

Simon Mallal

List of Publications by Year in descending order

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Version: 2024-02-01

129
papers

15,602
citations

38742

50
h-index

18647

119
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136
all docs

136
docs citations

136
times ranked

17364
citing authors

#	ARTICLE	IF	CITATIONS
1	SARS-CoV-2 vaccination induces immunological T cell memory able to cross-recognize variants from Alpha to Omicron. <i>Cell</i> , 2022, 185, 847-859.e11.	28.9	590
2	Immunodominant MHC-II (Major Histocompatibility Complex II) Restricted Epitopes in Human Apolipoprotein B. <i>Circulation Research</i> , 2022, 131, 258-276.	4.5	8
3	Altered subgenomic RNA abundance provides unique insight into SARS-CoV-2 B.1.1.7/Alpha variant infections. <i>Communications Biology</i> , 2022, 5, .	4.4	12
4	Deep Sequencing Analysis of Individual HIV-1 Proviruses Reveals Frequent Asymmetric Long Terminal Repeats. <i>Journal of Virology</i> , 2022, 96, .	3.4	8
5	The Proximal Airway Is a Reservoir for Adaptive Immunologic Memory in Idiopathic Subglottic Stenosis. <i>Laryngoscope</i> , 2021, 131, 610-617.	2.0	12
6	Human leukocyte antigen associations with protection against tuberculosis infection and disease in human immunodeficiency virus-1 infected individuals, despite household tuberculosis exposure and immune suppression. <i>Tuberculosis</i> , 2021, 126, 102023.	1.9	7
7	Comprehensive analysis of T cell immunodominance and immunoprevalence of SARS-CoV-2 epitopes in COVID-19 cases. <i>Cell Reports Medicine</i> , 2021, 2, 100204.	6.5	437
8	Subgenomic RNA identification in SARS-CoV-2 genomic sequencing data. <i>Genome Research</i> , 2021, 31, 645-658.	5.5	48
9	Anticytomegalovirus CD4 + T Cells Are Associated With Subclinical Atherosclerosis in Persons With HIV. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 1459-1473.	2.4	7
10	Visual Genomics Analysis Studio as a Tool to Analyze Multiomic Data. <i>Frontiers in Genetics</i> , 2021, 12, 642012.	2.3	14
11	The TCR repertoire of α -synuclein-specific T cells in Parkinson's disease is surprisingly diverse. <i>Scientific Reports</i> , 2021, 11, 302.	3.3	26
12	Relationship between CD4 T cell turnover, cellular differentiation and HIV persistence during ART. <i>PLoS Pathogens</i> , 2021, 17, e1009214.	4.7	25
13	Cross-Reactivity to Mutated Viral Immune Targets Can Influence CD8+ T Cell Functionality: An Alternative Viral Adaptation Strategy. <i>Frontiers in Immunology</i> , 2021, 12, 746986.	4.8	3
14	Generation of a Novel SARS-CoV-2 Sub-genomic RNA Due to the R203K/G204R Variant in Nucleocapsid: Homologous Recombination has Potential to Change SARS-CoV-2 at Both Protein and RNA Level. <i>Pathogens and Immunity</i> , 2021, 6, 27-49.	3.1	10
15	Generation of a Novel SARS-CoV-2 Sub-genomic RNA Due to the R203K/G204R Variant in Nucleocapsid: Homologous Recombination has Potential to Change SARS-CoV-2 at Both Protein and RNA Level. <i>Pathogens and Immunity</i> , 2021, 6, 27-49.	3.1	46
16	Severe COVID-19 Is Associated With an Altered Upper Respiratory Tract Microbiome. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 781968.	3.9	27
17	Identification of Novel Yellow Fever Class II Epitopes in YF-17D Vaccinees. <i>Viruses</i> , 2020, 12, 1300.	3.3	3
18	High Frequency of Shared Clonotypes in Human T Cell Receptor Repertoires. <i>Cell Reports</i> , 2020, 32, 107882.	6.4	39

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19	Selective and cross-reactive SARS-CoV-2 T cell epitopes in unexposed humans. <i>Science</i> , 2020, 370, 89-94.	12.6	1,036
20	Identification and Characterization of CD4 ⁺ T Cell Epitopes after Shingrix Vaccination. <i>Journal of Virology</i> , 2020, 94, .	3.4	18
21	Changes in Peripheral and Local Tumor Immunity after Neoadjuvant Chemotherapy Reshape Clinical Outcomes in Patients with Breast Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 5668-5681.	7.0	37
22	T Lymphocyte Subsets Associated With Prevalent Diabetes in Veterans With and Without Human Immunodeficiency Virus. <i>Journal of Infectious Diseases</i> , 2020, 222, 252-262.	4.0	11
23	Î±-Synuclein-specific T cell reactivity is associated with preclinical and early Parkinson's disease. <i>Nature Communications</i> , 2020, 11, 1875.	12.8	239
24	Characterization of Magnitude and Antigen Specificity of HLA-DP, DQ, and DRB3/4/5 Restricted DENV-Specific CD4 ⁺ T Cell Responses. <i>Frontiers in Immunology</i> , 2019, 10, 1568.	4.8	35
25	Single-cell transcriptomics reveal polyclonal memory T-cell responses in skin with positive abacavir patch test results. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 144, 1413-1416.e7.	2.9	19
26	Widespread Tau-Specific CD4 T Cell Reactivity in the General Population. <i>Journal of Immunology</i> , 2019, 203, 84-92.	0.8	36
27	343. T-cell Subsets Associated with Diabetes in Veterans with and without HIV. <i>Open Forum Infectious Diseases</i> , 2019, 6, S182-S183.	0.9	0
28	Clinical and evolutionary consequences of HIV adaptation to HLA. <i>Current Opinion in HIV and AIDS</i> , 2019, 14, 194-204.	3.8	6
29	Deep sequence analysis of HIV adaptation following vertical transmission reveals the impact of immune pressure on the evolution of HIV. <i>PLoS Pathogens</i> , 2019, 15, e1008177.	4.7	14
30	Deconstructing pulmonary fibrosis at single-cell resolution. <i>FASEB Journal</i> , 2019, 33, 847.3.	0.5	0
31	SJS/TEN 2017: Building Multidisciplinary Networks to Drive Science and Translation. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2018, 6, 38-69.	3.8	134
32	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 714 adults from Colombo, Sri Lanka. <i>Human Immunology</i> , 2018, 79, 87-88.	2.4	7
33	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 159 individuals from the Worcester region of the Western Cape province of South Africa. <i>Human Immunology</i> , 2018, 79, 143-144.	2.4	7
34	Pan-genomic analyses identify key <i>Helicobacter pylori</i> pathogenic loci modified by carcinogenic host microenvironments. <i>Gut</i> , 2018, 67, 1793-1804.	12.1	22
35	Differential Immunodominance Hierarchy of CD8 ⁺ T-Cell Responses in HLA-B*27:05- and -B*27:02-Mediated Control of HIV-1 Infection. <i>Journal of Virology</i> , 2018, 92, .	3.4	14
36	Adipose Tissue is Enriched for Activated and Late-Differentiated CD8 ⁺ T Cells and Shows Distinct CD8+ Receptor Usage, Compared With Blood in HIV-Infected Persons. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2018, 77, e14-e21.	2.1	37

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37	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 339 adults from Managua, Nicaragua. <i>Human Immunology</i> , 2018, 79, 1-2.	2.4	8
38	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 496 adults from San Diego, California, USA. <i>Human Immunology</i> , 2018, 79, 821-822.	2.4	10
39	Urinary Peptides As a Novel Source of T Cell Allergen Epitopes. <i>Frontiers in Immunology</i> , 2018, 9, 886.	4.8	16
40	Evidence of CD4+ T cell-mediated immune pressure on the Hepatitis C virus genome. <i>Scientific Reports</i> , 2018, 8, 7224.	3.3	16
41	Bacterial DNA is present in the fetal intestine and overlaps with that in the placenta in mice. <i>PLoS ONE</i> , 2018, 13, e0197439.	2.5	44
42	Rif1 inhibits replication fork progression and controls DNA copy number in <i>Drosophila</i> . <i>ELife</i> , 2018, 7, .	6.0	40
43	<i>Helicobacter pylori</i> genetic diversification in the Mongolian gerbil model. <i>PeerJ</i> , 2018, 6, e4803.	2.0	12
44	HIV integration sites in latently infected cell lines: evidence of ongoing replication. <i>Retrovirology</i> , 2017, 14, 2.	2.0	59
45	Patterns of Cellular Immunity Associated with Experimental Infection with rDEN2 ³⁰ (Tonga/74) Support Its Suitability as a Human Dengue Virus Challenge Strain. <i>Journal of Virology</i> , 2017, 91, .	3.4	24
46	Phenome-wide scanning identifies multiple diseases and disease severity phenotypes associated with HLA variants. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	105
47	T cells from patients with Parkinson's disease recognize α -synuclein peptides. <i>Nature</i> , 2017, 546, 656-661.	27.8	618
48	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 100 Luo infants from the Boro area of Nyanza Province, Kenya. <i>Human Immunology</i> , 2017, 78, 325-326.	2.4	6
49	Human CD4 ⁺ T Cell Responses to an Attenuated Tetravalent Dengue Vaccine Parallel Those Induced by Natural Infection in Magnitude, HLA Restriction, and Antigen Specificity. <i>Journal of Virology</i> , 2017, 91, .	3.4	83
50	Cytomegalovirus (CMV) Epitope-Specific CD4+ T Cells Are Inflated in HIV+ CMV+ Subjects. <i>Journal of Immunology</i> , 2017, 199, 3187-3201.	0.8	55
51	Prior Dengue Virus Exposure Shapes T Cell Immunity to Zika Virus in Humans. <i>Journal of Virology</i> , 2017, 91, .	3.4	148
52	Shared peptide binding of HLA Class I and II alleles associate with cutaneous nevirapine hypersensitivity and identify novel risk alleles. <i>Scientific Reports</i> , 2017, 7, 8653.	3.3	41
53	Influence of Human Leukocyte Antigen (HLA) Alleles and Killer Cell Immunoglobulin-Like Receptors (KIR) Types on Heparin-Induced Thrombocytopenia (HIT). <i>Pharmacotherapy</i> , 2017, 37, 1164-1171.	2.6	14
54	Potential for immune-driven viral polymorphisms to compromise antiretroviral-based preexposure prophylaxis for prevention of HIV-1 infection. <i>Aids</i> , 2017, 31, 1935-1943.	2.2	7

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55	Global Assessment of Dengue Virus-Specific CD4+ T Cell Responses in Dengue-Endemic Areas. <i>Frontiers in Immunology</i> , 2017, 8, 1309.	4.8	77
56	Definition of Human Epitopes Recognized in Tetanus Toxoid and Development of an Assay Strategy to Detect Ex Vivo Tetanus CD4+ T Cell Responses. <i>PLoS ONE</i> , 2017, 12, e0169086.	2.5	60
57	Comparison of HLA allelic imputation programs. <i>PLoS ONE</i> , 2017, 12, e0172444.	2.5	58
58	HIV-1 Mutation and Recombination Rates Are Different in Macrophages and T-cells. <i>Viruses</i> , 2016, 8, 118.	3.3	9
59	Impact of pre-adapted HIV transmission. <i>Nature Medicine</i> , 2016, 22, 606-613.	30.7	87
60	HLA-DRB1 Alleles Are Associated With Different Magnitudes of Dengue Virus-Specific CD4 ⁺ T-Cell Responses. <i>Journal of Infectious Diseases</i> , 2016, 214, 1117-1124.	4.0	88
61	Primer ID ultra-deep sequencing reveals dynamics of drug resistance-associated variants in breakthrough hepatitis C viruses: relevance to treatment outcome and resistance screening. <i>Antiviral Therapy</i> , 2016, 21, 567-577.	1.0	5
62	Abacavir-Reactive Memory T Cells Are Present in Drug Naïve Individuals. <i>PLoS ONE</i> , 2015, 10, e0117160.	2.5	73
63	Acyclovir Has Low but Detectable Influence on HLA-B*57:01 Specificity without Inducing Hypersensitivity. <i>PLoS ONE</i> , 2015, 10, e0124878.	2.5	11
64	Anti-Hepatitis C Virus T-Cell Immunity in the Context of Multiple Exposures to the Virus. <i>PLoS ONE</i> , 2015, 10, e0130420.	2.5	3
65	Polymorphisms of large effect explain the majority of the host genetic contribution to variation of HIV-1 virus load. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 14658-14663.	7.1	154
66	Evolving models of the immunopathogenesis of T _H 2-mediated drug allergy: The role of host, pathogens, and drug response. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 219-234.	2.9	185
67	A general method to eliminate laboratory induced recombinants during massive, parallel sequencing of cDNA library. <i>Virology Journal</i> , 2015, 12, 55.	3.4	14
68	Fibrosing mediastinitis complicating prior histoplasmosis is associated with human leukocyte antigen DQB1*04:02—a case control study. <i>BMC Infectious Diseases</i> , 2015, 15, 206.	2.9	15
69	<i>Helicobacter pylori</i> Adaptation <i>In Vivo</i> in Response to a High-Salt Diet. <i>Infection and Immunity</i> , 2015, 83, 4871-4883.	2.2	19
70	T Cell-Mediated Hypersensitivity Reactions to Drugs. <i>Annual Review of Medicine</i> , 2015, 66, 439-454.	12.2	109
71	Host-Specific Adaptation of HIV-1 Subtype B in the Japanese Population. <i>Journal of Virology</i> , 2014, 88, 4764-4775.	3.4	47
72	HLA class I drug-T cell receptor interactions in SJS/TEN. <i>Clinical and Translational Allergy</i> , 2014, 4, P2.	3.2	0

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73	Fever, Rash, and Systemic Symptoms: Understanding the Role of Virus and HLA in Severe Cutaneous Drug Allergy. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2014, 2, 21-33.	3.8	74
74	Influence of HLA-C Expression Level on HIV Control. <i>Science</i> , 2013, 340, 87-91.	12.6	352
75	Association Study of Common Genetic Variants and HIV-1 Acquisition in 6,300 Infected Cases and 7,200 Controls. <i>PLoS Pathogens</i> , 2013, 9, e1003515.	4.7	109
76	HIV escape mutations occur preferentially at HLA-binding sites of CD8 T-cell epitopes. <i>Aids</i> , 2013, 27, 899-905.	2.2	44
77	Drug hypersensitivity caused by alteration of the MHC-presented self-peptide repertoire. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 9959-9964.	7.1	354
78	Correlates of Protective Cellular Immunity Revealed by Analysis of Population-Level Immune Escape Pathways in HIV-1. <i>Journal of Virology</i> , 2012, 86, 13202-13216.	3.4	99
79	High avidity, high IFN γ -producing CD8 T cell responses following immune selection during HIV-1 infection. <i>Immunology and Cell Biology</i> , 2012, 90, 224-234.	2.3	36
80	HLA and pharmacogenetics of drug hypersensitivity. <i>Pharmacogenomics</i> , 2012, 13, 1285-1306.	1.3	161
81	The structural basis of HLA-associated drug hypersensitivity syndromes. <i>Immunological Reviews</i> , 2012, 250, 158-166.	6.0	31
82	Complications resulting from antiretroviral therapy for HIV infection. , 2012, , 177-191.		0
83	Extensive HLA-driven viral diversity following a narrow-source HIV-1 outbreak in rural China. <i>Blood</i> , 2011, 118, 98-106.	1.4	36
84	Mapping the Landscape of Host-Pathogen Coevolution: HLA Class I Binding and Its Relationship with Evolutionary Conservation in Human and Viral Proteins. <i>Journal of Virology</i> , 2011, 85, 1310-1321.	3.4	62
85	Translation of HLA-HIV Associations to the Cellular Level: HIV Adapts To Inflation CD8 T Cell Responses against Nef and HLA-Adapted Variant Epitopes. <i>Journal of Immunology</i> , 2011, 187, 2502-2513.	0.8	25
86	Adaptive Interactions between HLA and HIV-1: Highly Divergent Selection Imposed by HLA Class I Molecules with Common Supertype Motifs. <i>Journal of Immunology</i> , 2010, 184, 4368-4377.	0.8	39
87	Common Genetic Variation and the Control of HIV-1 in Humans. <i>PLoS Genetics</i> , 2009, 5, e1000791.	3.5	377
88	Divergent adaptation of hepatitis C virus genotypes 1 and 3 to human leukocyte antigen-restricted immune pressure. <i>Hepatology</i> , 2009, 50, 1017-1029.	7.3	60
89	Adaptation of HIV-1 to human leukocyte antigen class I. <i>Nature</i> , 2009, 458, 641-645.	27.8	408
90	Unique features of HLA-mediated HIV evolution in a Mexican cohort: a comparative study. <i>Retrovirology</i> , 2009, 6, 72.	2.0	33

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91	Successful Translation of Pharmacogenetics into the Clinic. <i>Molecular Diagnosis and Therapy</i> , 2009, 13, 1-9.	3.8	101
92	HLA-Associated Immune Escape Pathways in HIV-1 Subtype B Gag, Pol and Nef Proteins. <i>PLoS ONE</i> , 2009, 4, e6687.	2.5	148
93	HLA-B*5701 Screening for Hypersensitivity to Abacavir. <i>New England Journal of Medicine</i> , 2008, 358, 568-579.	27.0	1,665
94	Human Leukocyte Antigen Class I-Restricted Activation of CD8+ T Cells Provides the Immunogenetic Basis of a Systemic Drug Hypersensitivity. <i>Immunity</i> , 2008, 28, 822-832.	14.3	309
95	Marked Epitope- and Allele-Specific Differences in Rates of Mutation in Human Immunodeficiency Type 1 (HIV-1) Gag, Pol, and Nef Cytotoxic T-Lymphocyte Epitopes in Acute/Early HIV-1 Infection. <i>Journal of Virology</i> , 2008, 82, 9216-9227.	3.4	162
96	High Sensitivity of Human Leukocyte Antigen B*5701 as a Marker for Immunologically Confirmed Abacavir Hypersensitivity in White and Black Patients. <i>Clinical Infectious Diseases</i> , 2008, 46, 1111-1118.	5.8	384
97	Cytokine Profiling in Abacavir Hypersensitivity Patients. <i>Antiviral Therapy</i> , 2008, 13, 281-288.	1.0	22
98	Refining Abacavir Hypersensitivity Diagnoses using a Structured Clinical Assessment and Genetic Testing in the Swiss HIV Cohort Study. <i>Antiviral Therapy</i> , 2008, 13, 1019-1028.	1.0	29
99	Skewed association of polyfunctional antigen-specific CD8 T cell populations with HLA-B genotype. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 16233-16238.	7.1	118
100	HLA-B*5701 screening for susceptibility to abacavir hypersensitivity. <i>Journal of Antimicrobial Chemotherapy</i> , 2007, 59, 591-593.	3.0	61
101	Abacavir Hypersensitivity. , 2007, , 95-104.		1
102	Founder Effects in the Assessment of HIV Polymorphisms and HLA Allele Associations. <i>Science</i> , 2007, 315, 1583-1586.	12.6	234
103	Drug hypersensitivity in HIV. <i>Current Opinion in Allergy and Clinical Immunology</i> , 2007, 7, 324-330.	2.3	91
104	Immune responses to abacavir in antigen-presenting cells from hypersensitive patients. <i>Aids</i> , 2007, 21, 1233-1244.	2.2	48
105	A Whole-Genome Association Study of Major Determinants for Host Control of HIV-1. <i>Science</i> , 2007, 317, 944-947.	12.6	1,136
106	Extensive HLA class I allele promiscuity among viral CTL epitopes. <i>European Journal of Immunology</i> , 2007, 37, 2419-2433.	2.9	120
107	Leveraging Hierarchical Population Structure in Discrete Association Studies. <i>PLoS ONE</i> , 2007, 2, e591.	2.5	33
108	External Quality Assessment of HLA-B*5701 Reporting: An International Multicentre Survey. <i>Antiviral Therapy</i> , 2007, 12, 1027-1032.	1.0	40

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109	Prospective Genetic Screening Decreases the Incidence of Abacavir Hypersensitivity Reactions in the Western Australian HIV Cohort Study. <i>Clinical Infectious Diseases</i> , 2006, 43, 99-102.	5.8	301
110	Host genetics unplugged: removing the camouflage of viral adaptation. <i>Current Opinion in HIV and AIDS</i> , 2006, 1, 218-219.	3.8	0
111	A sensitive and rapid alternative to HLA typing as a genetic screening test for abacavir hypersensitivity syndrome. <i>Pharmacogenetics and Genomics</i> , 2006, 16, 353-357.	1.5	47
112	Control of human immunodeficiency virus replication by cytotoxic T lymphocytes targeting subdominant epitopes. <i>Nature Immunology</i> , 2006, 7, 173-178.	14.5	209
113	Host genetics and viral infections: immunology taught by viruses, virology taught by the immune system. <i>Current Opinion in Immunology</i> , 2006, 18, 413-421.	5.5	19
114	Predicting and diagnosing abacavir and nevirapine drug hypersensitivity: from bedside to bench and back again. <i>Pharmacogenomics</i> , 2006, 7, 15-23.	1.3	20
115	Impact of HLA-B Alleles, Epitope Binding Affinity, Functional Avidity, and Viral Coinfection on the Immunodominance of Virus-Specific CTL Responses. <i>Journal of Immunology</i> , 2006, 176, 4094-4101.	0.8	150
116	Transmission and accumulation of CTL escape variants drive negative associations between HIV polymorphisms and HLA. <i>Journal of Experimental Medicine</i> , 2005, 201, 891-902.	8.5	220
117	HIV/AIDS: HIV: Experiencing the Pressures of Modern Life. <i>Science</i> , 2005, 307, 1422-1424.	12.6	9
118	Adverse effects of antiretroviral therapy for HIV infection: a review of selected topics. <i>Expert Opinion on Drug Safety</i> , 2005, 4, 201-218.	2.4	43
119	HIV-1 Viral Escape in Infancy Followed by Emergence of a Variant-Specific CTL Response. <i>Journal of Immunology</i> , 2005, 174, 7524-7530.	0.8	109
120	Antiretroviral-therapy-associated lipotrophy: current status and future directions. <i>Sexual Health</i> , 2005, 2, 153.	0.9	39
121	Pharmacogenetics of antiretroviral therapy: genetic variation of response and toxicity. <i>Pharmacogenomics</i> , 2004, 5, 643-655.	1.3	33
122	Impact of host genetics on HIV disease progression and treatment. <i>Aids</i> , 2004, 18, 1231-1240.	2.2	20
123	Dominant influence of HLA-B in mediating the potential co-evolution of HIV and HLA. <i>Nature</i> , 2004, 432, 769-775.	27.8	784
124	Predisposition to abacavir hypersensitivity conferred by HLA-B*5701 and a haplotypic Hsp70-Hom variant. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 4180-4185.	7.1	451
125	Complications Associated with Nrti Therapy: Update on Clinical Features and Possible Pathogenic Mechanisms. <i>Antiviral Therapy</i> , 2004, 9, 849-863.	1.0	83
126	Contribution of Nucleoside-Analogue Reverse Transcriptase Inhibitor Therapy to Lipoatrophy from the Population to the Cellular Level. <i>Antiviral Therapy</i> , 2003, 8, 617-626.	1.0	101

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127	Antiretroviral Therapy and the Lipodystrophy Syndrome, Part 2: Concepts in Aetiopathogenesis. Antiviral Therapy, 2001, 6, 145-160.	1.0	54
128	The genetic basis for the association of the 8.1 ancestral haplotype (A1, B8, DR3) with multiple immunopathological diseases. Immunological Reviews, 1999, 167, 257-274.	6.0	506
129	Positive Selection of Mutations in the Helicobacter pylori <i>katA</i> 5' Untranslated Region in a Mongolian Gerbil Model of Gastric Disease. Infection and Immunity, 0, , .	2.2	3