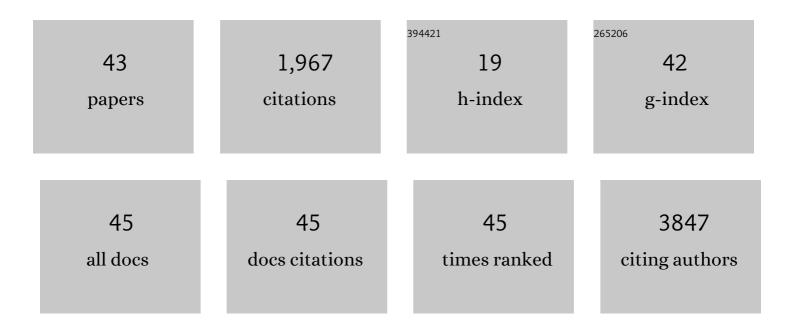
## Bianca Tesi

List of Publications by Year in descending order

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RIANCA TESL

#	Article	IF	CITATIONS
1	Rubella vaccine–induced granulomas are a novel phenotype with incomplete penetrance of genetic defects in cytotoxicity. Journal of Allergy and Clinical Immunology, 2022, 149, 388-399.e4.	2.9	11
2	Detection of germline mosaicism in fathers of children with intellectual disability syndromes caused by de novo variants. Molecular Genetics & Genomic Medicine, 2022, 10, e1880.	1.2	10
3	Childhood Kaposi sarcoma related to hypomorphic severe combined immunodeficiency caused by a novel <i>CORO1A</i> mutation. Pediatric Blood and Cancer, 2022, 69, e29487.	1.5	0
4	Patients with both Langerhans cell histiocytosis and Crohn's disease highlight a common role of interleukinâ€23. Acta Paediatrica, International Journal of Paediatrics, 2021, 110, 1315-1321.	1.5	8
5	Integration of whole genome sequencing into a healthcare setting: high diagnostic rates across multiple clinical entities in 3219 rare disease patients. Genome Medicine, 2021, 13, 40.	8.2	116
6	Clinical and laboratory signs of haemophagocytic lymphohistiocytosis associated with pandemic influenza A (H1N1) infection in patients needing extracorporeal membrane oxygenation. European Journal of Anaesthesiology, 2021, 38, 692-701.	1.7	1
7	Molecular Genetics Diversity of Primary Hemophagocytic Lymphohistiocytosis among Polish Pediatric Patients. Archivum Immunologiae Et Therapiae Experimentalis, 2021, 69, 31.	2.3	4
8	Adult-Onset Ataxia With Neuropathy and White Matter Abnormalities Due to a Novel SAMD9L Variant. Neurology: Genetics, 2021, 7, e628.	1.9	1
9	Whole Transcriptome Analysis Identifies Distinct Gene Expression Profiles between SF3B1mut and SF3B1 wt Myelodysplastic Syndrome with Ring Sideroblasts. Blood, 2021, 138, 3695-3695.	1.4	0
10	Multinational Study on the Clinical and Genetic Features of the ERCC6L2-Disease. Blood, 2021, 138, 864-864.	1.4	2
11	Alternative UNC13D Promoter Encodes a Functional Munc13-4 Isoform Predominantly Expressed in Lymphocytes and Platelets. Frontiers in Immunology, 2020, 11, 1154.	4.8	2
12	Diagnostic challenges for a novel SH2D1A mutation associated with Xâ€linked lymphoproliferative disease. Pediatric Blood and Cancer, 2020, 67, e28184.	1.5	4
13	Heterozygous variants in <i>DCC</i> . Neurology: Genetics, 2020, 6, e526.	1.9	4
14	Microdeletion of 7p12.1p13, including <i><scp>IKZF</scp>1</i> , causes intellectual impairment, overgrowth, and susceptibility to leukaemia. British Journal of Haematology, 2019, 185, 354-357.	2.5	4
15	Human DEF6 deficiency underlies an immunodeficiency syndrome with systemic autoimmunity and aberrant CTLA-4 homeostasis. Nature Communications, 2019, 10, 3106.	12.8	48
16	Early activating somatic <i>PIK3CA</i> mutations promote ectopic muscle development and upper limb overgrowth. Clinical Genetics, 2019, 96, 118-125.	2.0	14
17	Haploinsufficiency of <i>UNC13D</i> increases the risk of lymphoma. Cancer, 2019, 125, 1848-1854.	4.1	8
18	Nordic Guidelines for Germline Predisposition to Myeloid Neoplasms in Adults: Recommendations for Genetic Diagnosis, Clinical Management and Follow-up. HemaSphere, 2019, 3, e321.	2.7	51

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19	Fatal Central Nervous System Lymphocytic Vasculitis after Treatment for Burkitt Lymphoma in a Patient with a SH2D1A Mutation. Pediatric Infectious Disease Journal, 2019, 38, e29-e31.	2.0	4
20	Elevated ferritin and soluble CD25 in critically ill patients are associated with parameters of (hyper) inflammation and lymphocyte cytotoxicity. Minerva Anestesiologica, 2019, 85, 1289-1298.	1.0	13
21	A RAB27A 5′ untranslated region structural variant associated with late-onset hemophagocytic lymphohistiocytosis and normal pigmentation. Journal of Allergy and Clinical Immunology, 2018, 142, 317-321.e8.	2.9	22
22	HLH: genomics illuminates pathophysiological diversity. Blood, 2018, 132, 5-7.	1.4	11
23	ARID5B regulates metabolic programming in human adaptive NK cells. Journal of Experimental Medicine, 2018, 215, 2379-2395.	8.5	98
24	Targeting SAMHD1 with the Vpx protein to improve cytarabine therapy for hematological malignancies. Nature Medicine, 2017, 23, 256-263.	30.7	102
25	Gain-of-function SAMD9L mutations cause a syndrome of cytopenia, immunodeficiency, MDS, and neurological symptoms. Blood, 2017, 129, 2266-2279.	1.4	152
26	A Case of Familial Hemophagocytic Lymphohistiocytosis Type 4 With Involvement of the Central Nervous System Complicated With Infarct. Journal of Pediatric Hematology/Oncology, 2017, 39, e321-e324.	0.6	4
27	Differences in Granule Morphology yet Equally Impaired Exocytosis among Cytotoxic T Cells and NK Cells from Chediak–Higashi Syndrome Patients. Frontiers in Immunology, 2017, 8, 426.	4.8	26
28	Unperturbed Cytotoxic Lymphocyte Phenotype and Function in Myalgic Encephalomyelitis/Chronic Fatigue Syndrome Patients. Frontiers in Immunology, 2017, 8, 723.	4.8	24
29	Epigenetic Regulation of Adaptive NK Cell Diversification. Trends in Immunology, 2016, 37, 451-461.	6.8	60
30	Successful Hematopoietic Stem Cell Transplantation in a Patient with LPS-Responsive Beige-Like Anchor (LRBA) Gene Mutation. Journal of Clinical Immunology, 2016, 36, 480-489.	3.8	30
31	Cytopenia, Predisposition to Myelodysplastic Syndrome, Immunodeficiency, and Neurological Disease Caused By Gain-of-Function SAMD9L Mutations Is Frequently Ameliorated By Hematopoietic Revertant Mosaicism. Blood, 2016, 128, 4299-4299.	1.4	1
32	A Hemophagocytic Lymphohistiocytosis Case with Newly Defined UNC13D (c.175G>C; p.Ala59Pro) Mutation and a Rare Complication. Turkish Journal of Haematology, 2016, 32, 355-358.	0.5	4
33	Targeted high-throughput sequencing for genetic diagnostics of hemophagocytic lymphohistiocytosis. Genome Medicine, 2015, 7, 130.	8.2	37
34	The syndrome of hemophagocytic lymphohistiocytosis in primary immunodeficiencies: implications for differential diagnosis and pathogenesis. Haematologica, 2015, 100, 978-988.	3.5	161
35	Spectrum of Atypical Clinical Presentations in Patients with Biallelic <i>PRF1</i> Missense Mutations. Pediatric Blood and Cancer, 2015, 62, 2094-2100.	1.5	38
36	Hemophagocytic lymphohistiocytosis in 2 patients with underlying IFN-γ receptor deficiency. Journal of Allergy and Clinical Immunology, 2015, 135, 1638-1641.e5.	2.9	69

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37	Cytomegalovirus Infection Drives Adaptive Epigenetic Diversification of NK Cells with Altered Signaling and Effector Function. Immunity, 2015, 42, 443-456.	14.3	650
38	Diversification and Functional Specialization of Human NK Cell Subsets. Current Topics in Microbiology and Immunology, 2015, 395, 63-93.	1.1	56
39	An N-Terminal Missense Mutation in STX11 Causative of FHL4 Abrogates Syntaxin-11 Binding to Munc18-2. Frontiers in Immunology, 2014, 4, 515.	4.8	20
40	Pathophysiology and spectrum of diseases caused by defects in lymphocyte cytotoxicity. Experimental Cell Research, 2014, 325, 10-17.	2.6	38
41	Immunomodulatory activity of commonly used drugs on Fc-receptor-mediated human natural killer cell activation. Cancer Immunology, Immunotherapy, 2014, 63, 627-641.	4.2	33
42	Novel STAT3 Mutation Causing Hyper-IgE Syndrome: Studies of the Clinical Course and Immunopathology. Journal of Clinical Immunology, 2014, 34, 469-477.	3.8	11
43	An Indian boy with griscelli syndrome type 2: Case report and review of literature. Indian Journal of Dermatology, 2014, 59, 394.	0.3	9