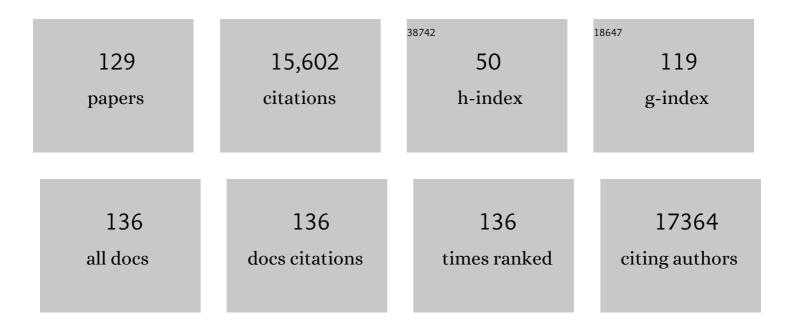
Simon Mallal

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	HLA-B*5701 Screening for Hypersensitivity to Abacavir. New England Journal of Medicine, 2008, 358, 568-579.	27.0	1,665
2	A Whole-Genome Association Study of Major Determinants for Host Control of HIV-1. Science, 2007, 317, 944-947.	12.6	1,136
3	Selective and cross-reactive SARS-CoV-2 T cell epitopes in unexposed humans. Science, 2020, 370, 89-94.	12.6	1,036
4	Dominant influence of HLA-B in mediating the potential co-evolution of HIV and HLA. Nature, 2004, 432, 769-775.	27.8	784
5	T cells from patients with Parkinson's disease recognize α-synuclein peptides. Nature, 2017, 546, 656-661.	27.8	618
6	SARS-CoV-2 vaccination induces immunological T cell memory able to cross-recognize variants from Alpha to Omicron. Cell, 2022, 185, 847-859.e11.	28.9	590
7	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
8	Predisposition to abacavir hypersensitivity conferred by HLA-B*5701 and a haplotypic Hsp70-Hom variant. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 4180-4185.	7.1	451
9	Comprehensive analysis of TÂcell immunodominance and immunoprevalence of SARS-CoV-2 epitopes in COVID-19 cases. Cell Reports Medicine, 2021, 2, 100204.	6.5	437
10	Adaptation of HIV-1 to human leukocyte antigen class I. Nature, 2009, 458, 641-645.	27.8	408
11	High Sensitivity of Human Leukocyte Antigen–B*5701 as a Marker for Immunologically Confirmed Abacavir Hypersensitivity in White and Black Patients. Clinical Infectious Diseases, 2008, 46, 1111-1118.	5.8	384
12	Common Genetic Variation and the Control of HIV-1 in Humans. PLoS Genetics, 2009, 5, e1000791.	3.5	377
13	Drug hypersensitivity caused by alteration of the MHC-presented self-peptide repertoire. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 9959-9964.	7.1	354
14	Influence of HLA-C Expression Level on HIV Control. Science, 2013, 340, 87-91.	12.6	352
15	Human Leukocyte Antigen Class I-Restricted Activation of CD8+ T Cells Provides the Immunogenetic Basis of a Systemic Drug Hypersensitivity. Immunity, 2008, 28, 822-832.	14.3	309
16	Prospective Genetic Screening Decreases the Incidence of Abacavir Hypersensitivity Reactions in the Western Australian HIV Cohort Study. Clinical Infectious Diseases, 2006, 43, 99-102.	5.8	301
17	α-Synuclein-specific T cell reactivity is associated with preclinical and early Parkinson's disease. Nature Communications, 2020, 11, 1875.	12.8	239
18	Founder Effects in the Assessment of HIV Polymorphisms and HLA Allele Associations. Science, 2007, 315, 1583-1586.	12.6	234

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19	Transmission and accumulation of CTL escape variants drive negative associations between HIV polymorphisms and HLA. Journal of Experimental Medicine, 2005, 201, 891-902.	8.5	220
20	Control of human immunodeficiency virus replication by cytotoxic T lymphocytes targeting subdominant epitopes. Nature Immunology, 2006, 7, 173-178.	14.5	209
21	Evolving models of the immunopathogenesis of TÂcell–mediated drug allergy: The role of host, pathogens, and drug response. Journal of Allergy and Clinical Immunology, 2015, 136, 219-234.	2.9	185
22	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. Journal of Virology, 2008, 82, 9216-9227.	3.4	162
23	HLA and pharmacogenetics of drug hypersensitivity. Pharmacogenomics, 2012, 13, 1285-1306.	1.3	161
24	Polymorphisms of large effect explain the majority of the host genetic contribution to variation of HIV-1 virus load. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 14658-14663.	7.1	154
25	Impact of HLA-B Alleles, Epitope Binding Affinity, Functional Avidity, and Viral Coinfection on the Immunodominance of Virus-Specific CTL Responses. Journal of Immunology, 2006, 176, 4094-4101.	0.8	150
26	Prior Dengue Virus Exposure Shapes T Cell Immunity to Zika Virus in Humans. Journal of Virology, 2017, 91, .	3.4	148
27	HLA-Associated Immune Escape Pathways in HIV-1 Subtype B Gag, Pol and Nef Proteins. PLoS ONE, 2009, 4, e6687.	2.5	148
28	SJS/TEN 2017: Building Multidisciplinary Networks to Drive Science and Translation. Journal of Allergy and Clinical Immunology: in Practice, 2018, 6, 38-69.	3.8	134
29	Extensive HLA class I allele promiscuity among viral CTL epitopes. European Journal of Immunology, 2007, 37, 2419-2433.	2.9	120
30	Skewed association of polyfunctional antigen-specific CD8 T cell populations with HLA-B genotype. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16233-16238.	7.1	118
31	HIV-1 Viral Escape in Infancy Followed by Emergence of a Variant-Specific CTL Response. Journal of Immunology, 2005, 174, 7524-7530.	0.8	109
32	Association Study of Common Genetic Variants and HIV-1 Acquisition in 6,300 Infected Cases and 7,200 Controls. PLoS Pathogens, 2013, 9, e1003515.	4.7	109
33	T Cell–Mediated Hypersensitivity Reactions to Drugs. Annual Review of Medicine, 2015, 66, 439-454.	12.2	109
34	Phenome-wide scanning identifies multiple diseases and disease severity phenotypes associated with HLA variants. Science Translational Medicine, 2017, 9, .	12.4	105
35	Successful Translation of Pharmacogenetics into the Clinic. Molecular Diagnosis and Therapy, 2009, 13, 1-9.	3.8	101
36	Contribution of Nucleoside-Analogue Reverse Transcriptase Inhibitor Therapy to Lipoatrophy from the Population to the Cellular Level. Antiviral Therapy, 2003, 8, 617-626.	1.0	101

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37	Correlates of Protective Cellular Immunity Revealed by Analysis of Population-Level Immune Escape Pathways in HIV-1. Journal of Virology, 2012, 86, 13202-13216.	3.4	99
38	Drug hypersensitivity in HIV. Current Opinion in Allergy and Clinical Immunology, 2007, 7, 324-330.	2.3	91
39	HLA-DRB1 Alleles Are Associated With Different Magnitudes of Dengue Virus–Specific CD4 ⁺ T-Cell Responses. Journal of Infectious Diseases, 2016, 214, 1117-1124.	4.0	88
40	Impact of pre-adapted HIV transmission. Nature Medicine, 2016, 22, 606-613.	30.7	87
41	Human CD4 ⁺ T Cell Responses to an Attenuated Tetravalent Dengue Vaccine Parallel Those Induced by Natural Infection in Magnitude, HLA Restriction, and Antigen Specificity. Journal of Virology, 2017, 91, .	3.4	83
42	Complications Associated with Nrti Therapy: Update on Clinical Features and Possible Pathogenic Mechanisms. Antiviral Therapy, 2004, 9, 849-863.	1.0	83
43	Global Assessment of Dengue Virus-Specific CD4+ T Cell Responses in Dengue-Endemic Areas. Frontiers in Immunology, 2017, 8, 1309.	4.8	77
44	Fever, Rash, and Systemic Symptoms: Understanding the Role of Virus and HLA in Severe Cutaneous Drug Allergy. Journal of Allergy and Clinical Immunology: in Practice, 2014, 2, 21-33.	3.8	74
45	Abacavir-Reactive Memory T Cells Are Present in Drug NaÃ⁻ve Individuals. PLoS ONE, 2015, 10, e0117160.	2.5	73
46	Mapping the Landscape of Host-Pathogen Coevolution: HLA Class I Binding and Its Relationship with Evolutionary Conservation in Human and Viral Proteins. Journal of Virology, 2011, 85, 1310-1321.	3.4	62
47	HLA-B*5701 screening for susceptibility to abacavir hypersensitivity. Journal of Antimicrobial Chemotherapy, 2007, 59, 591-593.	3.0	61
48	Divergent adaptation of hepatitis C virus genotypes 1 and 3 to human leukocyte antigen-restricted immune pressure. Hepatology, 2009, 50, 1017-1029.	7.3	60
49	Definition of Human Epitopes Recognized in Tetanus Toxoid and Development of an Assay Strategy to Detect Ex Vivo Tetanus CD4+ T Cell Responses. PLoS ONE, 2017, 12, e0169086.	2.5	60
50	HIV integration sites in latently infected cell lines: evidence of ongoing replication. Retrovirology, 2017, 14, 2.	2.0	59
51	Comparison of HLA allelic imputation programs. PLoS ONE, 2017, 12, e0172444.	2.5	58
52	Cytomegalovirus (CMV) Epitope–Specific CD4+ T Cells Are Inflated in HIV+ CMV+ Subjects. Journal of Immunology, 2017, 199, 3187-3201.	0.8	55
53	Antiretoviral Therapy and the Lipodystrophy Syndrome, Part 2: Concepts in Aetiopathogenesis. Antiviral Therapy, 2001, 6, 145-160.	1.0	54
54	Immune responses to abacavir in antigen-presenting cells from hypersensitive patients. Aids, 2007, 21, 1233-1244.	2.2	48

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55	Subgenomic RNA identification in SARS-CoV-2 genomic sequencing data. Genome Research, 2021, 31, 645-658.	5.5	48
56	A sensitive and rapid alternative to HLA typing as a genetic screening test for abacavir hypersensitivity syndrome. Pharmacogenetics and Genomics, 2006, 16, 353-357.	1.5	47
57	Host-Specific Adaptation of HIV-1 Subtype B in the Japanese Population. Journal of Virology, 2014, 88, 4764-4775.	3.4	47
58	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. Pathogens and Immunity, 2021, 6, 27-49.	3.1	46
59	HIV escape mutations occur preferentially at HLA-binding sites of CD8 T-cell epitopes. Aids, 2013, 27, 899-905.	2.2	44
60	Bacterial DNA is present in the fetal intestine and overlaps with that in the placenta in mice. PLoS ONE, 2018, 13, e0197439.	2.5	44
61	Adverse effects of antiretroviral therapy for HIV infection: a review of selected topics. Expert Opinion on Drug Safety, 2005, 4, 201-218.	2.4	43
62	Shared peptide binding of HLA Class I and II alleles associate with cutaneous nevirapine hypersensitivity and identify novel risk alleles. Scientific Reports, 2017, 7, 8653.	3.3	41
63	Rif1 inhibits replication fork progression and controls DNA copy number in Drosophila. ELife, 2018, 7, .	6.0	40
64	External Quality Assessment of <i>HLA-B*5701</i> Reporting: An International Multicentre Survey. Antiviral Therapy, 2007, 12, 1027-1032.	1.0	40
65	Adaptive Interactions between HLA and HIV-1: Highly Divergent Selection Imposed by HLA Class I Molecules with Common Supertype Motifs. Journal of Immunology, 2010, 184, 4368-4377.	0.8	39
66	High Frequency of Shared Clonotypes in Human T Cell Receptor Repertoires. Cell Reports, 2020, 32, 107882.	6.4	39
67	Antiretroviral-therapy-associated lipoatrophy: current status and future directions. Sexual Health, 2005, 2, 153.	0.9	39
68	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. Journal of Acquired Immune Deficiency Syndromes (1999), 2018, 77, e14-e21.	2.1	37
69	Changes in Peripheral and Local Tumor Immunity after Neoadjuvant Chemotherapy Reshape Clinical Outcomes in Patients with Breast Cancer. Clinical Cancer Research, 2020, 26, 5668-5681.	7.0	37
70	Extensive HLA-driven viral diversity following a narrow-source HIV-1 outbreak in rural China. Blood, 2011, 118, 98-106.	1.4	36
71	Highâ€avidity, highâ€IFNγâ€producing CD8 Tâ€cell responses following immune selection during HIVâ€1 infectio Immunology and Cell Biology, 2012, 90, 224-234.	on. 2.3	36
72	Widespread Tau-Specific CD4 T Cell Reactivity in the General Population. Journal of Immunology, 2019, 203, 84-92.	0.8	36

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73	Characterization of Magnitude and Antigen Specificity of HLA-DP, DQ, and DRB3/4/5 Restricted DENV-Specific CD4+ T Cell Responses. Frontiers in Immunology, 2019, 10, 1568.	4.8	35
74	Pharmacogenetics of antiretroviral therapy: genetic variation of response and toxicity. Pharmacogenomics, 2004, 5, 643-655.	1.3	33
75	Unique features of HLA-mediated HIV evolution in a Mexican cohort: a comparative study. Retrovirology, 2009, 6, 72.	2.0	33
76	Leveraging Hierarchical Population Structure in Discrete Association Studies. PLoS ONE, 2007, 2, e591.	2.5	33
77	The structural basis of <scp>HLA</scp> â€associated drug hypersensitivity syndromes. Immunological Reviews, 2012, 250, 158-166.	6.0	31
78	Refining Abacavir Hypersensitivity Diagnoses using a Structured Clinical Assessment and Genetic Testing in the Swiss HIV Cohort Study. Antiviral Therapy, 2008, 13, 1019-1028.	1.0	29
79	Severe COVID-19 Is Associated With an Altered Upper Respiratory Tract Microbiome. Frontiers in Cellular and Infection Microbiology, 2021, 11, 781968.	3.9	27
80	The TCR repertoire of α-synuclein-specific T cells in Parkinson's disease is surprisingly diverse. Scientific Reports, 2021, 11, 302.	3.3	26
81	Translation of HLA–HIV Associations to the Cellular Level: HIV Adapts To Inflate CD8 T Cell Responses against Nef and HLA-Adapted Variant Epitopes. Journal of Immunology, 2011, 187, 2502-2513.	0.8	25
82	Relationship between CD4 T cell turnover, cellular differentiation and HIV persistence during ART. PLoS Pathogens, 2021, 17, e1009214.	4.7	25
83	Patterns of Cellular Immunity Associated with Experimental Infection with rDEN2Δ30 (Tonga/74) Support Its Suitability as a Human Dengue Virus Challenge Strain. Journal of Virology, 2017, 91, .	3.4	24
84	Pan-genomic analyses identify key <i>Helicobacter pylori</i> pathogenic loci modified by carcinogenic host microenvironments. Gut, 2018, 67, 1793-1804.	12.1	22
85	Cytokine Profiling in Abacavir Hypersensitivity Patients. Antiviral Therapy, 2008, 13, 281-288.	1.0	22
86	Impact of host genetics on HIV disease progression and treatment. Aids, 2004, 18, 1231-1240.	2.2	20
87	Predicting and diagnosing abacavir and nevirapine drug hypersensitivity: from bedside to bench and back again. Pharmacogenomics, 2006, 7, 15-23.	1.3	20
88	Host genetics and viral infections: immunology taught by viruses, virology taught by the immune system. Current Opinion in Immunology, 2006, 18, 413-421.	5.5	19
89	Helicobacter pylori Adaptation <i>In Vivo</i> in Response to a High-Salt Diet. Infection and Immunity, 2015, 83, 4871-4883.	2.2	19
90	Single-cell transcriptomics reveal polyclonal memory T-cell responses in skin with positive abacavir patch test results. Journal of Allergy and Clinical Immunology, 2019, 144, 1413-1416.e7.	2.9	19

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91	Identification and Characterization of CD4 ⁺ T Cell Epitopes after Shingrix Vaccination. Journal of Virology, 2020, 94, .	3.4	18
92	Urinary Peptides As a Novel Source of T Cell Allergen Epitopes. Frontiers in Immunology, 2018, 9, 886.	4.8	16
93	Evidence of CD4+ T cell-mediated immune pressure on the Hepatitis C virus genome. Scientific Reports, 2018, 8, 7224.	3.3	16
94	Fibrosing mediastinitis complicating prior histoplasmosis is associated with human leukocyte antigen DQB1*04:02 â^' a case control study. BMC Infectious Diseases, 2015, 15, 206.	2.9	15
95	A general method to eliminate laboratory induced recombinants during massive, parallel sequencing of cDNA library. Virology Journal, 2015, 12, 55.	3.4	14
96	Influence of Human Leukocyte Antigen (<scp>HLA</scp>) Alleles and Killer Cell Immunoglobulinâ€Like Receptors (<scp>KIR</scp>) Types on Heparinâ€Induced Thrombocytopenia (<scp>HIT</scp>). Pharmacotherapy, 2017, 37, 1164-1171.	2.6	14
97	Differential Immunodominance Hierarchy of CD8 ⁺ T-Cell Responses in HLA-B*27:05- and -B*27:02-Mediated Control of HIV-1 Infection. Journal of Virology, 2018, 92, .	3.4	14
98	Deep sequence analysis of HIV adaptation following vertical transmission reveals the impact of immune pressure on the evolution of HIV. PLoS Pathogens, 2019, 15, e1008177.	4.7	14
99	Visual Genomics Analysis Studio as a Tool to Analyze Multiomic Data. Frontiers in Genetics, 2021, 12, 642012.	2.3	14
100	The Proximal Airway Is a Reservoir for Adaptive Immunologic Memory in Idiopathic Subglottic Stenosis. Laryngoscope, 2021, 131, 610-617.	2.0	12
101	<i>Helicobacter pylori</i> genetic diversification in the Mongolian gerbil model. PeerJ, 2018, 6, e4803.	2.0	12
102	Altered subgenomic RNA abundance provides unique insight into SARS-CoV-2 B.1.1.7/Alpha variant infections. Communications Biology, 2022, 5, .	4.4	12
103	Acyclovir Has Low but Detectable Influence on HLA-B*57:01 Specificity without Inducing Hypersensitivity. PLoS ONE, 2015, 10, e0124878.	2.5	11
104	T Lymphocyte Subsets Associated With Prevalent Diabetes in Veterans With and Without Human Immunodeficiency Virus. Journal of Infectious Diseases, 2020, 222, 252-262.	4.0	11
105	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 496 adults from San Diego, California, USA. Human Immunology, 2018, 79, 821-822.	2.4	10
106	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. Pathogens and Immunity, 2021, 6, 27-49.	3.1	10
107	HIV/AIDS: HIV: Experiencing the Pressures of Modern Life. Science, 2005, 307, 1422-1424.	12.6	9
108	HIV-1 Mutation and Recombination Rates Are Different in Macrophages and T-cells. Viruses, 2016, 8, 118.	3.3	9

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109	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 339 adults from Managua, Nicaragua. Human Immunology, 2018, 79, 1-2.	2.4	8
110	Immunodominant MHC-II (Major Histocompatibility Complex II) Restricted Epitopes in Human Apolipoprotein B. Circulation Research, 2022, 131, 258-276.	4.5	8
111	Deep Sequencing Analysis of Individual HIV-1 Proviruses Reveals Frequent Asymmetric Long Terminal Repeats. Journal of Virology, 2022, 96, .	3.4	8
112	Potential for immune-driven viral polymorphisms to compromise antiretroviral-based preexposure prophylaxis for prevention of HIV-1 infection. Aids, 2017, 31, 1935-1943.	2.2	7
113	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 714 adults from Colombo, Sri Lanka. Human Immunology, 2018, 79, 87-88.	2.4	7
114	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. Human Immunology, 2018, 79, 143-144.	2.4	7
115	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. Tuberculosis, 2021, 126, 102023.	1.9	7
116	Anticytomegalovirus CD4 + T Cells Are Associated With Subclinical Atherosclerosis in Persons With HIV. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 1459-1473.	2.4	7
117	Sequence-based HLA-A, B, C, DP, DQ, and DR typing of 100 Luo infants from the Boro area of Nyanza Province, Kenya. Human Immunology, 2017, 78, 325-326.	2.4	6
118	Clinical and evolutionary consequences of HIV adaptation to HLA. Current Opinion in HIV and AIDS, 2019, 14, 194-204.	3.8	6
119	Primer ID ultra-deep sequencing reveals dynamics of drug resistance-associated variants in breakthrough hepatitis C viruses: relevance to treatment outcome and resistance screening. Antiviral Therapy, 2016, 21, 567-577.	1.0	5
120	Anti-Hepatitis C Virus T-Cell Immunity in the Context of Multiple Exposures to the Virus. PLoS ONE, 2015, 10, e0130420.	2.5	3
121	Identification of Novel Yellow Fever Class II Epitopes in YF-17D Vaccinees. Viruses, 2020, 12, 1300.	3.3	3
122	Cross-Reactivity to Mutated Viral Immune Targets Can Influence CD8+ T Cell Functionality: An Alternative Viral Adaptation Strategy. Frontiers in Immunology, 2021, 12, 746986.	4.8	3
123	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
124	Abacavir Hypersensitivity. , 2007, , 95-104.		1
125	Host genetics unplugged: removing the camouflage of viral adaptation. Current Opinion in HIV and AIDS, 2006, 1, 218-219.	3.8	0
126	Complications resulting from antiretroviral therapy for HIV infection. , 2012, , 177-191.		0

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127	HLA class lâ€drugâ€Tâ€cell receptor interactions in SJS/TEN. Clinical and Translational Allergy, 2014, 4, P2.	3.2	ο
128	343. T-cell Subsets Associated with Diabetes in Veterans with and without HIV. Open Forum Infectious Diseases, 2019, 6, S182-S183.	0.9	0
129	Deconstructing pulmonary fibrosis at singleâ€eell resolution. FASEB Journal, 2019, 33, 847.3.	0.5	Ο