Lynn R Goldin

List of Publications by Year
in descending order

106	6,629	57758	64796
papers	citations	44 h-index	79 g-index

106	106	106	8028
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Detectable clonal mosaicism and its relationship to aging and cancer. Nature Genetics, 2012, 44, 651-658.	21.4	519
2	Rare missense variants in POT1 predispose to familial cutaneous malignant melanoma. Nature Genetics, 2014, 46, 482-486.	21.4	283
3	Risk of monoclonal gammopathy of undetermined significance (MGUS) and subsequent multiple myeloma among African American and white veterans in the United States. Blood, 2006, 107, 904-906.	1.4	280
4	Chronic Immune Stimulation Might Act As a Trigger for the Development of Acute Myeloid Leukemia or Myelodysplastic Syndromes. Journal of Clinical Oncology, 2011, 29, 2897-2903.	1.6	239
5	Increased risks of polycythemia vera, essential thrombocythemia, and myelofibrosis among 24â€‰577 first-degree relatives of 11â€‰039 patients with myeloproliferative neoplasms in Sweden. Blood, 2008, 112, 2199-2204.	1.4	226
6	Arterial and venous thrombosis in monoclonal gammopathy of undetermined significance and multiple myeloma: a population-based study. Blood, 2010, 115, 4991-4998.	1.4	204
7	Familial risk of lymphoproliferative tumors in families of patients with chronic lymphocytic leukemia: results from the Swedish Family-Cancer Database. Blood, 2004, 104, 1850-1854.	1.4	189
8	Autoimmunity and Susceptibility to Hodgkin Lymphoma: A Population-Based Caseâ€“Control Study in Scandinavia. Journal of the National Cancer Institute, 2006, 98, 1321-1330.	6.3	179
9	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. Nature Genetics, 2013, 45, 868-876.	21.4	179
10	Familial aggregation of Hodgkin lymphoma and related tumors. Cancer, 2004, 100, 1902-1908.	4.1	155
11	Autoimmunity and the risk of myeloproliferative neoplasms. Haematologica, 2010, 95, 1216-1220.	3.5	151
12	Risk of lymphoproliferative disorders among first-degree relatives of lymphoplasmacytic lymphoma/WaldenstrÃ¶m macroglobulinemia patients: a population-based study in Sweden. Blood, 2008, 112, 3052-3056.	1.4	143
13	Prevalence of Monoclonal Gammopathy of Undetermined Significance Among Men in Ghana. Mayo Clinic Proceedings, 2007, 82, 1468-1473.	3.0	142
14	Obesity is associated with an increased risk of monoclonal gammopathy of undetermined significance among black and white women. Blood, 2010, 116, 1056-1059.	1.4	137
15	Risk of plasma cell and lymphoproliferative disorders among 14621 first-degree relatives of 4458 patients with monoclonal gammopathy of undetermined significance in Sweden. Blood, 2009, 114, 791-795.	1.4	133
16	Familial characteristics of autoimmune and hematologic disorders in 8,406 multiple myeloma patients: A population-based case-control study. International Journal of Cancer, 2006, 118, 3095-3098.	5.1	125
17	Genome-wide association study identifies a novel susceptibility locus at 6p21.3 among familial CLL. Blood, 2011, 117, 1911-1916.	1.4	118
18	B-cell monoclonal lymphocytosis and B-cell abnormalities in the setting of familial B-cell chronic lymphocytic leukemia. Cytometry, 2003, 52B, 1-12.	1.8	114

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19	Elevated risk of chronic lymphocytic leukemia and other indolent non-Hodgkin's lymphomas among relatives of patients with chronic lymphocytic leukemia. Haematologica, 2009, 94, 647-653.	3.5	113
20	Autoimmune disease in individuals and close family members and susceptibility to nonâ€Hodgkin's lymphoma. Arthritis and Rheumatism, 2008, 58, 657-666.	6.7	106
21	Ascertainment and diagnostic accuracy for hematopoietic lymphoproliferative malignancies in Sweden 1964â€“2003. International Journal of Cancer, 2007, 121, 2260-2266.	5.1	104
22	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. American Journal of Human Genetics, 2015, 96, 487-497.	6.2	101
23	Patterns of survival and causes of death following a diagnosis of monoclonal gammopathy of undetermined significance: a population-based study. Haematologica, 2009, 94, 1714-1720.	3.5	95
24	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. Nature Communications, 2016, 7, 10933.	12.8	94
25	Respiratory tract infections and subsequent risk of chronic lymphocytic leukemia. Blood, 2007, 109, 2198-2201.	1.4	89
26	Autoimmunity and lymphomagenesis. International Journal of Cancer, 2009, 124, 1497-1502.	5.1	89
27	Lung Cancer Prognosis Before and After Recurrence in a Population-Based Setting. Journal of the National Cancer Institute, 2015, 107, djv059.	6.3	86
28	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	12.8	86
29	Highly increased familial risks for specific lymphoma subtypes. British Journal of Haematology, 2009, 146, 91-94.	2.5	85
30	Immune-Related and Inflammatory Conditions and Risk of Lymphoplasmacytic Lymphoma or Waldenstrom Macroglobulinemia. Journal of the National Cancer Institute, 2010, 102, 557-567.	6.3	83
31	A high-density SNP genome-wide linkage search of 206 families identifies susceptibility loci for chronic lymphocytic leukemia. Blood, 2007, 110, 3326-3333.	1.4	79
32	Common variation at 6p21.31 (BAK1) influences the risk of chronic lymphocytic leukemia. Blood, 2012, 120, 843-846.	1.4	76
33	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. Nature Communications, 2017, 8, 14175.	12.8	75
34	Genomewide Linkage Screen for WaldenstrÃ¶m Macroglobulinemia Susceptibility Loci in High-Risk Families. American Journal of Human Genetics, 2006, 79, 695-701.	6.2	72
35	Clinical Characteristics of Familial B-CLL in the National Cancer Institute Familial Registry. Leukemia and Lymphoma, 2001, 42, 99-108.	1.3	71
36	Patterns of autoimmunity and subsequent chronic lymphocytic leukemia in Nordic countries. Blood, 2006, 108, 292-296.	1.4	63

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37	Patterns of hematologic malignancies and solid tumors among 37,838 firstâ€degree relatives of 13,896 patients with multiple myeloma in Sweden. International Journal of Cancer, 2009, 125, 2147-2150.	5.1	63
38	Familial Aggregation and Heterogeneity of Non-Hodgkin Lymphoma in Population-Based Samples. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 2402-2406.	2.5	61
39	Identification of a novel chromosome region, 13q21.33-q22.2, for susceptibility genes in familial chronic lymphocytic leukemia. Blood, 2007, 109, 916-925.	1.4	61
40	Common occurrence of monoclonal Bâ€cell lymphocytosis among members of highâ€risk CLL families. British Journal of Haematology, 2010, 151, 152-158.	2.5	61
41	Mutations in a gene encoding a midbody kelch protein in familial and sporadic classical Hodgkin lymphoma lead to binucleated cells. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 14920-14925.	7.1	59
42	No Evidence for Anticipation in Lymphoproliferative Tumors in Population-Based Samples. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 1245-1250.	2.5	51
43	Anticipation in Familial Chronic Lymphocytic Leukemia. American Journal of Human Genetics, 1999, 65, 265-268.	6.2	50
44	A genome scan of 18 families with chronic lymphocytic leukaemia. British Journal of Haematology, 2003, 121, 866-873.	2.5	48
45	Germline mutations in Protection of Telomeres 1 in two families with Hodgkin lymphoma. British Journal of Haematology, 2018, 181, 372-377.	2.5	48
46	CXCR4 expression is associated with survival in familial chronic lymphocytic leukemia, but CD38 expression is not. Blood, 2002, 100, 1100-1101.	1.4	43
47	Whole exome sequencing reveals a C-terminal germline variant in CEBPA-associated acute myeloid leukemia: 45-year follow up of a large family. Haematologica, 2016, 101, 846-852.	3.5	42
48	Whole exome sequencing in families at high risk for Hodgkin lymphoma: identification of a predisposing mutation in the KDR gene. Haematologica, 2016, 101, 853-860.	3.5	40
49	Risk of second malignant neoplasms among lymphoma patients with a family history of cancer. International Journal of Cancer, 2006, 120, 1099-1102.	5.1	39
50	Current Insight on Trends, Causes, and Mechanisms of Hodgkinâ€™s Lymphoma. Cancer Journal (Sudbury,) Tj ETQq0 0 0 rgBT /Overlock	2.0	39
51	Common genetic variants in candidate genes and risk of familial lymphoid malignancies. British Journal of Haematology, 2009, 146, 418-423.	2.5	38
52	Chronic lymphocytic leukaemia genetics overview. British Journal of Haematology, 2007, 139, 630-634.	2.5	36
53	Familial Aggregation of Acute Myeloid Leukemia and Myelodysplastic Syndromes. Journal of Clinical Oncology, 2012, 30, 179-183.	1.6	35
54	Respiratory tract infections in the pathway to multiple myeloma: a population-based study in Scandinavia. Haematologica, 2006, 91, 1697-700.	3.5	35

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55	Familial chronic lymphocytic leukemia. Current Opinion in Hematology, 2010, 17, 350-355.	2.5	33
56	Genetic Susceptibility Variants for Chronic Lymphocytic Leukemia. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1098-1102.	2.5	31
57	Genetic and immune-related factors in the pathogenesis of lymphoproliferative and plasma cell malignancies. Haematologica, 2009, 94, 1581-1589.	3.5	30
58	Evolution of multiple cell clones over a 29-year period of a CLL patient. Nature Communications, 2016, 7, 13765.	12.8	29
59	Autoimmunity and risk for Hodgkin's lymphoma by subtype. Haematologica, 2009, 94, 1468-1469.	3.5	28
60	Perspectives on familial chronic lymphocytic leukemia: genes and the environment. Seminars in Hematology, 2004, 41, 201-206.	3.4	27
61	Familial CLL: Genes and Environment. Hematology American Society of Hematology Education Program, 2007, 2007, 339-345.	2.5	27
62	Genetic Susceptibility to Chronic Lymphocytic Leukemia. Seminars in Hematology, 2013, 50, 296-302.	3.4	26
63	Family studies in chronic lymphocytic leukaemia and other lymphoproliferative tumours. British Journal of Haematology, 2007, 139, 774-779.	2.5	23
64	Sex-related DNA methylation differences in B cell chronic lymphocytic leukemia. Biology of Sex Differences, 2019, 10, 2.	4.1	23
65	No familial aggregation in chronic myeloid leukemia. Blood, 2013, 122, 460-461.	1.4	22
66	Population-based study on the impact of the familial form of WaldenstrÃ¶m macroglobulinemia on overall survival. Blood, 2015, 125, 2174-2175.	1.4	21
67	Association of polygenic risk score with the risk of chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. Blood, 2018, 131, 2541-2551.	1.4	21
68	A genome-wide linkage scan for body mass index on Framingham Heart Study families. BMC Genetics, 2003, 4, S97.	2.7	20
69	History of autoimmune disease is associated with impaired survival in multiple myeloma and monoclonal gammopathy of undetermined significance: a population-based study. Annals of Hematology, 2017, 96, 261-269.	1.8	20
70	Familial Aggregation of Lymphoplasmacytic Lymphoma/WaldenstrÃ¶m Macroglobulinemia with Solid Tumors and Myeloid Malignancies. Acta Haematologica, 2012, 127, 173-177.	1.4	19
71	Telomere length and heavy-chain mutation status in familial chronic lymphocytic leukemia. Leukemia Research, 2002, 26, 791-794.	0.8	18
72	Personal and family history of autoimmune diabetes mellitus and susceptibility to young-adult-onset Hodgkin lymphoma. International Journal of Cancer, 2006, 118, 449-452.	5.1	17

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73	Genetics- and Immune-Related Factors in the Pathogenesis of Lymphoplasmacytic Lymphoma/WaldenstrÃ¶m's Macroglobulinemia. Clinical Lymphoma and Myeloma, 2009, 9, 23-26.	1.4	16
74	Re: Familial Clustering of Hodgkin Lymphoma and Multiple Sclerosis. Journal of the National Cancer Institute, 2005, 97, 543-544.	6.3	15
75	High-density mapping and follow-up studies on chromosomal regions 1, 3, 6, 12, 13 and 17 in 28 families with chronic lymphocytic leukaemia. British Journal of Haematology, 2006, 133, 060120074427018.	2.5	15
76	Precursors to Lymphoproliferative Malignancies. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 533-539.	2.5	15
77	Whole exome sequencing in families with CLL detects a variant in Integrin Î² 2 associated with disease susceptibility. Blood, 2016, 128, 2261-2263.	1.4	15
78	Mosaic chromosome 20q deletions are more frequent in the aging population. Blood Advances, 2017, 1, 380-385.	5.2	15
79	Prevalence of pathogenic/likely pathogenic variants in the 24 cancer genes of the ACMG Secondary Findings v2.0 list in a large cancer cohort and ethnicity-matched controls. Genome Medicine, 2018, 10, 99.	8.2	15
80	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for WaldenstrÃ¶m macroglobulinemia. Nature Communications, 2018, 9, 4182.	12.8	15
81	Regional inference with averagedPvalues increases the power to detect linkage. , 1999, 17, 157-164.		14
82	Analysis of metabolic syndrome phenotypes in Framingham Heart Study families from Genetic Analysis Workshop 13. Genetic Epidemiology, 2003, 25, S78-S89.	1.3	14
83	Risk of solid tumors and myeloid hematological malignancies among first-degree relatives of patients with monoclonal gammopathy of undetermined significance. Haematologica, 2009, 94, 1179-1181.	3.5	14
84	Mosaic 13q14 deletions in peripheral leukocytes of non-hematologic cancer cases and healthy controls. Journal of Human Genetics, 2016, 61, 411-418.	2.3	13
85	Familial Aspects of Chronic Lymphocytic Leukemia, Monoclonal B-Cell Lymphocytosis (MBL), and Related Lymphomas. European Journal of Clinical & Medical Oncology, 2010, 2, 119-126.	0.0	12
86	Association of elevated serumfree light chains with chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. Blood Cancer Journal, 2019, 9, 59.	6.2	9
87	ATM mutations and protein expression are not associated with familial B-CLL cases. Leukemia Research, 2003, 27, 973-975.	0.8	8
88	Infection in infancy and subsequent risk of developing lymphoma in children and young adults. Blood, 2011, 117, 1670-1672.	1.4	8
89	Survival in patients with familial and sporadic myeloproliferative neoplasms. Blood, 2015, 125, 3665-3666.	1.4	8
90	Germline and somatic JAK2 mutations and susceptibility to chronic myeloproliferative neoplasms. Genome Medicine, 2009, 1, 55.	8.2	7

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91	Monoclonal B cell lymphocytosis: Clinical and population perspectives. Cytometry Part B - Clinical Cytometry, 2010, 78B, S115-9.	1.5	7
92	No association of ARLTS1 polymorphisms and risk for familial chronic lymphocytic leukaemia. British Journal of Haematology, 2007, 137, 173-175.	2.5	4
93	KLHDC8B Is a Novel, Mitotically-Regulated Classical Hodgkinâ€™s Lymphoma Candidate Susceptibility Gene.. Blood, 2006, 108, 473-473.	1.4	4
94	GeneÃ—Environment Interaction from Caseâ€Control and Caseâ€Case Approaches. Genetic Epidemiology, 2001, 21, S825-30.	1.3	3
95	Use of Weighted pâ€Values in Regional Inference Procedures. Genetic Epidemiology, 2001, 21, S484-9.	1.3	3
96	Combined somatic mutation and copy number analysis in the survival of familial <scp>CLL</scp>. British Journal of Haematology, 2018, 181, 604-613.	2.5	3
97	In search of genetic factors predisposing to familial hairy cell leukemia (HCL): exome-sequencing of four multiplex HCL pedigrees. Leukemia, 2020, 34, 1934-1938.	7.2	3
98	Comparison of two linkage inference procedures for genes related to the P300 component of the event related potential. Genetic Epidemiology, 1999, 17, S163-7.	1.3	2
99	Monoclonal B-Cell Lymphocytosis Is Commonly Observed Among Unaffected Members of High Risk CLL Families.. Blood, 2009, 114, 1232-1232.	1.4	2
100	Arterial and Venous Thrombosis in Monoclonal Gammopathy of Undetermined Significance and Multiple Myeloma: A Population-Based Study.. Blood, 2009, 114, 1872-1872.	1.4	2
101	Risk of Monoclonal Gammopathy of Undetermined Significance (MGUS) and Subsequent Multiple Myeloma among African-American and White Veterans in the U.S.. Blood, 2005, 106, 1541-1541.	1.4	1
102	Assessment of estimation procedures for risk and onset hazard with dependent data. Genetic Epidemiology, 1999, 17, S97-S102.	1.3	0
103	Introduction: Linkage Analysis of Quantitative Traits. Genetic Epidemiology, 2001, 21, S459-60.	1.3	0
104	Genetics in Lymphomagenesis. , 2013, , 835-847.		0
105	Parental longevity and survival among patients with multiple myeloma and monoclonal gammopathy of undetermined significance: a populationâ€based study. British Journal of Haematology, 2019, 186, 37-44.	2.5	0
106	Circulating B Cell Clones in Familial Waldenstroì†m Macroglobulinemia.. Blood, 2012, 120, 2703-2703.	1.4	0