Brenton G Mar

List of Publications by Year in descending order

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RDENTON C. MAD

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Whole exome sequencing of a breast tumor in a patient with Diamond Blackfan anemia. Blood Cells, Molecules, and Diseases, 2021, 89, 102566. | 0.6 | 0 |
| 2 | Psychometric evaluation of the Advanced Systemic Mastocytosis Symptom Assessment Form (AdvSM-SAF). Leukemia Research, 2021, 108, 106606. | 0.4 | 6 |
| 3 | Effective Control of Advance Systemic Mastocytosis with Avapritinib: Mutational Analysis from the Explorer Clinical Study. Blood, 2021, 138, 318-318. | 0.6 | 16 |
| 4 | PIONEER: A Randomized, Double-Blind, Placebo-Controlled, Phase 2 Study of Avapritinib in Patients with Indolent or Smoldering Systemic Mastocytosis (SM) With Symptoms Inadequately Controlled by Standard Therapy. Journal of Allergy and Clinical Immunology, 2020, 145, AB336. | 1.5 | 15 |
| 5 | Pioneer: A Randomized, Double-Blind, Placebo-Controlled, Phase 2 Study of Avapritinib in Patients with Indolent or Smoldering Systemic Mastocytosis with Symptoms Inadequately Controlled with Standard Therapy. Blood, 2019, 134, 2950-2950. | 0.6 | 2 |
| 6 | Diagnosis and Treatment of Acute Myeloid Leukemia in Children. , 2018, , 359-374. | | 0 |
| 7 | PPM1D-truncating mutations confer resistance to chemotherapy and sensitivity to PPM1D inhibition in hematopoietic cells. Blood, 2018, 132, 1095-1105. | 0.6 | 160 |
| 8 | Clonal Hematopoiesis Associated With Adverse Outcomes After Autologous Stem-Cell Transplantation for Lymphoma. Journal of Clinical Oncology, 2017, 35, 1598-1605. | 0.8 | 339 |
| 9 | Deletion of ribosomal protein genes is a common vulnerability in human cancer, especially in concert with <i> <scp>TP</scp> 53 </i> mutations. EMBO Molecular Medicine, 2017, 9, 498-507. | 3.3 | 86 |
| 10 | Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation. New England Journal of Medicine, 2017, 376, 536-547. | 13.9 | 586 |
| 11 | The relative utilities of genome-wide, gene panel, and individual gene sequencing in clinical practice. Blood, 2017, 130, 433-439. | 0.6 | 50 |
| 12 | The EMT regulator ZEB2 is a novel dependency of human and murine acute myeloid leukemia. Blood, 2017, 129, 497-508. | 0.6 | 65 |
| 13 | SETD2 alterations impair DNA damage recognition and lead to resistance to chemotherapy in leukemia. Blood, 2017, 130, 2631-2641. | 0.6 | 102 |
| 14 | Functionally identifiable apoptosis-insensitive subpopulations determine chemoresistance in acute myeloid leukemia. Journal of Clinical Investigation, 2016, 126, 3827-3836. | 3.9 | 40 |
| 15 | Genetic Alterations Predict Outcomes in Patients with Myelodysplastic Syndrome Receiving Allogeneic Hematopoietic Stem Cell Transplantation. Blood, 2016, 128, 69-69. | 0.6 | 2 |
| 16 | Clonal Hematopoiesis Associated with Adverse Outcomes Following Autologous Stem Cell Transplantation for Non-Hodgkin Lymphoma. Blood, 2016, 128, 986-986. | 0.6 | 3 |
| 17 | Acute myeloid leukemia ontogeny is defined by distinct somatic mutations. Blood, 2015, 125, 1367-1376. | 0.6 | 747 |
| 18 | SETD2 Heterozygous Loss in Leukemia Leads to Chemotherapy Resistance through Attenuation of the DNA Damage Response. Blood, 2015, 126, 2626-2626. | 0.6 | 0 |

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|----|---|------|-----------|
| 19 | Age-Related Clonal Hematopoiesis Associated with Adverse Outcomes. New England Journal of Medicine, 2014, 371, 2488-2498. | 13.9 | 3,474 |
| 20 | Somatic Mutations Predict Poor Outcome in Patients With Myelodysplastic Syndrome After Hematopoietic Stem-Cell Transplantation. Journal of Clinical Oncology, 2014, 32, 2691-2698. | 0.8 | 359 |
| 21 | Mutations in epigenetic regulators including SETD2 are gained during relapse in paediatric acute lymphoblastic leukaemia. Nature Communications, 2014, 5, 3469. | 5.8 | 171 |
| 22 | Ontogeny-Specific Patterns of Genetic Alterations in Acute Myeloid Leukemia. Blood, 2014, 124, 18-18. | 0.6 | 0 |
| 23 | Clonal Hematopoiesis with Somatic Mutations Is a Common, Age-Related Condition Associated with Adverse Outcomes. Blood, 2014, 124, 840-840. | 0.6 | 1 |
| 24 | Sequencing histone-modifying enzymes identifies UTX mutations in acute lymphoblastic leukemia. Leukemia, 2012, 26, 1881-1883. | 3.3 | 70 |
| 25 | The controversial role of the Hedgehog pathway in normal and malignant hematopoiesis. Leukemia, 2011, 25, 1665-1673. | 3.3 | 58 |
| 26 | Self-renewal related signaling in myeloid leukemia stem cells. International Journal of Hematology, 2011, 94, 109-117. | 0.7 | 41 |
| 27 | CCR7 signalling as an essential regulator of CNS infiltration in T-cell leukaemia. Nature, 2009, 459, 1000-1004. | 13.7 | 227 |
| 28 | Knockdown of CCR7 or Its Ligands Causes a Loss of Central Nervous System Involvement in Notch1 Induced T-ALL. Blood, 2008, 112, 199-199. | 0.6 | 4 |
| 29 | Genetic Engineering and Significant Ex-Vivo Expansion of Cord Blood Natural Killer Cells: Implications for Post-Transplant Adoptive Cellular Immunotherapy. Blood, 2008, 112, 209-209. | 0.6 | 1 |
| 30 | Membrane-Associated and Secreted Genes in Breast Cancer. Cancer Research, 2004, 64, 8682-8687. | 0.4 | 17 |
| 31 | Novel transcription factors in human CD34 antigen–positive hematopoietic cells. Blood, 2002, 100, 107-119 | 0.6 | 32 |
| 32 | A novel nuclear protein, 5qNCA (LOC51780) is a candidate for the myeloid leukemia tumor suppressor gene on chromosome 5 band q31. Oncogene, 2001, 20, 6946-6954. | 2.6 | 71 |