

# Barbara L Kee

## List of Publications by Year in descending order

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63  
papers

4,037  
citations

147801

31  
h-index

144013

57  
g-index

69  
all docs

69  
docs citations

69  
times ranked

4973  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mature natural killer cell and lymphoid tissueâ€“inducing cell development requires Id2-mediated suppression of E protein activity. <i>Journal of Experimental Medicine</i> , 2007, 204, 1119-1130.	8.5	331
2	The Transcriptional Regulation of B Cell Lineage Commitment. <i>Immunity</i> , 2007, 26, 715-725.	14.3	322
3	E and ID proteins branch out. <i>Nature Reviews Immunology</i> , 2009, 9, 175-184.	22.7	275
4	Induction of Early B Cell Factor (EBF) and Multiple B Lineage Genes by the Basic Helix-Loop-Helix Transcription Factor E12. <i>Journal of Experimental Medicine</i> , 1998, 188, 699-713.	8.5	231
5	Development of innate lymphoid cells. <i>Nature Immunology</i> , 2016, 17, 775-782.	14.5	188
6	E2A Proteins Promote Development of Lymphoid-Primed Multipotent Progenitors. <i>Immunity</i> , 2008, 29, 217-227.	14.3	187
7	Epigenetic repression of the Igk locus by STAT5-mediated recruitment of the histone methyltransferase Ezh2. <i>Nature Immunology</i> , 2011, 12, 1212-1220.	14.5	169
8	Id3 inhibits B lymphocyte progenitor growth and survival in response to TGF- $\beta$ 2. <i>Nature Immunology</i> , 2001, 2, 242-247.	14.5	156
9	NFIL3 Orchestrates the Emergence of Common Helper Innate Lymphoid Cell Precursors. <i>Cell Reports</i> , 2015, 10, 2043-2054.	6.4	154
10	Early B Cell Factor Promotes B Lymphopoiesis with Reduced Interleukin 7 Responsiveness in the Absence of E2A. <i>Journal of Experimental Medicine</i> , 2004, 199, 1689-1700.	8.5	148
11	Gene Deregulation and Chronic Activation in Natural Killer Cells Deficient in the Transcription Factor ETS1. <i>Immunity</i> , 2012, 36, 921-932.	14.3	118
12	SAP Protein-Dependent Natural Killer T-like Cells Regulate the Development of CD8+ T Cells with Innate Lymphocyte Characteristics. <i>Immunity</i> , 2010, 33, 203-215.	14.3	107
13	E2A proteins: essential regulators at multiple stages of B-cell development. <i>Immunological Reviews</i> , 2000, 175, 138-149.	6.0	106
14	Innate Lymphoid Cells Control Early Colonization Resistance against Intestinal Pathogens through ID2-Dependent Regulation of the Microbiota. <i>Immunity</i> , 2015, 42, 731-743.	14.3	102
15	Cryptic activation of an Irf8 enhancer governs cDC1 fate specification. <i>Nature Immunology</i> , 2019, 20, 1161-1173.	14.5	100
16	Inhibitor of DNA Binding 3 Limits Development of Murine Slam-Associated Adaptor Protein-Dependent Innate $\beta$ 1 T cells. <i>PLoS ONE</i> , 2010, 5, e9303.	2.5	83
17	E proteins and the regulation of early lymphocyte development. <i>Immunological Reviews</i> , 2010, 238, 93-109.	6.0	79
18	Essential Functions for ID Proteins at Multiple Checkpoints in Invariant NKT Cell Development. <i>Journal of Immunology</i> , 2013, 191, 5973-5983.	0.8	76

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19	The ETS1 transcription factor is required for the development and cytokine-induced expansion of ILC2. <i>Journal of Experimental Medicine</i> , 2016, 213, 687-696.	8.5	75
20	Gut Microbiota Regulates K/BxN Autoimmune Arthritis through Follicular Helper T but Not Th17 Cells. <i>Journal of Immunology</i> , 2016, 196, 1550-1557.	0.8	72
21	Interferon-producing killer dendritic cells (IKDCs) arise via a unique differentiation pathway from primitive c-kit <sup>hi</sup> CD62L <sup>+</sup> lymphoid progenitors. <i>Blood</i> , 2007, 109, 4825-4931.	1.4	71
22	The transcription factor lymphoid enhancer factor 1 controls invariant natural killer T cell expansion and Th2-type effector differentiation. <i>Journal of Experimental Medicine</i> , 2015, 212, 793-807.	8.5	68
23	E2A transcription factors limit expression of Gata3 to facilitate T lymphocyte lineage commitment. <i>Blood</i> , 2013, 121, 1534-1542.	1.4	65
24	Batf Pioneers the Reorganization of Chromatin in Developing Effector T Cells via Ets1-Dependent Recruitment of Ctcf. <i>Cell Reports</i> , 2019, 29, 1203-1220.e7.	6.4	63
25	IL-7R $\alpha$ and E47: independent pathways required for development of multipotent lymphoid progenitors. <i>EMBO Journal</i> , 2002, 21, 103-113.	7.8	52
26	Notch1 promotes survival of E2A-deficient T cell lymphomas through pre-T cell receptor-dependent and -independent mechanisms. <i>Blood</i> , 2006, 107, 4115-4121.	1.4	51
27	Defining innate and innate-like lymphoid cells. <i>Immunological Reviews</i> , 2014, 261, 177-197.	6.0	48
28	Transcription factor ID2 prevents E proteins from enforcing a naive T lymphocyte gene program during NK cell development. <i>Science Immunology</i> , 2018, 3, .	11.9	47
29	Notch1 co-opts lymphoid enhancer factor 1 for survival of murine T-cell lymphomas. <i>Blood</i> , 2007, 110, 2650-2658.	1.4	45
30	Transcription factor regulation of B lineage commitment. <i>Current Opinion in Immunology</i> , 2001, 13, 180-185.	5.5	40
31	Growth factor independent 1B (Gfi1b) is an E2A target gene that modulates Gata3 in T-cell lymphomas. <i>Blood</i> , 2007, 109, 4406-4414.	1.4	40
32	Transcriptional regulation of lymphocyte development. <i>Current Opinion in Genetics and Development</i> , 2008, 18, 441-448.	3.3	29
33	EZH2 Regulates the Developmental Timing of Effectors of the Pre-T Antigen Receptor Checkpoints. <i>Journal of Immunology</i> , 2017, 198, 4682-4691.	0.8	29
34	In vitro tracking of IL-7 responsiveness and gene expression during commitment of bipotent B-cell/macrophage progenitors. <i>Current Biology</i> , 1996, 6, 1159-1169.	3.9	27
35	Extrinsic and intrinsic regulation of early natural killer cell development. <i>Immunologic Research</i> , 2008, 40, 193-207.	2.9	26
36	Id3 Induces Growth Arrest and Caspase-2-Dependent Apoptosis in B Lymphocyte Progenitors. <i>Journal of Immunology</i> , 2005, 175, 4518-4527.	0.8	25

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37	Transcriptional regulation of natural killer cell development. <i>Current Opinion in Immunology</i> , 2010, 22, 193-198.	5.5	25
38	Differential Roles for the E2A Activation Domains in B Lymphocytes and Macrophages. <i>Journal of Immunology</i> , 2008, 180, 1694-1703.	0.8	22
39	The transcription factor BCL-6 controls early development of innate-like T cells. <i>Nature Immunology</i> , 2020, 21, 1058-1069.	14.5	20
40	Transcriptional and epigenetic regulation of innate-like T lymphocyte development. <i>Current Opinion in Immunology</i> , 2018, 51, 39-45.	5.5	19
41	Inhibitors of DNA Binding Proteins Restrict T Cell Potential by Repressing Notch1 Expression in Flt3-Negative Common Lymphoid Progenitors. <i>Journal of Immunology</i> , 2012, 189, 3822-3830.	0.8	18
42	Murine thymic NK cells are distinct from ILC1s and have unique transcription factor requirements. <i>European Journal of Immunology</i> , 2017, 47, 800-805.	2.9	18
43	The transcriptional repressor ID2 supports natural killer cell maturation by controlling TCF1 amplitude. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	17
44	Repression of <i>Ccr9</i> Transcription in Mouse T Lymphocyte Progenitors by the Notch Signaling Pathway. <i>Journal of Immunology</i> , 2015, 194, 3191-3200.	0.8	16
45	Transcriptional regulation of natural killer cell development and maturation. <i>Advances in Immunology</i> , 2020, 146, 1-28.	2.2	14
46	Combinatorial ETS1-Dependent Control of Oncogenic NOTCH1 Enhancers in T-cell Leukemia. <i>Blood Cancer Discovery</i> , 2020, 1, 178-197.	5.0	11
47	The HOX11/TLX1 Transcription Factor Oncogene Induces Chromosomal Aneuploidy in T-ALL. <i>Blood</i> , 2009, 114, 142-142.	1.4	8
48	Oncogenic and Tumor Suppressor Functions for Lymphoid Enhancer Factor 1 in E2a <sup>-/-</sup> T Acute Lymphoblastic Leukemia. <i>Frontiers in Immunology</i> , 2022, 13, 845488.	4.8	8
49	Genomic and Transcriptional Mechanisms Governing Innate-like T Lymphocyte Development. <i>Journal of Immunology</i> , 2022, 209, 208-216.	0.8	7
50	A Comprehensive Transcriptional Landscape of Human Hematopoiesis. <i>Cell Stem Cell</i> , 2011, 8, 122-124.	11.1	5
51	Analysis of GzmbCre as a Model System for Gene Deletion in the Natural Killer Cell Lineage. <i>PLoS ONE</i> , 2015, 10, e0125211.	2.5	4
52	Cutting Edge: Lymphomyeloid-Primed Progenitor Cell Fates Are Controlled by the Transcription Factor Tal1. <i>Journal of Immunology</i> , 2019, 202, 2837-2842.	0.8	4
53	E Protein Transcription Factors as Suppressors of T Lymphocyte Acute Lymphoblastic Leukemia. <i>Frontiers in Immunology</i> , 2022, 13, 885144.	4.8	4
54	Ezh2 Represses Transcription of Innate Lymphoid Genes in B Lymphocyte Progenitors and Maintains the B-2 Cell Fate. <i>Journal of Immunology</i> , 2020, 204, 1760-1769.	0.8	3

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55	Development of Natural Killer Cells and ILC1. , 2016, , 140-148.		2
56	Applying the TOR(C)QUE in iNKT cells: A new twist in an old tale. European Journal of Immunology, 2017, 47, 454-457.	2.9	2
57	Lncâ€™ing Id2 to ILC1. Immunity, 2017, 47, 389-390.	14.3	2
58	A s-myly Route toward Lymphoid Differentiation. Immunity, 2009, 30, 474-476.	14.3	1
59	Itâ€™s a Phase That EBF1 Is Going Through. Immunity, 2020, 53, 1123-1125.	14.3	1
60	E2A and the Development of B and T Lymphocytes. , 0, , 255-270.		0
61	Identification of Oncogenic Pathways of T-Acute Lymphoblastic Leukemia (T-ALL) through Gene Expression Profiling of Mouse Tumor Models.. Blood, 2006, 108, 2234-2234.	1.4	0
62	Oncogenic Transcriptional Programs Controlled by TLX1/HOX11 and TLX3/HOX11L2 in T-ALL.. Blood, 2009, 114, 676-676.	1.4	0
63	BCL11B Mutations In T-Cell Acute Lymphoblastic Leukemia. Blood, 2010, 116, 471-471.	1.4	0