Joost L Kluiver

List of Publications by Year
in descending order

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68	3,099	185998 28	155451 55
papers	citations	h-index	g-index

72	72	72	4706
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	BIC and miR-155 are highly expressed in Hodgkin, primary mediastinal and diffuse large B cell lymphomas. Journal of Pathology, 2005, 207, 243-249.	2.1	640
2	High expression of B-cell receptor inducible geneBIC in all subtypes of Hodgkin lymphoma. Genes Chromosomes and Cancer, 2003, 37, 20-28.	1.5	224
3	Lack ofBIC and microRNA miR-155 expression in primary cases of Burkitt lymphoma. Genes Chromosomes and Cancer, 2006, 45, 147-153.	1.5	219
4	Immunoâ€miRs: critical regulators of Tâ€cell development, function and ageing. Immunology, 2015, 144, 1-10.	2.0	141
5	Regulation of pri-microRNA BIC transcription and processing in Burkitt lymphoma. Oncogene, 2007, 26, 3769-3776.	2.6	131
6	Rapid Generation of MicroRNA Sponges for MicroRNA Inhibition. PLoS ONE, 2012, 7, e29275.	1.1	125
7	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.	2.2	111
8	Generation of miRNA sponge constructs. Methods, 2012, 58, 113-117.	1.9	95
9	The role of microRNAs in normal hematopoiesis and hematopoietic malignancies. Leukemia, 2006, 20, 1931-1936.	3.3	92
10	The mutational landscape of Hodgkin lymphoma cell lines determined by whole-exome sequencing. Leukemia, 2014, 28, 2248-2251.	3.3	74
11	MiRNA profiling in B nonâ€Hodgkin lymphoma: a MYCâ€related miRNA profile characterizes Burkitt lymphoma. British Journal of Haematology, 2010, 149, 896-899.	1.2	71
12	Long noncoding RNAs as a novel component of the Myc transcriptional network. FASEB Journal, 2015, 29, 2338-2346.	0.2	67
13	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.	2.3	66
14	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.	1.1	61
15	miR-24-3p Is Overexpressed in Hodgkin Lymphoma and Protects Hodgkin and Reed-Sternberg Cells from Apoptosis. American Journal of Pathology, 2017, 187, 1343-1355.	1.9	46
16	Intricate crosstalk between <scp>MYC</scp> and nonâ€coding <scp>RNA</scp>s regulates hallmarks of cancer. Molecular Oncology, 2019, 13, 26-45.	2.1	45
17	Analysis of serum immune markers in seropositive and seronegative rheumatoid arthritis and in high-risk seropositive arthralgia patients. Scientific Reports, 2016, 6, 26021.	1.6	44
18	Non-Coding RNAs in Cancer Radiosensitivity: MicroRNAs and lncRNAs as Regulators of Radiation-Induced Signaling Pathways. Cancers, 2020, 12, 1662.	1.7	44

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19	MicroRNA profiling of human primary macrophages exposed to dengue virus identifies miRNA-3614-5p as antiviral and regulator of ADAR1 expression. PLoS Neglected Tropical Diseases, 2017, 11, e0005981.	1.3	43
20	Inhibition of the miR-155 target NIAM phenocopies the growth promoting effect of miR-155 in B-cell lymphoma. Oncotarget, 2016, 7, 2391-2400.	0.8	43
21	ZDHHC11 and ZDHHC11B are critical novel components of the oncogenic MYC-miR-150-MYB network in Burkitt lymphoma. Leukemia, 2017, 31, 1470-1473.	3.3	39
22	Emerging roles for long noncoding RNAs in B-cell development and malignancy. Critical Reviews in Oncology/Hematology, 2017, 120, 77-85.	2.0	37
23	T-cell Activation Induces Dynamic Changes in miRNA Expression Patterns in CD4 and CD8 T-cell Subsets. MicroRNA (Shariqah, United Arab Emirates), 2015, 4, 117-122.	0.6	37
24	Long Noncoding RNA Expression Profiling in Normal B-Cell Subsets and Hodgkin Lymphoma Reveals Hodgkin and Reed-Sternberg Cellâ€“Specific Long Noncoding RNAs. American Journal of Pathology, 2016, 186, 2462-2472.	1.9	36
25	MiRâ€17/106b seed family regulates p21 in Hodgkin's lymphoma. Journal of Pathology, 2011, 225, 609-617.	2.1	35
26	Age-related gene and miRNA expression changes in airways of healthy individuals. Scientific Reports, 2019, 9, 3765.	1.6	34
27	Identification of transforming growth factor-beta-regulated microRNAs and the microRNA-targetomes in primary lung fibroblasts. PLoS ONE, 2017, 12, e0183815.	1.1	34
28	BCL6 alternative breakpoint region break and homozygous deletion of 17q24 in the nodular lymphocyte predominance type of Hodgkin's lymphomaâ€“derived cell line DEV. Human Pathology, 2006, 37, 675-683.	1.1	29
29	Gene expression analysis of dendritic/Langerhans cells and Langerhans cell histiocytosis. Journal of Pathology, 2006, 209, 474-483.	2.1	27
30	MicroRNAs regulate B-cell receptor signaling-induced apoptosis. Genes and Immunity, 2012, 13, 239-244.	2.2	27
31	The entire miRâ€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.	1.5	26
32	Global correlation of genome and transcriptome changes in classical Hodgkin lymphoma. Hematological Oncology, 2007, 25, 21-29.	0.8	24
33	Cellular Localization and Processing of Primary Transcripts of Exonic MicroRNAs. PLoS ONE, 2013, 8, e76647.	1.1	24
34	Functional Studies on Primary Tubular Epithelial Cells Indicate a Tumor Suppressor Role of SETD2 in Clear Cell Renal Cell Carcinoma. Neoplasia, 2016, 18, 339-346.	2.3	23
35	Differential miRNA Expression Profiles in Cumulus and Mural Granulosa Cells from Human Pre-ovulatory Follicles. MicroRNA (Shariqah, United Arab Emirates), 2018, 8, 61-67.	0.6	23
36	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.	1.1	23

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37	MicroRNA High Throughput Loss-of-Function Screening Reveals an Oncogenic Role for miR-21-5p in Hodgkin Lymphoma. Cellular Physiology and Biochemistry, 2018, 49, 144-159.	1.1	20
38	Report: workshop on mediastinal grey zone lymphoma. European Journal of Haematology, 2005, 75, 45-52.	1.1	19
39	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. Cancers, 2020, 12, 1464.	1.7	19
40	Studying MicroRNAs in Lymphoma. Methods in Molecular Biology, 2013, 971, 265-276.	0.4	17
41	Marked TGF-Î²-regulated miRNA expression changes in both COPD and control lung fibroblasts. Scientific Reports, 2019, 9, 18214.	1.6	16
42	Current Smoking is Associated with Decreased Expression of miR-335-5p in Parenchymal Lung Fibroblasts. International Journal of Molecular Sciences, 2019, 20, 5176.	1.8	15
43	Mir-155 Enhances B-Cell Lymphoma Growth By Targeting TBRG1. Blood, 2015, 126, 4820-4820.	0.6	14
44	Involvement of MicroRNAs in the Aging-Related Decline of CD28 Expression by Human T Cells. Frontiers in Immunology, 2018, 9, 1400.	2.2	13
45	Actionability of on-target ALK Resistance Mutations in Patients With Non-Small Cell Lung Cancer: Local Experience and Review of the Literature. Clinical Lung Cancer, 2022, 23, e104-e115.	1.1	13
46	MiR-378a-3p Is Critical for Burkitt Lymphoma Cell Growth. Cancers, 2020, 12, 3546.	1.7	12
47	AAV8-mediated gene transfer of microRNA-132 improves beta cell function in mice fed a high-fat diet. Journal of Endocrinology, 2019, 240, 123-132.	1.2	12
48	Small RNA sequencing reveals a comprehensive miRNA signature ofBRCA1-associated high-grade serous ovarian cancer. Journal of Clinical Pathology, 2016, 69, 979-985.	1.0	11
49	Circulating miRNAs in patients with Barrettâ€™s esophagus, high-grade dysplasia and esophageal adenocarcinoma. Journal of Gastrointestinal Oncology, 2018, 9, 1150-1156.	0.6	11
50	Tuberous sclerosis complex is required for tumor maintenance in MYCâ€driven Burkitt's lymphoma. EMBO Journal, 2018, 37, .	3.5	10
51	Argonaute 2 immunoprecipitation revealed large tumor suppressor kinase 1 as a novel proapoptotic target of miRâ€21 in T cells. FEBS Journal, 2017, 284, 555-567.	2.2	7
52	Argonaute 2 RNA Immunoprecipitation Reveals Distinct miRNA Targetomes of Primary Burkitt Lymphoma Tumors and Normal B Cells. American Journal of Pathology, 2018, 188, 1289-1299.	1.9	7
53	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. PLoS ONE, 2019, 14, e0223260.	1.1	4
54	NGS-Based High-Throughput Screen to Identify MicroRNAs Regulating Growth of B-Cell Lymphoma. Methods in Molecular Biology, 2019, 1956, 269-282.	0.4	3

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55	BIC and miR-155 Are Highly Expressed in Hodgkin, Primary Mediastinal and Diffuse Large B Cell Lymphomas.. Blood, 2005, 106, 970-970.	0.6	3
56	The Role of the MYC/miR-150/MYB/ZDHHC11 Network in Hodgkin Lymphoma and Diffuse Large B-Cell Lymphoma. Genes, 2022, 13, 227.	1.0	3
57	Nonâ€smallâ€cell lung cancer infiltrated with chronic myelomonocytic leukaemia: a molecular diagnostic challenge to recognise mixed cancers in a single biopsy. Histopathology, 2021, 78, 1043-1046.	1.6	2
58	Long Non-Coding RNAs Are Commonly Deregulated In Hodgkin Lymphoma. Blood, 2013, 122, 628-628.	0.6	2
59	High Expression of Micro-RNA BIC / miR155 in All Subtypes of Hodgkin Lymphoma.. Blood, 2004, 104, 430-430.	0.6	1
60	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.	0.6	1
61	Long Non-coding RNAs in the Development and Maintenance of Lymphoid Malignancies. , 2019, , 127-149.		0
62	Serial Analysis of Gene Expression Revealed Consistent Downregulation of More Than 100 Genes in Hodgkin Lymphoma.. Blood, 2004, 104, 4288-4288.	0.6	0
63	Lack of BIC and microRNA miR-155 Expression in Primary Cases of Burkitt Lymphoma.. Blood, 2005, 106, 1922-1922.	0.6	0
64	Regulation of pri-miRNA BIC Transcription and Processing in Burkitt Lymphoma.. Blood, 2006, 108, 2380-2380.	0.6	0
65	Long Non-Coding RNAs As Components Of The MYC Network In B Cell Lymphoma. Blood, 2013, 122, 1260-1260.	0.6	0
66	Hitting the brake: miR-31 regulates CD8 T cell effector function. Non-coding RNA Investigation, 0, 1, 8-8.	0.6	0
67	Target gene identification of TGF-Î²-induced miR-455-3p and miR-21-3p in lung fibroblasts. , 2017, , .		0
68	Age-related gene and microRNA expression changes in the airways of healthy individuals. , 2018, , .		0