

Lukasz P Gondek

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

807
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858243

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1288
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#	ARTICLE	IF	CITATIONS
1	Nonmyeloablative Allogeneic Transplantation With Post-Transplant Cyclophosphamide for Acute Myeloid Leukemia With IDH Mutations: A Single Center Experience. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2022, 22, 260-269.	0.2	4
2	Donor Clonal Hematopoiesis and Recipient Outcomes After Transplantation. <i>Journal of Clinical Oncology</i> , 2022, 40, 189-201.	0.8	79
3	Post-Transplantation Cyclophosphamide-Based Graft- versus-Host Disease Prophylaxis with Nonmyeloablative Conditioning for Blood or Marrow Transplantation for Myelofibrosis. <i>Transplantation and Cellular Therapy</i> , 2022, 28, 259.e1-259.e11.	0.6	11
4	Impact of diagnostic genetics on remission MRD and transplantation outcomes in older patients with AML. <i>Blood</i> , 2022, 139, 3546-3557.	0.6	37
5	Genomic landscape of myelodysplastic/myeloproliferative neoplasm can predict response to hypomethylating agent therapy. <i>Leukemia and Lymphoma</i> , 2022, 63, 1942-1948.	0.6	8
6	The role of the atypical chemokine receptor CCRL2 in myelodysplastic syndrome and secondary acute myeloid leukemia. <i>Science Advances</i> , 2022, 8, eab18952.	4.7	7
7	Allelic complexity of <i>KMT2A</i> partial tandem duplications in acute myeloid leukemia and myelodysplastic syndromes. <i>Blood Advances</i> , 2022, 6, 4236-4240.	2.5	6
8	Clonal Hematopoiesis and the Risk of Hematologic Malignancies after Curative Therapies for Sickle Cell Disease. <i>Journal of Clinical Medicine</i> , 2022, 11, 3160.	1.0	2
9	Whole-exome sequencing identifies functional classes of gene mutations associated with bone marrow failure in pediatric Fanconi Anemia patients. <i>European Journal of Haematology</i> , 2021, 107, 293-294.	1.1	0
10	Deep learning for diagnosis of acute promyelocytic leukemia via recognition of genomically imprinted morphologic features. <i>Npj Precision Oncology</i> , 2021, 5, 38.	2.3	18
11	Gender-related differences in the outcomes and genomic landscape of patients with myelodysplastic syndrome/myeloproliferative neoplasm overlap syndromes. <i>British Journal of Haematology</i> , 2021, 193, 1142-1150.	1.2	21
12	Clonal hematopoiesis and bone marrow failure syndromes. <i>Best Practice and Research in Clinical Haematology</i> , 2021, 34, 101273.	0.7	6
13	CHIP: is clonal hematopoiesis a surrogate for aging and other disease?. <i>Hematology American Society of Hematology Education Program</i> , 2021, 2021, 384-389.	0.9	8
14	Assessing clonal haematopoiesis: clinical burdens and benefits of diagnosing myelodysplastic syndrome precursor states. <i>Lancet Haematology</i> , 2020, 7, e73-e81.	2.2	45
15	Stem cell donors should be screened for CHIP. <i>Blood Advances</i> , 2020, 4, 784-788.	2.5	28
16	The diagnostic utility of targeted gene panel sequencing in discriminating etiologies of cytopenia. <i>American Journal of Hematology</i> , 2019, 94, 1141-1148.	2.0	33
17	Fibrinogen consumption and use of heparin are risk factors for delayed bleeding during acute promyelocytic leukemia induction. <i>Leukemia Research</i> , 2019, 83, 106174.	0.4	8
18	Hedgehog/GLI1 activation leads to leukemic transformation of myelodysplastic syndrome in vivo and GLI1 inhibition results in antitumor activity. <i>Oncogene</i> , 2019, 38, 687-698.	2.6	21

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19	Hotspot SF3B1 mutations induce metabolic reprogramming and vulnerability to serine deprivation. <i>Journal of Clinical Investigation</i> , 2019, 129, 4708-4723.	3.9	41
20	Anthracycline-induced acute myocarditis and ventricular fibrillation arrest. <i>American Journal of Hematology</i> , 2018, 93, 469-470.	2.0	4
21	Donor cell leukemia arising from clonal hematopoiesis after bone marrow transplantation. <i>Leukemia</i> , 2016, 30, 1916-1920.	3.3	79
22	Integration of Hedgehog and mutant FLT3 signaling in myeloid leukemia. <i>Science Translational Medicine</i> , 2015, 7, 291ra96.	5.8	50
23	I Walk the Line: How to Tell MDS From Other Bone Marrow Failure Conditions. <i>Current Hematologic Malignancy Reports</i> , 2014, 9, 389-399.	1.2	9
24	250K Single Nucleotide Polymorphism Array Karyotyping Identifies Acquired Uniparental Disomy and Homozygous Mutations, Including Novel Missense Substitutions of <i>c-Cbl</i> , in Myeloid Malignancies. <i>Cancer Research</i> , 2008, 68, 10349-10357.	0.4	282