# Kostas Stamatopoulos

List of Publications by Year
in descending order



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|---|---|---|---|
| 333 papers | 9,286 citations | 47006 47 h-index | 49909 87 g-index |
| 335 all docs | 335 docs citations | 335 times ranked | 8882 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Precision diagnostics in lymphomas â€“ Recent developments and future directions. Seminars in Cancer Biology, 2022, 84, 170-183. | 9.6 | 13 |
| 2 | The Genomics of Hairy Cell Leukaemia and Splenic Diffuse Red Pulp Lymphoma. Cancers, 2022, 14, 697. | 3.7 | 9 |
| 3 | Cytogenetics in Chronic Lymphocytic Leukemia: ERIC Perspectives and Recommendations. HemaSphere, 2022, 6, e707. | 2.7 | 17 |
| 4 | The TÎ”p63/BCL2 axis represents a novel mechanism of clinical aggressiveness in chronic lymphocytic leukemia. Blood Advances, 2022, 6, 2646-2656. | 5.2 | 1 |
| 5 | Chronic Graft-Versus-Host Disease Immunoprofiling Reveals T Cell Clonal Dynamics That Correlate with Disease Activity: A Novel Molecular Markerâ€?. Transplantation and Cellular Therapy, 2022, 28, S273-S274. | 1.2 | 0 |
| 6 | Immunoglobulin gene sequence analysis in chronic lymphocytic leukemia: the 2022 update of the recommendations by ERIC, the European Research Initiative on CLL. Leukemia, 2022, 36, 1961-1968. | 7.2 | 34 |
| 7 | The EHA Research Roadmap: Malignant Lymphoid Diseases. HemaSphere, 2022, 6, e726. | 2.7 | 1 |
| 8 | Three-dimensional co-culture model of chronic lymphocytic leukemia bone marrow microenvironment predicts patient-specific response to mobilizing agents. Haematologica, 2021, 106, 2334-2344. | 3.5 | 18 |
| 9 | Comparative analysis of targeted next-generation sequencing panels for the detection of gene mutations in chronic lymphocytic leukemia: an ERIC multi-center study. Haematologica, 2021, 106, 682-691. | 3.5 | 10 |
| 10 | Stem cell factor is implicated in microenvironmental interactions and cellular dynamics of chronic lymphocytic leukemia. Haematologica, 2021, 106, 692-700. | 3.5 | 4 |
| 11 | Infrequent â€œchronic lymphocytic leukemia-specificâ€•immunoglobulin stereotypes in aged individuals with or without low-count monoclonal B-cell lymphocytosis. Haematologica, 2021, 106, 1178-1181. | 3.5 | 8 |
| 12 | Higher-order connections between stereotyped subsets: implications for improved patient classification in CLL. Blood, 2021, 137, 1365-1376. | 1.4 | 72 |
| 13 | Triggering interferon signaling in T cells with avadomide sensitizes CLL to anti-PD-L1/PD-1 immunotherapy. Blood, 2021, 137, 216-231. | 1.4 | 40 |
| 14 | Higher-order immunoglobulin repertoire restrictions in CLL: the illustrative case of stereotyped subsets 2 and 169. Blood, 2021, 137, 1895-1904. | 1.4 | 21 |
| 15 | Exploiting B-cell Receptor Stereotypy to Design Tailored Immunotherapy in Chronic Lymphocytic Leukemia. Clinical Cancer Research, 2021, 27, 729-739. | 7.0 | 5 |
| 16 | The Calcitriol/Vitamin D Receptor System Regulates Key Immune Signaling Pathways in Chronic Lymphocytic Leukemia. Cancers, 2021, 13, 285. | 3.7 | 3 |
| 17 | Control of PD-L1 expression in CLL-cells by stromal triggering of the Notch-c-Myc-EZH2 oncogenic signaling axis. , 2021, 9, e001889. | | 15 |
| 18 | MyPal-Child study protocol: an observational prospective clinical feasibility study of the MyPal ePRO-based early palliative care digital system in paediatric oncology patients. BMJ Open, 2021, 11, e045226. | 1.9 | 9 |

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| 19 | Myeloid-derived suppressor cell subtypes differentially influence T-cell function, T-helper subset differentiation, and clinical course in CLL. Leukemia, 2021, 35, 3163-3175. | 7.2 | 25 |
| 20 | In CLL, epigenetics also points to the BCR. Blood, 2021, 137, 2863-2865. | 1.4 | 0 |
| 21 | T Cell Defects and Immunotherapy in Chronic Lymphocytic Leukemia. Cancers, 2021, 13, 3255. | 3.7 | 6 |
| 22 | MyPal: Designing and Evaluating Digital Patient-Reported Outcome Systems for Cancer Palliative Care in Europe. Journal of Palliative Medicine, 2021, 24, 962-964. | 1.1 | 5 |
| 23 | RPS15 mutations rewire RNA translation in chronic lymphocytic leukemia. Blood Advances, 2021, 5, 2788-2792. | 5.2 | 12 |
| 24 | TAp63 and BCL2 expression are co-affected by cell-extrinsic signals in chronic lymphocytic leukemia. Leukemia and Lymphoma, 2021, 62, 1-4. | 1.3 | 1 |
| 25 | The Significance of B-cell Receptor Stereotypy in Chronic Lymphocytic Leukemia. Hematology/Oncology Clinics of North America, 2021, 35, 687-702. | 2.2 | 3 |
| 26 | Comparison of different strategies for the triage to colposcopy of women tested high-risk HPV positive on self-collected cervicovaginal samples. Gynecologic Oncology, 2021, 162, 560-568. | 1.4 | 0 |
| 27 | MyPal ADULT study protocol: a randomised clinical trial of the MyPal ePRO-based early palliative care system in adult patients with haematological malignancies. BMJ Open, 2021, 11, e050256. | 1.9 | 8 |
| 28 | Distinctive Signaling Profiles With Distinct Biological and Clinical Implications in Aggressive CLL Subsets With Stereotyped B-Cell Receptor Immunoglobulin. Frontiers in Oncology, 2021, 11, 771454. | 2.8 | 8 |
| 29 | COVID-19 severity and mortality in patients with CLL: an update of the international ERIC and Campus CLL study. Leukemia, 2021, 35, 3444-3454. | 7.2 | 57 |
| 30 | Validation of the EuroClonality-NGS DNA capture panel as an integrated genomic tool for lymphoproliferative disorders. Blood Advances, 2021, 5, 3188-3198. | 5.2 | 2 |
| 31 | Understanding Monoclonal B Cell Lymphocytosis: An Interplay of Genetic and Microenvironmental Factors. Frontiers in Oncology, 2021, 11, 769612. | 2.8 | 10 |
| 32 | Specific T Cell Receptor Gene Repertoire Profiles in Subgroups of CLL Patients with Distinct Genomic Aberrations. Blood, 2021, 138, 3749-3749. | 1.4 | 0 |
| 33 | Different Prognostic Impact of Recurrent Gene Mutations in IGHV-Mutated and IGHV-Unmutated Chronic Lymphocytic Leukemia: A Retrospective, Multi-Center Cohort Study By Eric, the European Research Initiative on CLL, in Harmony. Blood, 2021, 138, 2617-2617. | 1.4 | 1 |
| 34 | The Clonotypic BCR IG of CLL Patients Contain Predicted T-Cell Class I Epitopes with Shared Structural Properties. Blood, 2021, 138, 1540-1540. | 1.4 | 2 |
| 35 | Distinct Modes of Ongoing Antigen Interactions Shape Intraclonal Dynamics in Splenic Marginal Zone Lymphoma. Blood, 2021, 138, 1330-1330. | 1.4 | 0 |
| 36 | Acceptability of Self-Sampling for Human Papillomavirus-Based Cervical Cancer Screening. Journal of Women's Health, 2020, 29, 1447-1456. | 3.3 | 10 |

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| 37 | Congenital and Acquired Chronic Neutropenias: Challenges, Perspectives and Implementation of the EuNetâ€INNOCHRON Action. HemaSphere, 2020, 4, e406. | 2.7 | 2 |
| 38 | Chronic lymphocytic leukemias with trisomy 12 show a distinct DNA methylation profile linked to altered chromatin activation. Haematologica, 2020, 105, 2864-2867. | 3.5 | 11 |
| 39 | Immunoglobulin gene analysis in chronic lymphocytic leukemia in the era of next generation sequencing. Leukemia, 2020, 34, 2545-2551. | 7.2 | 29 |
| 40 | Tracing CLL-biased stereotyped immunoglobulin gene rearrangements in normal B cell subsets using a high-throughput immunogenetic approach. Molecular Medicine, 2020, 26, 25. | 4.4 | 17 |
| 41 | COVID-19 severity and mortality in patients with chronic lymphocytic leukemia: a joint study by ERIC, the European Research Initiative on CLL, and CLL Campus. Leukemia, 2020, 34, 2354-2363. | 7.2 | 198 |
| 42 | T-Cell Dynamics in Chronic Lymphocytic Leukemia under Different Treatment Modalities. Clinical Cancer Research, 2020, 26, 4958-4969. | 7.0 | 18 |
| 43 | Prognostic impact of prevalent chronic lymphocytic leukemia stereotyped subsets: analysis within prospective clinical trials of the German CLL Study Group (GCLLSG). Haematologica, 2020, 105, 2598-2607. | 3.5 | 44 |
| 44 | B Cell Receptor Immunogenetics in B Cell Lymphomas: Immunoglobulin Genes as Key to Ontogeny and Clinical Decision Making. Frontiers in Oncology, 2020, 10, 67. | 2.8 | 26 |
| 45 |  IGLV3-21 * 01  is an inherited risk factor for CLL through the acquisition of a single-point mutation enabling autonomous BCR signaling. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 4320-4327. | 7.1 | 55 |
| 46 | High-throughput analysis of the T cell receptor gene repertoire in low-count monoclonal B cell lymphocytosis reveals a distinct profile from chronic lymphocytic leukemia. Haematologica, 2020, 105, e515. | 3.5 | 3 |
| 47 | Increased frequency of the single nucleotide polymorphism of the <scp>DARC</scp>/<scp>ACKR1</scp> gene associated with ethnic neutropenia in a cohort of European patients with chronic idiopathic neutropenia. American Journal of Hematology, 2020, 95, E163-E166. | 4.1 | 8 |
| 48 | Pretransplant Genetic Susceptibility: Clinical Relevance in Transplant-Associated Thrombotic Microangiopathy. Thrombosis and Haemostasis, 2020, 120, 638-646. | 3.4 | 33 |
| 49 | T Cells in Chronic Lymphocytic Leukemia: A Two-Edged Sword. Frontiers in Immunology, 2020, 11, 612244. | 4.8 | 31 |
| 50 | Immunoglobulin heavy variable somatic hyper mutation status in chronic lymphocytic leukaemia: on the threshold of a new era?. British Journal of Haematology, 2020, 189, 809-810. | 2.5 | 6 |
| 51 | Primary vitreoretinal lymphomas display a remarkably restricted immunoglobulin gene repertoire. Blood Advances, 2020, 4, 1357-1366. | 5.2 | 29 |
| 52 | TRIP - T cell receptor/immunoglobulin profiler. BMC Bioinformatics, 2020, 21, 422. | 2.6 | 11 |
| 53 | Genomic arrays identify high-risk chronic lymphocytic leukemia with genomic complexity: a multi-center study. Haematologica, 2020, 106, 87-97. | 3.5 | 43 |
| 54 | Challenges and Solutions for Collecting and Analyzing Real World Data: The Eric CLL Database as an Illustrative Example. HemaSphere, 2020, 4, e425. | 2.7 | 2 |

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| 55 | Development of a ePRO-Based Palliative Care Intervention for Cancer Patients: A Participatory Design Approach. Studies in Health Technology and Informatics, 2020, 270, 941-945. | 0.3 | 3 |
| 56 | Worldwide Examination of Patients with CLL Hospitalized for COVID-19. Blood, 2020, 136, 45-49. | 1.4 | 2 |
| 57 | T Cell Immunoprofiling of Patients with Relapsed and/or Refractory Myeloma Who Receive Daratumumab Monotherapy: Longitudinal Analysis during 7 Cycle Follow-up of the Rebuild Phase 2 Study. Blood, 2020, 136, 28-28. | 1.4 | 1 |
| 58 | Implementation of HPV-based Cervical Cancer Screening Combined with Self-sampling Using a Midwifery Network Across Rural Greece: The GRECOSELF Study. Cancer Prevention Research, 2019, 12, 701-710. | 1.5 | 17 |
| 59 | Standardized next-generation sequencing of immunoglobulin and T-cell receptor gene recombinations for MRD marker identification in acute lymphoblastic leukaemia; a EuroClonality-NGS validation study. Leukemia, 2019, 33, 2241-2253. | 7.2 | 177 |
| 60 | Approaching Empowerment Holistically: are Physicians Willing And Able?. International Journal of Reliable and Quality E-Healthcare, 2019, 8, 11-22. | 1.1 | 0 |
| 61 | Different time-dependent changes of risk for evolution in chronic lymphocytic leukemia with mutated or unmutated antigen B cell receptors. Leukemia, 2019, 33, 1801-1805. | 7.2 | 5 |
| 62 | Quality control and quantification in IG/TR next-generation sequencing marker identification: protocols and bioinformatic functionalities by EuroClonality-NGS. Leukemia, 2019, 33, 2254-2265. | 7.2 | 70 |
| 63 | EZH2 upregulates the PI3K/AKT pathway through IGF1R and MYC in clinically aggressive chronic lymphocytic leukaemia. Epigenetics, 2019, 14, 1125-1140. | 2.7 | 24 |
| 64 | Next-generation sequencing of immunoglobulin gene rearrangements for clonality assessment: a technical feasibility study by EuroClonality-NGS. Leukemia, 2019, 33, 2227-2240. | 7.2 | 92 |
| 65 | Study of gene expressions' correlation structures in subgroups of Chronic Lymphocytic Leukemia Patients. Journal of Biomedical Informatics, 2019, 95, 103211. | 4.3 | 2 |
| 66 | Stereotyped B Cell Receptor Immunoglobulins in B Cell Lymphomas. Methods in Molecular Biology, 2019, 1956, 139-155. | 0.9 | 17 |
| 67 | Inhibition of EZH2 and immune signaling exerts synergistic antitumor effects in chronic lymphocytic leukemia. Blood Advances, 2019, 3, 1891-1896. | 5.2 | 10 |
| 68 | DNA methylation profiles in chronic lymphocytic leukemia patients treated with chemoimmunotherapy. Clinical Epigenetics, 2019, 11, 177. | 4.1 | 15 |
| 69 | Cytogenetic complexity in chronic lymphocytic leukemia: definitions, associations, and clinical impact. Blood, 2019, 133, 1205-1216. | 1.4 | 164 |
| 70 | Integrated epigenomic and transcriptomic analysis reveals TP63 as a novel player in clinically aggressive chronic lymphocytic leukemia. International Journal of Cancer, 2019, 144, 2695-2706. | 5.1 | 24 |
| 71 | Dichotomous Toll-like receptor responses in chronic lymphocytic leukemia patients under ibrutinib treatment. Leukemia, 2019, 33, 1030-1051. | 7.2 | 4 |
| 72 | Skewing of the T-cell receptor repertoire in patients receiving rituximab after allogeneic hematopoietic cell transplantation: what lies beneath?. Leukemia and Lymphoma, 2019, 60, 1685-1692. | 1.3 | 5 |

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| 73 | Immunoglobulin Gene Analysis in Chronic Lymphocytic Leukemia. Methods in Molecular Biology, 2019, 1881, 51-62. | 0.9 | 2 |
| 74 | Diseaseâ€biased and shared characteristics of the immunoglobulin gene repertoires in marginal zone B cell lymphoproliferations. Journal of Pathology, 2019, 247, 416-421. | 4.5 | 25 |
| 75 | Tailored approaches grounded on immunogenetic features for refined prognostication in chronic lymphocytic leukemia. Haematologica, 2019, 104, 360-369. | 3.5 | 42 |
| 76 | Euroclonality-NGS DNA Capture Panel for Integrated Analysis of IG/TR Rearrangements, Translocations, Copy Number and Sequence Variation in Lymphoproliferative Disorders. Blood, 2019, 134, 888-888. | 1.4 | 4 |
| 77 | VH CDR3-Focused Somatic Hypermutation in CLL IGHV-IGHD-IGHJ Gene Rearrangements with 100% IGHV Germline Identity. Blood, 2019, 134, 4277-4277. | 1.4 | 3 |
| 78 | Detailed Functional Characterization of Splenic Marginal Zone Lymphoma: Uncovering Links between the Epigenetic and the Signaling Machinery. Blood, 2019, 134, 1512-1512. | 1.4 | 0 |
| 79 | Genome-Wide Histone Acetylation Profiling in Chronic Lymphocytic Leukemia Reveals a Distinctive Signature in Stereotyped Subset #8. Blood, 2019, 134, 1241-1241. | 1.4 | 0 |
| 80 | Higher Order Restrictions of the Immunoglobulin Repertoire in CLL: The Illustrative Case of Stereotyped Subsets #2 and #169. Blood, 2019, 134, 5453-5453. | 1.4 | 1 |
| 81 | Functional Calcitriol/Vitamin D Receptor Signaling in Chronic Lymphocytic Leukemia. Blood, 2019, 134, 3019-3019. | 1.4 | 0 |
| 82 | Longitudinal T Cell Immunoprofiling of Patients with Relapsed and/or Refractory Myeloma Who Receive Daratumumab Monotherapy: A Subanalysis of a Phase 2 Study (the REBUILD Study). Blood, 2019, 134, 3167-3167. | 1.4 | 2 |
| 83 | Changes in N-Glycosylation Induced By Somatic Hypermutation Modulate the Antigen Reactivity of the Immunoglobulin Receptors in CLL Stereotyped Subset #201. Blood, 2019, 134, 1733-1733. | 1.4 | 1 |
| 84 | Highly similar genomic landscapes in monoclonal B-cell lymphocytosis and ultra-stable chronic lymphocytic leukemia with low frequency of driver mutations. Haematologica, 2018, 103, 865-873. | 3.5 | 47 |
| 85 | Restricted T cell receptor repertoire in CLL-like monoclonal B cell lymphocytosis and early stage CLL. OncoImmunology, 2018, 7, e1432328. | 4.6 | 20 |
| 86 | No improvement in long-term survival over time for chronic lymphocytic leukemia patients in stereotyped subsets #1 and #2 treated with chemo(immuno)therapy. Haematologica, 2018, 103, e158-e161. | 3.5 | 16 |
| 87 | Automated shape-based clustering of 3D immunoglobulin protein structures in chronic lymphocytic leukemia. BMC Bioinformatics, 2018, 19, 414. | 2.6 | 9 |
| 88 | Immunoglobulin Gene Sequence Analysis In Chronic Lymphocytic Leukemia: From Patient Material To Sequence Interpretation. Journal of Visualized Experiments, 2018, , . | 0.3 | 6 |
| 89 | IRProfiler â€“ a software toolbox for high throughput immune receptor profiling. BMC Bioinformatics, 2018, 19, 144. | 2.6 | 7 |
| 90 | Eliciting Anti-Tumor T Cell Immunity in Chronic Lymphocytic Leukemia (CLL) with PD-L1/PD-1 Blockade Is Enhanced By Avadomide Immunotherapy through the Triggering of Immunogenic Interferon Signaling. Blood, 2018, 132, 237-237. | 1.4 | 2 |

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| 91 | RPS15 mutations Repress mRNA Translation in Chronic Lymphocytic Leukemia Cells. Blood, 2018, 132, 1843-1843. | 1.4 | 1 |
| 92 | Remarkable Functional Constraints on the Antigen Receptors of CLL Stereotyped Subset #2: High-Throughput Immunogenetic Evidence. Blood, 2018, 132, 1839-1839. | 1.4 | 5 |
| 93 | A novel ex vivo high-throughput assay reveals antiproliferative effects of idelalisib and ibrutinib in chronic lymphocytic leukemia. Oncotarget, 2018, 9, 26019-26031. | 1.8 | 8 |
| 94 | Evidence for Epitope-Specific T Cell Responses in HIV-Associated Non Neoplastic Lymphadenopathy: High-Throughput Immunogenetic Evidence. Blood, 2018, 132, 1117-1117. | 1.4 | 2 |
| 95 | IGHV Gene Replacement: A Potential Mechanism for Establishing Stereotypy in Certain Cases of Chronic Lymphocytic Leukemia. Blood, 2018, 132, 1841-1841. | 1.4 | 0 |
| 96 | The Transcription Factor TAp63 Exerts Pro-Survival Effects in Chronic Lymphocytic Leukemia Acting through the BCL2 Pathway. Blood, 2018, 132, 3110-3110. | 1.4 | 0 |
| 97 | Pre-Transplant Genetic Susceptibility in Adult Allogeneic Hematopoietic Cell Transplant Recipients: Incidence and Clinical Relevance in Transplant-Associated Thrombotic Microangiopathy. Blood, 2018, 132, 3401-3401. | 1.4 | 0 |
| 98 | Longitudinal High-Throughput T Cell Repertoire Profiling of Chronic Lymphocytic Leukemia Patients Under Different Types of Treatment: Implications for Combination Strategies. Blood, 2018, 132, 4400-4400. | 1.4 | 0 |
| 99 | Splenic diffuse red pulp small B-cell lymphoma displays increased expression of cyclin D3 and recurrent CCND3 mutations. Blood, 2017, 129, 1042-1045. | 1.4 | 52 |
| 100 | Comprehensive translocation and clonality detection in lymphoproliferative disorders by next-generation sequencing. Haematologica, 2017, 102, e57-e60. | 3.5 | 35 |
| 101 | 3D Protein-Structure-Oriented Discovery of Clinical Relation Across Chronic Lymphocytic Leukemia Patients. Lecture Notes in Computer Science, 2017, , 139-150. | 1.3 | 0 |
| 102 | High-Throughput Immunogenetics for Clinical and Research Applications in Immunohematology: Potential and Challenges. Journal of Immunology, 2017, 198, 3765-3774. | 0.8 | 61 |
| 103 | Numerous Ontogenetic Roads to Mantle Cell Lymphoma. American Journal of Pathology, 2017, 187, 1454-1458. | 3.8 | 11 |
| 104 | Distinct homotypic B-cell receptor interactions shape the outcome of chronic lymphocytic leukaemia. Nature Communications, 2017, 8, 15746. | 12.8 | 93 |
| 105 | Cytotoxic T cells in chronic idiopathic neutropenia express restricted antigen receptors. Leukemia and Lymphoma, 2017, 58, 2926-2933. | 1.3 | 6 |
| 106 | Chronic Lymphocytic Leukemia with Mutated IGHV4-34 Receptors: Shared and Distinct Immunogenetic Features and Clinical Outcomes. Clinical Cancer Research, 2017, 23, 5292-5301. | 7.0 | 27 |
| 107 | Immunoglobulin genes in chronic lymphocytic leukemia: key to understanding the disease and improving risk stratification. Haematologica, 2017, 102, 968-971. | 3.5 | 28 |
| 108 | The inhibitory receptor toll interleukin-1R 8 (TIR8/IL-1R8/SIGIRR) is downregulated in chronic lymphocytic leukemia. Leukemia and Lymphoma, 2017, 58, 2419-2425. | 1.3 | 9 |

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| 109 | Clonal B-cell lymphocytosis of marginal zone origin. Best Practice and Research in Clinical Haematology, 2017, 30, 77-83. | 1.7 | 12 |
| 110 | Chronic Lymphocytic Leukemia Patient Clustering Based on Somatic Hypermutation (SHM) Analysis. Advances in Experimental Medicine and Biology, 2017, 988, 127-138. | 1.6 | 4 |
| 111 | A gene is known by the company it keeps: enrichment ofTNFAIP3gene aberrations in MALT lymphomas expressing IGHV4-34 antigen receptors. Journal of Pathology, 2017, 243, 403-406. | 4.5 | 7 |
| 112 | Calreticulin as a novel B-cell receptor antigen in chronic lymphocytic leukemia. Haematologica, 2017, 102, e394-e396. | 3.5 | 10 |
| 113 | Monoclonal B-cell lymphocytosis in a hospital-based UK population and a rural Ugandan population: a cross-sectional study. Lancet Haematology,the, 2017, 4, e334-e340. | 4.6 | 12 |
| 114 | Tp53gene p72R polymorphism in chronic lymphocytic leukemia: incidence and clinical significance amongst cases with unmutated immunoglobulin receptors. Leukemia and Lymphoma, 2017, 58, 726-728. | 1.3 | 2 |
| 115 | Binding of CLL Subset 4 B Cell Receptor Immunoglobulins to Viable Human Memory B Lymphocytes Requires a Distinctive IGKV Somatic Mutation. Molecular Medicine, 2017, 23, 1-12. | 4.4 | 14 |
| 116 | T cells in chronic lymphocytic leukemia: can they fight?. Oncotarget, 2017, 8, 99209-99210. | 1.8 | 4 |
| 117 | Karyotypic complexity rather than chromosome 8 abnormalities aggravates the outcome of chronic lymphocytic leukemia patients with TP53 aberrations. Oncotarget, 2016, 7, 80916-80924. | 1.8 | 29 |
| 118 | Additional trisomies amongst patients with chronic lymphocytic leukemia carrying trisomy 12: the accompanying chromosome makes a difference. Haematologica, 2016, 101, e299-e302. | 3.5 | 35 |
| 119 | Frequent NFKBIE deletions are associated with poor outcome in primary mediastinal B-cell lymphoma. Blood, 2016, 128, 2666-2670. | 1.4 | 82 |
| 120 | ATM mutations in major stereotyped subsets of chronic lymphocytic leukemia: enrichment in subset #2 is associated with markedly short telomeres. Haematologica, 2016, 101, e369-e373. | 3.5 | 16 |
| 121 | Different spectra of recurrent gene mutations in subsets of chronic lymphocytic leukemia harboring stereotyped B-cell receptors. Haematologica, 2016, 101, 959-967. | 3.5 | 57 |
| 122 | Innovation in the prognostication of chronic lymphocytic leukemia: how far beyond TP53 gene analysis can we go?. Haematologica, 2016, 101, 263-265. | 3.5 | 19 |
| 123 | B Cell Anergy Modulated by TLR1/2 and the miR-17âˆ¼92 Cluster Underlies the Indolent Clinical Course of Chronic Lymphocytic Leukemia Stereotyped Subset #4. Journal of Immunology, 2016, 196, 4410-4417. | 0.8 | 30 |
| 124 | Whole-exome sequencing in relapsing chronic lymphocytic leukemia: clinical impact of recurrent RPS15 mutations. Blood, 2016, 127, 1007-1016. | 1.4 | 130 |
| 125 | Clinical impact of recurrently mutated genes on lymphoma diagnostics: state-of-the-art and beyond. Haematologica, 2016, 101, 1002-1009. | 3.5 | 43 |
| 126 | NF-ÎºB activation in chronic lymphocytic leukemia: A point of convergence of external triggers and intrinsic lesions. Seminars in Cancer Biology, 2016, 39, 40-48. | 9.6 | 60 |

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| 127 | B-cell malignancies: All roads lead to NF-Î²B activation. Seminars in Cancer Biology, 2016, 39, 1-2. | 9.6 | 4 |
| 128 | Toll-like receptors signaling: A complex network for NF-Î²B activation in B-cell lymphoid malignancies. Seminars in Cancer Biology, 2016, 39, 15-25. | 9.6 | 65 |
| 129 | Integrating multiple immunogenetic data sources for feature extraction and mining somatic hypermutation patterns: the case of â€œtowards analysisâ€•in chronic lymphocytic leukaemia. BMC Bioinformatics, 2016, 17, 173. | 2.6 | 1 |
| 130 | An Immunogenetic Signature of Ongoing Antigen Interactions in Splenic Marginal Zone Lymphoma Expressing IGHV1-2*04 Receptors. Clinical Cancer Research, 2016, 22, 2032-2040. | 7.0 | 17 |
| 131 | Antigen Selection Shapes the T-cell Repertoire in Chronic Lymphocytic Leukemia. Clinical Cancer Research, 2016, 22, 167-174. | 7.0 | 43 |
| 132 | Longitudinal Assessment of CLL Patients Under Ibrutinib Treatment Reveals Maintained Capacity to Respond to Microenvironmental Stimuli through the Toll-like Receptors. Blood, 2016, 128, 2025-2025. | 1.4 | 1 |
| 133 | CLL Stereotyped IGHV-D-J Rearrangements Can Be Detected Throughout Normal B-Cell Developmental Stages in Aged People When Using Ultra-Deep, Next Generation Sequencing Techniques. Blood, 2016, 128, 2028-2028. | 1.4 | 3 |
| 134 | Distinct Immunogenetic Signatures in IgA Versus IgG Multiple Myeloma. Blood, 2016, 128, 2062-2062. | 1.4 | 4 |
| 135 | In CLL, Myeloid-Derived Suppressor Cells and Their Monocytic and Granulocytic Varieties Differ in T-Cell Subset Association and Polarization Induction. Blood, 2016, 128, 4350-4350. | 1.4 | 1 |
| 136 | Automated Clustering Analysis of Immunoglobulin Sequences in Chronic Lymphocytic Leukemia Based on 3D Structural Descriptors. Blood, 2016, 128, 4365-4365. | 1.4 | 2 |
| 137 | Reappraising Immunoglobulin Repertoire Restrictions in Chronic Lymphocytic Leukemia: Focus on Major Stereotyped Subsets and Closely Related Satellites. Blood, 2016, 128, 4376-4376. | 1.4 | 1 |
| 138 | The histone methyltransferase EZH2 as a novel prosurvival factor in clinically aggressive chronic lymphocytic leukemia. Oncotarget, 2016, 7, 35946-35959. | 1.8 | 29 |
| 139 | Molecular Immunoprofiling the T Cell Repertoire after Rituximab Administration Reveals Frequent Oligoclonality Albeit with Different Patterns Depending on the Clinical Context. Blood, 2016, 128, 5792-5792. | 1.4 | 0 |
| 140 | ÎFÎšÎ'Î™Î• Deletions: A Novel Marker of Clinical Aggressiveness in Primary Mediastinal B-Cell Lymphoma. Blood, 2016, 128, 609-609. | 1.4 | 0 |
| 141 | Not all IGHV3-21 chronic lymphocytic leukemias are equal: prognostic considerations. Blood, 2015, 125, 856-859. | 1.4 | 70 |
| 142 | Excessive antigen reactivity may underlie the clinical aggressiveness of chronic lymphocytic leukemia stereotyped subset #8. Blood, 2015, 125, 3580-3587. | 1.4 | 49 |
| 143 | Prognostic relevance of MYD88 mutations in CLL: the jury is still out. Blood, 2015, 126, 1043-1044. | 1.4 | 32 |
| 144 | Functional loss of Iº Bÿ leads to NF-Î²B deregulation in aggressive chronic lymphocytic leukemia. Journal of Experimental Medicine, 2015, 212, 833-843. | 8.5 | 85 |

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| 145 | AEGLE: A big bio-data analytics framework for integrated health-care services. , 2015, , . | | 4 |
| 146 | Toll-like receptor stimulation in splenic marginal zone lymphoma can modulate cell signaling, activation and proliferation. Haematologica, 2015, 100, 1460-1468. | 3.5 | 19 |
| 147 | Ofatumumab in poor-prognosis chronic lymphocytic leukemia: a Phase IV, non-interventional, observational study from the European Research Initiative on Chronic Lymphocytic Leukemia. Haematologica, 2015, 100, 511-516. | 3.5 | 42 |
| 148 | Molecular Evidence for Antigen Drive in the Natural History of Mantle Cell Lymphoma. American Journal of Pathology, 2015, 185, 1740-1748. | 3.8 | 13 |
| 149 | Non-coding recurrent mutations in chronic lymphocytic leukaemia. Nature, 2015, 526, 519-524. | 27.8 | 749 |
| 150 | Genetics and Prognostication in Splenic Marginal Zone Lymphoma: Revelations from Deep Sequencing. Clinical Cancer Research, 2015, 21, 4174-4183. | 7.0 | 129 |
| 151 | ARResT/AssignSubsets: a novel application for robust subclassification of chronic lymphocytic leukemia based on B cell receptor IG stereotypy. Bioinformatics, 2015, 31, 3844-3846. | 4.1 | 43 |
| 152 | Targeted next-generation sequencing in chronic lymphocytic leukemia: a high-throughput yet tailored approach will facilitate implementation in a clinical setting. Haematologica, 2015, 100, 370-376. | 3.5 | 57 |
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