

# Megan E Mcnerney

## List of Publications by Year in descending order

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Version: 2024-02-01

37  
papers

1,729  
citations

471509

17  
h-index

414414

32  
g-index

37  
all docs

37  
docs citations

37  
times ranked

3165  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Therapy-related myeloid neoplasms: when genetics and environment collide. <i>Nature Reviews Cancer</i> , 2017, 17, 513-527.  | 28.4 | 270       |
| 2  | 2B4 Acts As a Non-MHC Major Histocompatibility Complex Binding Inhibitory Receptor on Mouse Natural Killer Cells. <i>Journal of Experimental Medicine</i> , 2004, 199, 1245-1254.  | 8.5  | 179       |
| 3  | A new self: MHC-class-I-independent Natural-killer-cell self-tolerance. <i>Nature Reviews Immunology</i> , 2005, 5, 363-374.   | 22.7 | 156       |
| 4  | CUX1 is a haploinsufficient tumor suppressor gene on chromosome 7 frequently inactivated in acute myeloid leukemia. <i>Blood</i> , 2013, 121, 975-983.   | 1.4  | 130       |
| 5  | Clinical Validation of a Next-Generation Sequencing Genomic Oncology Panel via Cross-Platform Benchmarking against Established Amplicon Sequencing Assays. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 43-56.                      | 2.8  | 105       |
| 6  | Regulatory Defects in Cbl and Mitogen-Activated Protein Kinase (Extracellular Signal-Related Kinase) Pathways Cause Persistent Hyperexpression of CD40 Ligand in Human Lupus T Cells. <i>Journal of Immunology</i> , 2000, 165, 6627-6634. | 0.8  | 90        |
| 7  | 2B4 (CD244) is a non-MHC binding receptor with multiple functions on natural killer cells and CD8+ T cells. <i>Molecular Immunology</i> , 2005, 42, 489-494.   | 2.2  | 90        |
| 8  | Targeted Disruption of the <i>2B4</i> Gene in Mice Reveals an In Vivo Role of 2B4 (CD244) in the Rejection of B16 Melanoma Cells. <i>Journal of Immunology</i> , 2005, 174, 800-807.   | 0.8  | 88        |
| 9  | Requirement of homotypic NK-cell interactions through 2B4(CD244)/CD48 in the generation of NK effector functions. <i>Blood</i> , 2006, 107, 3181-3188.   | 1.4  | 78        |
| 10 | Bionimbus: a cloud for managing, analyzing and sharing large genomics datasets. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2014, 21, 969-975.   | 4.4  | 66        |
| 11 | Widespread genetic epistasis among cancer genes. <i>Nature Communications</i> , 2014, 5, 4828.   | 12.8 | 63        |
| 12 | Dominant Role of Oncogene Dosage and Absence of Tumor Suppressor Activity in <i>Nras</i> -Driven Hematopoietic Transformation. <i>Cancer Discovery</i> , 2013, 3, 993-1001.  | 9.4  | 60        |
| 13 | Robust stratification of breast cancer subtypes using differential patterns of transcript isoform expression. <i>PLoS Genetics</i> , 2017, 13, e1006589.   | 3.5  | 53        |
| 14 | 2B4 (CD244)-CD48 interactions provide a novel MHC class I-independent system for NK-cell self-tolerance in mice. <i>Blood</i> , 2005, 106, 1337-1340.  | 1.4  | 50        |
| 15 | Cooperative loss of RAS feedback regulation drives myeloid leukemogenesis. <i>Nature Genetics</i> , 2015, 47, 539-543.   | 21.4 | 39        |
| 16 | Gene dosage effect of CUX1 in a murine model disrupts HSC homeostasis and controls the severity and mortality of MDS. <i>Blood</i> , 2018, 131, 2682-2697.   | 1.4  | 36        |
| 17 | The spectrum of somatic mutations in high-risk acute myeloid leukaemia with $\Delta 7/\text{del}(7q)$ . <i>British Journal of Haematology</i> , 2014, 166, 550-556.  | 2.5  | 29        |
| 18 | An Integrated Genomic Approach to the Assessment and Treatment of Acute Myeloid Leukemia. <i>Seminars in Oncology</i> , 2011, 38, 215-224.   | 2.2  | 21        |

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|----|---|------|-----------|
| 19 | The haploinsufficient tumor suppressor, CUX1, acts as an analog transcriptional regulator that controls target genes through distal enhancers that loop to target promoters. <i>Nucleic Acids Research</i> , 2017, 45, 6350-6361.               | 14.5 | 21        |
| 20 | Cytotoxic Therapyâ€™Induced Effects on Both Hematopoietic and Marrow Stromal Cells Promotes Therapy-Related Myeloid Neoplasms. <i>Blood Cancer Discovery</i> , 2020, 1, 32-47.  | 5.0  | 16        |
| 21 | Retroviral insertional mutagenesis identifies the del(5q) genes, CXXC5, TIFAB and ETF1, as well as the Wnt pathway, as potential targets in del(5q) myeloid neoplasms. <i>Haematologica</i> , 2016, 101, e232-e236.                             | 3.5  | 13        |
| 22 | Loss of a 7q gene, <i>CUX1</i> , disrupts epigenetically driven DNA repair and drives therapy-related myeloid neoplasms. <i>Blood</i> , 2021, 138, 790-805.   | 1.4  | 13        |
| 23 | Deficiency of <i>Cux1</i> , Encoded on Human Chromosome 7q, Causes Aberrant Hematopoietic Stem Cell Function and Spontaneous Myeloproliferative Disease in Mice. <i>Blood</i> , 2017, 130, 789-789.   | 1.4  | 10        |
| 24 | A phase 1 study of azacitidine with high-dose cytarabine and mitoxantrone in high-risk acute myeloid leukemia. <i>Blood Advances</i> , 2020, 4, 599-606.  | 5.2  | 9         |
| 25 | System for Informatics in the Molecular Pathology Laboratory. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 522-532.  | 2.8  | 8         |
| 26 | Venetoclax imparts distinct cell death sensitivity and adaptivity patterns in T cells. <i>Cell Death and Disease</i> , 2021, 12, 1005.  | 6.3  | 8         |
| 27 | Development of warm auto- and allo-antibodies in a 3-year old boy with sickle cell haemoglobinopathy following his first transfusion of a single unit of red blood cells. <i>Blood Transfusion</i> , 2010, 8, 126-8.                            | 0.4  | 7         |
| 28 | The significance of CUX1 and chromosome 7 in myeloid malignancies. <i>Current Opinion in Hematology</i> , 2022, 29, 92-102.   | 2.5  | 6         |
| 29 | Deficiency of CUX1, Encoded on 7q, Blocks the Normal HSC DNA Damage Response and Drives Highly Penetrant Therapy-Related Myeloid Neoplasms in Mice. <i>Blood</i> , 2019, 134, 641-641.  | 1.4  | 5         |
| 30 | The Harmful Consequences of Increased Fitness in Hematopoietic Stem Cells. <i>Cell Stem Cell</i> , 2018, 23, 634-635.   | 11.1 | 4         |
| 31 | CRISPR screening in human hematopoietic stem and progenitor cells reveals an enrichment for tumor suppressor genes within chromosome 7 commonly deleted regions. <i>Leukemia</i> , 2022, 36, 1421-1425.   | 7.2  | 3         |
| 32 | Next-Generation Sequencing Analysis of 23 Therapy-Related Acute Myeloid Leukemia Transcriptomes. <i>Blood</i> , 2010, 116, 850-850.   | 1.4  | 2         |
| 33 | Genomic studies controvert the existence of the CUX1 p75 isoform. <i>Scientific Reports</i> , 2022, 12, 151.  | 3.3  | 1         |
| 34 | Natural killer cell subsets in allograft rejection and tolerance. <i>Current Opinion in Organ Transplantation</i> , 2007, 12, 10-16.  | 1.6  | 0         |
| 35 | Retroviral Insertional Mutagenesis In <i>Egr1</i> +/- mice, Haploinsufficient For a Human Del(5q) Myeloid Leukemia Gene, Develop Myeloid Neoplasms With Proviral Insertions In Genes Syntenic To Human 5q. <i>Blood</i> , 2013, 122, 1275-1275. | 1.4  | 0         |
| 36 | Enhancer-Promoter Looping Deciphers Dosage of the Haploinsufficient Transcription Factor, CUX1. <i>Blood</i> , 2016, 128, 2700-2700.  | 1.4  | 0         |

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|----|---|-----|-----------|
| 37 | CUX1 Deficiency Potentiates RAS Signaling to Drive Malignancy. Blood, 2021, 138, 1159-1159. | 1.4 | 0         |