Paul A Lacaze

List of Publications by Year in Descending Order

67 papers	1,298 citations	18 h-index	35 g-index
94 ext. papers	1,782 ext. citations	5.8 avg, IF	4.14 L-index

#	Paper	IF	Citations
67	Population DNA screening for medically actionable disease risk in adults.. *Medical Journal of Australia*, **2022**,	4	2
66	Co-designing genomics research with a large group of donor-conceived siblings.. *Research Involvement and Engagement*, **2021**, 7, 89	4.4	0
65	Population genomic screening of young adults for familial hypercholesterolaemia: a cost-effectiveness analysis. *European Heart Journal*, **2021**,	9.5	3
64	Genetic testing in dementia-A medical genetics perspective. *International Journal of Geriatric Psychiatry*, **2021**, 36, 1158-1170	3.9	2
63	Rare Germline Pathogenic Variants Identified by Multigene Panel Testing and the Risk of Aggressive Prostate Cancer. *Cancers*, **2021**, 13,	6.6	1
62	Genomic Risk Score for Melanoma in a Prospective Study of Older Individuals. *Journal of the National Cancer Institute*, **2021**, 113, 1379-1385	9.7	4
61	Essentials of a new clinical practice guidance on familial hypercholesterolaemia for physicians. *Internal Medicine Journal*, **2021**, 51, 769-779	1.6	0
60	Involving elderly research participants in the co-design of a future multi-generational cohort study. *Research Involvement and Engagement*, **2021**, 7, 23	4.4	1
59	Study protocol: the Australian genetics and life insurance moratorium-monitoring the effectiveness and response (A-GLIMMER) project. *BMC Medical Ethics*, **2021**, 22, 63	2.9	1
58	Effect of APOE and a polygenic risk score on incident dementia and cognitive decline in a healthy older population. *Aging Cell*, **2021**, 20, e13384	9.9	3
57	Synopsis of an integrated guidance for enhancing the care of familial hypercholesterolaemia: an Australian perspective. *American Journal of Preventive Cardiology*, **2021**, 6, 100151	1.9	0
56	Genetic variants associated with inherited cardiovascular disorders among 13,131 asymptomatic older adults of European descent. *Npj Genomic Medicine*, **2021**, 6, 51	6.2	3
55	Genomic Risk Prediction for Breast Cancer in Older Women. *Cancers*, **2021**, 13,	6.6	1
54	Ethical and practical implications of returning genetic research results: two Australian case studies. *Medical Journal of Australia*, **2021**, 214, 259-262.e1	4	
53	Integrated Guidance for Enhancing the Care of Familial Hypercholesterolaemia in Australia. *Heart Lung and Circulation*, **2021**, 30, 324-349	1.8	20
52	Polygenic score modifies risk for Alzheimer's disease in ε4 homozygotes at phenotypic extremes. *Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring*, **2021**, 13, e12226	5.2	0
51	Monitoring the genetic testing and life insurance moratorium in Australia: a national research project. *Medical Journal of Australia*, **2021**, 214, 157-159.e1	4	7

50	Involving people affected by a rare condition in shaping future genomic research. *Research Involvement and Engagement*, **2021**, 7, 14	4.4	1
49	Predictive Performance of a Polygenic Risk Score for Incident Ischemic Stroke in a Healthy Older Population. *Stroke*, **2021**, 52, 2882-2891	6.7	3
48	Prognostic Value of a Polygenic Risk Score for Coronary Heart Disease in Individuals Aged 70 Years and Older.. *Circulation Genomic and Precision Medicine*, **2021**, CIRCGEN121003429	5.2	0
47	Population-based estimates of breast cancer risk for carriers of pathogenic variants identified by gene-panel testing. *Npj Breast Cancer*, **2021**, 7, 153	7.8	1
46	A polygenic risk score predicts mosaic loss of chromosome Y in circulating blood cells.. *Cell and Bioscience*, **2021**, 11, 205	9.8	0
45	DNA methylation analysis of candidate genes associated with dementia in peripheral blood. *Epigenomics*, **2020**, 12, 2109-2123	4.4	3
44	Recovery of natural killer cell cytotoxicity in a p.A91V perforin homozygous patient following severe haemophagocytic lymphohistiocytosis. *British Journal of Haematology*, **2020**, 190, 458-461	4.5	
43	Familial Hypercholesterolemia in a Healthy Elderly Population. *Circulation Genomic and Precision Medicine*, **2020**, 13, e002938	5.2	2
42	Genetic Variation in PEAR1, Cardiovascular Outcomes and Effects of Aspirin in a Healthy Elderly Population. *Clinical Pharmacology and Therapeutics*, **2020**, 108, 1289-1298	6.1	5
41	Impact of direct-to-consumer genetic testing on Australian clinical genetics services. *European Journal of Medical Genetics*, **2020**, 63, 103968	2.6	6
40	Medically actionable pathogenic variants in a population of 13,131 healthy elderly individuals. *Genetics in Medicine*, **2020**, 22, 1883-1886	8.1	10
39	Genetic associations with clozapine-induced myocarditis in patients with schizophrenia. *Translational Psychiatry*, **2020**, 10, 37	8.6	10
38	The Medical Genome Reference Bank contains whole genome and phenotype data of 2570 healthy elderly. *Nature Communications*, **2020**, 11, 435	17.4	20
37	Blood DNA methylation signatures to detect dementia prior to overt clinical symptoms. *Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring*, **2020**, 12, e12056	5.2	2
36	Disclosing genetic information to family members without consent: Five Australian case studies. *European Journal of Medical Genetics*, **2020**, 63, 104035	2.6	2
35	Baseline characteristics and age-related macular degeneration in participants of the "ASPirin in Reducing Events in the Elderly" (ASPREE)-AMD trial. *Contemporary Clinical Trials Communications*, **2020**, 20, 100667	1.8	1
34	Genetic discrimination by Australian insurance companies: a survey of consumer experiences. *European Journal of Human Genetics*, **2020**, 28, 108-113	5.3	14
33	Prevalence and disease predisposition of p.A91V perforin in an aged population of European ancestry. *Blood*, **2020**, 135, 582-584	2.2	2

32	Genetic resilience to Alzheimer's disease in APOE ε4 homozygotes: A systematic review. *Alzheimer's and Dementia*, **2019**, 15, 1612-1623	1.2	12
31	Public Involvement in Global Genomics Research: A Scoping Review. *Frontiers in Public Health*, **2019**, 7, 79	6	21
30	Implementation of public health genomics in Pakistan. *European Journal of Human Genetics*, **2019**, 27, 1485-1492	5.3	5
29	Response to Veenstra et al. *Genetics in Medicine*, **2019**, 21, 2842-2843	8.1	4
28	Genetic testing and insurance in Australia. *Australian Journal of General Practice*, **2019**, 48, 96-99	1.5	9
27	Population genomic screening of all young adults in a health-care system: a cost-effectiveness analysis. *Genetics in Medicine*, **2019**, 21, 1958-1968	8.1	34
26	The Medical Genome Reference Bank: a whole-genome data resource of 4000 healthy elderly individuals. Rationale and cohort design. *European Journal of Human Genetics*, **2019**, 27, 308-316	5.3	17
25	Regulation of Internet-based Genetic Testing: Challenges for Australia and Other Jurisdictions. *Frontiers in Public Health*, **2018**, 6, 24	6	7
24	Phenotypic Heterogeneity in Dementia: A Challenge for Epidemiology and Biomarker Studies. *Frontiers in Public Health*, **2018**, 6, 181	6	27
23	Blood DNA methylation as a potential biomarker of dementia: A systematic review. *Alzheimer's and Dementia*, **2018**, 14, 81-103	1.2	44
22	Genetics, Insurance and Professional Practice: Survey of the Australasian Clinical Genetics Workforce. *Frontiers in Public Health*, **2018**, 6, 333	6	6
21	The genomic potential of the Aspirin in Reducing Events in the Elderly and Statins in Reducing Events in the Elderly studies. *Internal Medicine Journal*, **2017**, 47, 461-463	1.6	7
20	Pathogenic variants in the healthy elderly: unique ethical and practical challenges. *Journal of Medical Ethics*, **2017**, 43, 714-722	2.5	9
19	Penetrance and the Healthy Elderly. *Genetic Testing and Molecular Biomarkers*, **2017**, 21, 637-640	1.6	2
18	Rare disease registries: a call to action. *Internal Medicine Journal*, **2017**, 47, 1075-1079	1.6	22
17	The Dangers of Direct-to-Consumer Genetic Testing for Alzheimer's Disease : Comment on "Personal Genomic Testing, Genetic Inheritance, and Uncertainty". *Journal of Bioethical Inquiry*, **2017**, 14, 585-587	1.9	2
16	Genetics and Insurance in Australia: Concerns around a Self-Regulated Industry. *Public Health Genomics*, **2017**, 20, 247-256	1.9	11
15	Should Australia Ban the Use of Genetic Test Results in Life Insurance?. *Frontiers in Public Health*, **2017**, 5, 330	6	13

14	CD200 receptor restriction of myeloid cell responses antagonizes antiviral immunity and facilitates cytomegalovirus persistence within mucosal tissue. *PLoS Pathogens*, **2015**, 11, e1004641	7.6	12
13	ID4 controls mammary stem cells and marks breast cancers with a stem cell-like phenotype. *Nature Communications*, **2015**, 6, 6548	17.4	40
12	Whole blood gene expression profiling of neonates with confirmed bacterial sepsis. *Genomics Data*, **2015**, 3, 41-8		24
11	Identification of a human neonatal immune-metabolic network associated with bacterial infection. *Nature Communications*, **2014**, 5, 4649	17.4	84
10	The transcription factor STAT-1 couples macrophage synthesis of 25-hydroxycholesterol to the interferon antiviral response. *Immunity*, **2013**, 38, 106-18	32.3	258
9	Temporal profiling of the coding and noncoding murine cytomegalovirus transcriptomes. *Journal of Virology*, **2011**, 85, 6065-76	6.6	24
8	Host defense against viral infection involves interferon mediated down-regulation of sterol biosynthesis. *PLoS Biology*, **2011**, 9, e1000598	9.7	181
7	Reversible inhibition of murine cytomegalovirus replication by gamma interferon (IFN-γ) in primary macrophages involves a primed type I IFN-signaling subnetwork for full establishment of an immediate-early antiviral state. *Journal of Virology*, **2011**, 85, 10286-99	6.6	25
6	Consolidation of the cancer genome into domains of repressive chromatin by long-range epigenetic silencing (LRES) reduces transcriptional plasticity. *Nature Cell Biology*, **2010**, 12, 235-46	23.4	153
5	Construction of a large scale integrated map of macrophage pathogen recognition and effector systems. *BMC Systems Biology*, **2010**, 4, 63	3.5	32
4	Combined genome-wide expression profiling and targeted RNA interference in primary mouse macrophages reveals perturbation of transcriptional networks associated with interferon signalling. *BMC Genomics*, **2009**, 10, 372	4.5	18
3	A logic-based diagram of signalling pathways central to macrophage activation. *BMC Systems Biology*, **2008**, 2, 36	3.5	46
2	Quantitative assessment of human whole blood RNA as a potential biomarker for infectious disease. *Analyst, The*, **2007**, 132, 1200-9	5	11
1	The Medical Genome Reference Bank: Whole genomes and phenotype of 2,570 healthy elderly		1