9	54
34	Where have all the cell lines gone?. International Journal of Cancer, 2013, 132, 1232-1234.	5.1	4
35	Kaposi's sarcomaâ€derived cell line SLK is not of endothelial origin, but is a contaminant from a known renal carcinoma cell line. International Journal of Cancer, 2013, 132, 1954-1958.	5.1	80
36	Beware imposters: MAâ€1, a novel MALT lymphoma cell line, is misidentified and corresponds to Pfeiffer, a diffuse large Bâ€cell lymphoma cell line. Genes Chromosomes and Cancer, 2013, 52, 986-988.	2.8	5

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37	Stable expression of MutLÎ ³ in human cells reveals no specific response to mismatched DNA, but distinct recruitment to damage sites. Journal of Cellular Biochemistry, 2013, 114, 2405-2414.	2.6	21
38	BCR–ABL1expression in multiple myeloma cells: A case of mistaken identity?. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E270-E271.	7.1	4
39	U-2932: two clones in one cell line, a tool for the study of clonal evolution. Leukemia, 2013, 27, 1155-1164.	7.2	22
40	Online Verification of Human Cell Line Identity by STR DNA Typing. Methods in Molecular Biology, 2011, 731, 45-55.	0.9	19
41	XRCC4 controls nuclear import and distribution of Ligase IV and exchanges faster at damaged DNA in complex with Ligase IV. DNA Repair, 2011, 10, 1232-1242.	2.8	18
42	53BP1 and MDC1 foci formation in HT-1080 cells for low- and high-LET microbeam irradiations. Radiation and Environmental Biophysics, 2011, 50, 345-352.	1.4	20
43	Online imaging of initial DNA damages at the PTB microbeam. Radiation Protection Dosimetry, 2011, 143, 349-352.	0.8	9
44	Recommendation of short tandem repeat profiling for authenticating human cell lines, stem cells, and tissues. In Vitro Cellular and Developmental Biology - Animal, 2010, 46, 727-732.	1.5	103
45	Cell line crossâ€contamination initiative: An interactive reference database of STR profiles covering common cancer cell lines. International Journal of Cancer, 2010, 126, 303-304.	5.1	83
46	Quality of Cell Products: Authenticity, Identity, Genomic Stability and Status of Differentiation. Transfusion Medicine and Hemotherapy, 2010, 37, 2-2.	1.6	18
47	Hypomethylation and expression of BEX2, IGSF4 and TIMP3 indicative of MLL translocations in Acute Myeloid Leukemia. Molecular Cancer, 2009, 8, 86.	19.2	29
48	One falsehood leads easily to another. International Journal of Cancer, 2008, 122, 2165-2168.	5.1	8
49	Human Leukemia and Lymphoma Cell Lines as Models and Resources. Current Medicinal Chemistry, 2008, 15, 339-359.	2.4	45
50	The oncoprotein NPM-ALK of anaplastic large-cell lymphoma induces JUNB transcription via ERK1/2 and JunB translation via mTOR signaling. Blood, 2007, 110, 3374-3383.	1.4	90
51	Cell line OCI/AML3 bears exon-12 NPM gene mutation-A and cytoplasmic expression of nucleophosmin. Leukemia, 2005, 19, 1760-1767.	7.2	139
52	Short tandem repeat DNA typing provides an international reference standard for authentication of human cell lines. ALTEX: Alternatives To Animal Experimentation, 2005, 22, 103-9.	1.5	51
53	Expression ofHOXGenes in Acute Leukemia Cell Lines with and withoutMLLTranslocations. Leukemia and Lymphoma, 2004, 45, 567-574.	1.3	51
54	Tumor necrosis factor receptor-associated factor (TRAF) 4 is a new binding partner for the p70S6 serine/threonine kinase. Leukemia Research, 2003, 27, 687-694.	0.8	28

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55	False leukemia–lymphoma cell lines: an update on over 500 cell lines. Leukemia, 2003, 17, 416-426.	7.2	187
56	BLADDER CARCINOMA CELL LINE ECV304 IS NOT A MODEL SYSTEM FOR ENDOTHELIAL CELLS. In Vitro Cellular and Developmental Biology - Animal, 2002, 38, 185.	1.5	36
57	Expression and functional analysis of the anaplastic lymphoma kinase (ALK) gene in tumor cell lines. International Journal of Cancer, 2002, 100, 49-56.	5.1	110
58	DNA profiling and cytogenetic analysis of cell line WSU-CLL reveal cross-contamination with cell line REH (pre B-ALL). Leukemia, 2002, 16, 1868-1870.	7.2	12
59	Short tandem repeat profiling provides an international reference standard for human cell lines. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 8012-8017.	7.1	428
60	Cross-contamination: HS-Sultan is not a myeloma but a Burkitt lymphoma cell line. Blood, 2001, 98, 3495-3496.	1.4	15
61	Tumor necrosis factor-α–induced proliferation requires synthesis of granulocyte-macrophage colony-stimulating factor. Experimental Hematology, 2000, 28, 1008-1015.	0.4	15
62	The Human Tartrate-Resistant Acid Phosphatase (TRAP): Involvement of the Hemin Responsive Elements (HRE) in Transcriptional Regulation. Leukemia and Lymphoma, 2000, 36, 603-612.	1.3	6
63	False human hematopoietic cell lines: cross-contaminations and misinterpretations. Leukemia, 1999, 13, 1601-1607.	7.2	113
64	ECV304 (endothelial) is really T24 (bladder carcinoma): Cell line cross-contamination at source. In Vitro Cellular and Developmental Biology - Animal, 1999, 35, 558-559.	1.5	128
65	Expression of the growth arrest-specific gene 6 (GAS6) in leukemia and lymphoma cell lines. Leukemia Research, 1999, 23, 643-651.	0.8	35
66	Widespread intraspecies cross-contamination of human tumor cell lines arising at source. , 1999, 83, 555-563.		321
67	Identity of original and late passage Dami megakaryocytes with HEL erythroleukemia cells shown by combined cytogenetics and DNA fingerprinting. Leukemia, 1997, 11, 2032-2038.	7.2	35
68	Expression and function of CD95 (FAS/APO-1) in leukaemia-lymphoma tumour lines. British Journal of Haematology, 1997, 96, 584-593.	2.5	64
69	A multifunctional vector family for gene expression in mammalian cells. Gene, 1994, 149, 387-388.	2.2	37
70	A new hybrid promoter directs transcription at identical start points in mammalian cells and in vitro. Gene, 1994, 149, 389-390.	2.2	13
71	Dicistronic transcription units for gene expression in mammalian cells. Gene, 1993, 128, 247-249.	2.2	170
72	DSMZCellDive: Diving into high-throughput cell line data. F1000Research, 0, 11, 420.	1.6	3