

Joost L Kluiver

List of Publications by Year in descending order

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Version: 2024-02-01

68
papers

3,099
citations

186265

28
h-index

155660

55
g-index

72
all docs

72
docs citations

72
times ranked

4706
citing authors

#	ARTICLE	IF	CITATIONS
1	Actionability of on-target ALK Resistance Mutations in Patients With Non-Small Cell Lung Cancer: Local Experience and Review of the Literature. <i>Clinical Lung Cancer</i> , 2022, 23, e104-e115.	2.6	13
2	The Role of the MYC/miR-150/MYB/ZDHHC11 Network in Hodgkin Lymphoma and Diffuse Large B-Cell Lymphoma. <i>Genes</i> , 2022, 13, 227.	2.4	3
3	Non-small cell lung cancer infiltrated with chronic myelomonocytic leukaemia: a molecular diagnostic challenge to recognise mixed cancers in a single biopsy. <i>Histopathology</i> , 2021, 78, 1043-1046.	2.9	2
4	MiR-378a-3p Is Critical for Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020, 12, 3546.	3.7	12
5	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020, 12, 1464.	3.7	19
6	Non-Coding RNAs in Cancer Radiosensitivity: MicroRNAs and lncRNAs as Regulators of Radiation-Induced Signaling Pathways. <i>Cancers</i> , 2020, 12, 1662.	3.7	44
7	Long Non-coding RNAs in the Development and Maintenance of Lymphoid Malignancies. , 2019, , 127-149.		0
8	A super-SILAC based proteomics analysis of diffuse large B-cell lymphoma-NOS patient samples to identify new proteins that discriminate GCB and non-GCB lymphomas. <i>PLoS ONE</i> , 2019, 14, e0223260.	2.5	4
9	Current Smoking is Associated with Decreased Expression of miR-335-5p in Parenchymal Lung Fibroblasts. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5176.	4.1	15
10	Age-related gene and miRNA expression changes in airways of healthy individuals. <i>Scientific Reports</i> , 2019, 9, 3765.	3.3	34
11	NGS-Based High-Throughput Screen to Identify MicroRNAs Regulating Growth of B-Cell Lymphoma. <i>Methods in Molecular Biology</i> , 2019, 1956, 269-282.	0.9	3
12	Marked TGF- β 2-regulated miRNA expression changes in both COPD and control lung fibroblasts. <i>Scientific Reports</i> , 2019, 9, 18214.	3.3	16
13	Intricate crosstalk between MYC and non-coding RNA regulates hallmarks of cancer. <i>Molecular Oncology</i> , 2019, 13, 26-45.	4.6	45
14	AAV8-mediated gene transfer of microRNA-132 improves beta cell function in mice fed a high-fat diet. <i>Journal of Endocrinology</i> , 2019, 240, 123-132.	2.6	12
15	Argonaute 2 RNA Immunoprecipitation Reveals Distinct miRNA Targetomes of Primary Burkitt Lymphoma Tumors and Normal B Cells. <i>American Journal of Pathology</i> , 2018, 188, 1289-1299.	3.8	7
16	Circulating miRNAs in patients with Barrett's esophagus, high-grade dysplasia and esophageal adenocarcinoma. <i>Journal of Gastrointestinal Oncology</i> , 2018, 9, 1150-1156.	1.4	11
17	Differential miRNA Expression Profiles in Cumulus and Mural Granulosa Cells from Human Pre-ovulatory Follicles. <i>MicroRNA (Sharjah, United Arab Emirates)</i> , 2018, 8, 61-67.	1.2	23
18	Tuberous sclerosis complex is required for tumor maintenance in MYC-driven Burkitt's lymphoma. <i>EMBO Journal</i> , 2018, 37, .	7.8	10

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19	MicroRNA High Throughput Loss-of-Function Screening Reveals an Oncogenic Role for miR-21-5p in Hodgkin Lymphoma. <i>Cellular Physiology and Biochemistry</i> , 2018, 49, 144-159.	1.6	20
20	Involvement of MicroRNAs in the Aging-Related Decline of CD28 Expression by Human T Cells. <i>Frontiers in Immunology</i> , 2018, 9, 1400.	4.8	13
21	Age-related gene and microRNA expression changes in the airways of healthy individuals. , 2018, , .		0
22	Argonaute 2 immunoprecipitation revealed large tumor suppressor kinase 1 as a novel proapoptotic target of miR-21 in T cells. <i>FEBS Journal</i> , 2017, 284, 555-567.	4.7	7
23	miR-24-3p Is Overexpressed in Hodgkin Lymphoma and Protects Hodgkin and Reed-Sternberg Cells from Apoptosis. <i>American Journal of Pathology</i> , 2017, 187, 1343-1355.	3.8	46
24	ZDHHC11 and ZDHHC11B are critical novel components of the oncogenic MYC-miR-150-MYB network in Burkitt lymphoma. <i>Leukemia</i> , 2017, 31, 1470-1473.	7.2	39
25	Emerging roles for long noncoding RNAs in B-cell development and malignancy. <i>Critical Reviews in Oncology/Hematology</i> , 2017, 120, 77-85.	4.4	37
26	MicroRNA profiling of human primary macrophages exposed to dengue virus identifies miRNA-3614-5p as antiviral and regulator of ADAR1 expression. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005981.	3.0	43
27	Identification of transforming growth factor-beta-regulated microRNAs and the microRNA-targetomes in primary lung fibroblasts. <i>PLoS ONE</i> , 2017, 12, e0183815.	2.5	34
28	Target gene identification of TGF- β -induced miR-455-3p and miR-21-3p in lung fibroblasts. , 2017, , .		0
29	Small RNA sequencing reveals a comprehensive miRNA signature of BRCA1-associated high-grade serous ovarian cancer. <i>Journal of Clinical Pathology</i> , 2016, 69, 979-985.	2.0	11
30	Analysis of serum immune markers in seropositive and seronegative rheumatoid arthritis and in high-risk seropositive arthralgia patients. <i>Scientific Reports</i> , 2016, 6, 26021.	3.3	44
31	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cell-Specific Long Noncoding RNAs. <i>American Journal of Pathology</i> , 2016, 186, 2462-2472.	3.8	36
32	Functional Studies on Primary Tubular Epithelial Cells Indicate a Tumor Suppressor Role of SETD2 in Clear Cell Renal Cell Carcinoma. <i>Neoplasia</i> , 2016, 18, 339-346.	5.3	23
33	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. <i>Oncotarget</i> , 2016, 7, 2391-2400.	1.8	43
34	Long noncoding RNAs as a novel component of the Myc transcriptional network. <i>FASEB Journal</i> , 2015, 29, 2338-2346.	0.5	67
35	Immuno-miRs: critical regulators of T-cell development, function and ageing. <i>Immunology</i> , 2015, 144, 1-10.	4.4	141
36	Mir-155 Enhances B-Cell Lymphoma Growth By Targeting TBRC1. <i>Blood</i> , 2015, 126, 4820-4820.	1.4	14

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37	Age-Associated Differences in MiRNA Signatures Are Restricted to CD45RO Negative T Cells and Are Associated with Changes in the Cellular Composition, Activation and Cellular Ageing. PLoS ONE, 2015, 10, e0137556.	2.5	23
38	T-cell Activation Induces Dynamic Changes in miRNA Expression Patterns in CD4 and CD8 T-cell Subsets. MicroRNA (Sharjah, United Arab Emirates), 2015, 4, 117-122.	1.2	37
39	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. Leukemia, 2014, 28, 2248-2251.	7.2	74
40	Comprehensive analysis of miRNA expression in T-cell subsets of rheumatoid arthritis patients reveals defined signatures of naive and memory Tregs. Genes and Immunity, 2014, 15, 115-125.	4.1	111
41	The entire miR-200 seed family is strongly deregulated in clear cell renal cell cancer compared to the proximal tubular epithelial cells of the kidney. Genes Chromosomes and Cancer, 2013, 52, 165-173.	2.8	26
42	Studying MicroRNAs in Lymphoma. Methods in Molecular Biology, 2013, 971, 265-276.	0.9	17
43	Long Non-Coding RNAs Are Commonly Deregulated In Hodgkin Lymphoma. Blood, 2013, 122, 628-628.	1.4	2
44	Dual Role of miR-21 in CD4+ T-Cells: Activation-Induced miR-21 Supports Survival of Memory T-Cells and Regulates CCR7 Expression in Naive T-Cells. PLoS ONE, 2013, 8, e76217.	2.5	61
45	Cellular Localization and Processing of Primary Transcripts of Exonic MicroRNAs. PLoS ONE, 2013, 8, e76647.	2.5	24
46	Long Non-Coding RNAs As Components Of The MYC Network In B Cell Lymphoma. Blood, 2013, 122, 1260-1260.	1.4	0
47	Generation of miRNA sponge constructs. Methods, 2012, 58, 113-117.	3.8	95
48	Rapid Generation of MicroRNA Sponges for MicroRNA Inhibition. PLoS ONE, 2012, 7, e29275.	2.5	125
49	MicroRNAs regulate B-cell receptor signaling-induced apoptosis. Genes and Immunity, 2012, 13, 239-244.	4.1	27
50	MiR-17/106b seed family regulates p21 in Hodgkin's lymphoma. Journal of Pathology, 2011, 225, 609-617.	4.5	35
51	MiRNA profiling in B non-Hodgkin lymphoma: a MYC-related miRNA profile characterizes Burkitt lymphoma. British Journal of Haematology, 2010, 149, 896-899.	2.5	71
52	Global correlation of genome and transcriptome changes in classical Hodgkin lymphoma. Hematological Oncology, 2007, 25, 21-29.	1.7	24
53	Regulation of pri-microRNA BIC transcription and processing in Burkitt lymphoma. Oncogene, 2007, 26, 3769-3776.	5.9	131
54	BCL6 alternative breakpoint region break and homozygous deletion of 17q24 in the nodular lymphocyte predominance type of Hodgkin's lymphoma-derived cell line DEV. Human Pathology, 2006, 37, 675-683.	2.0	29

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55	Gene expression analysis of dendritic/Langerhans cells and Langerhans cell histiocytosis. Journal of Pathology, 2006, 209, 474-483.	4.5	27
56	The role of microRNAs in normal hematopoiesis and hematopoietic malignancies. Leukemia, 2006, 20, 1931-1936.	7.2	92
57	Lack of <i>BIC</i> and microRNA miR-155 expression in primary cases of Burkitt lymphoma. Genes Chromosomes and Cancer, 2006, 45, 147-153.	2.8	219
58	Regulation of pri-miRNA BIC Transcription and Processing in Burkitt Lymphoma.. Blood, 2006, 108, 2380-2380.	1.4	0
59	Report: workshop on mediastinal grey zone lymphoma. European Journal of Haematology, 2005, 75, 45-52.	2.2	19
60	BIC and miR-155 are highly expressed in Hodgkin, primary mediastinal and diffuse large B cell lymphomas. Journal of Pathology, 2005, 207, 243-249.	4.5	640
61	Polymorphisms and Lack of or Aberrant Expression of HLA Class I and II May Influence Antigen Presentation in Classical Hodgkin Lymphoma.. Blood, 2005, 106, 20-20.	1.4	1
62	BIC and miR-155 Are Highly Expressed in Hodgkin, Primary Mediastinal and Diffuse Large B Cell Lymphomas.. Blood, 2005, 106, 970-970.	1.4	3
63	Lack of BIC and microRNA miR-155 Expression in Primary Cases of Burkitt Lymphoma.. Blood, 2005, 106, 1922-1922.	1.4	0
64	High Expression of Micro-RNA BIC / miR155 in All Subtypes of Hodgkin Lymphoma.. Blood, 2004, 104, 430-430.	1.4	1
65	Serial Analysis of Gene Expression Revealed Consistent Downregulation of More Than 100 Genes in Hodgkin Lymphoma.. Blood, 2004, 104, 4288-4288.	1.4	0
66	High expression of B cell receptor inducible gene <i>BIC</i> in all subtypes of Hodgkin lymphoma. Genes Chromosomes and Cancer, 2003, 37, 20-28.	2.8	224
67	Common and differential chemokine expression patterns in rs cells of NLP, EBV positive and negative classical hodgkin lymphomas. International Journal of Cancer, 2002, 99, 665-672.	5.1	66
68	Hitting the brake: miR-31 regulates CD8 T cell effector function. Non-coding RNA Investigation, 0, 1, 8-8.	0.6	0