

Naoto Keicho

List of Publications by Year in descending order

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102
papers

3,366
citations

109311

35
h-index

175241

52
g-index

107
all docs

107
docs citations

107
times ranked

4182
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel Screening System of Virulent Strains for the Establishment of a <i>Mycobacterium avium</i> Complex Lung Disease Mouse Model Using Whole-Genome Sequencing. <i>Microbiology Spectrum</i> , 2022, 10, e0045122.	3.0	4
2	Phenotypic and genotypic features of the <i>Mycobacterium tuberculosis</i> lineage 1 subgroup in central Vietnam. <i>Scientific Reports</i> , 2021, 11, 13609.	3.3	2
3	Primary ciliary dyskinesia caused by a large homozygous deletion including exons 1–4 of <i>DRC1</i> in Japanese patients with recurrent sinopulmonary infection. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1033.	1.2	23
4	Genotyping of <i>Mycobacterium tuberculosis</i> spreading in Hanoi, Vietnam using conventional and whole genome sequencing methods. <i>Infection, Genetics and Evolution</i> , 2020, 78, 104107.	2.3	11
5	Membrane-associated mucins of the ocular surface: New genes, new protein functions and new biological roles in human and mouse. <i>Progress in Retinal and Eye Research</i> , 2020, 75, 100777.	15.5	30
6	Coordinated In Vitro Release of Granulysin, Perforin and IFN- γ in TB and HIV/TB Co-Infection Associated with Clinical Outcomes before and after Anti-TB Treatment. <i>Pathogens</i> , 2020, 9, 655.	2.8	4
7	Can Interferon- γ Release Assays Be Useful for Monitoring the Response to Anti-tuberculosis Treatment?: A Systematic Review and Meta-analysis. <i>Archivum Immunologiae Et Therapiae Experimentalis</i> , 2020, 68, 4.	2.3	7
8	Primary ciliary dyskinesia in Japan: systematic review and meta-analysis. <i>BMC Pulmonary Medicine</i> , 2019, 19, 135.	2.0	14
9	Recurring large deletion in <i>DRC1</i> (<i>CCDC164</i>) identified as causing primary ciliary dyskinesia in two Asian patients. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e838.	1.2	30
10	Whole-genome sequencing-based epidemiological analysis of anti-tuberculosis drug resistance genes in Japan in 2007: Application of the Genome Research for Asian Tuberculosis (GRaT) database. <i>Scientific Reports</i> , 2019, 9, 12823.	3.3	10
11	Whole genome sequencing, analyses of drug resistance-conferring mutations, and correlation with transmission of <i>Mycobacterium tuberculosis</i> carrying <i>katG</i> -S315T in Hanoi, Vietnam. <i>Scientific Reports</i> , 2019, 9, 15354.	3.3	20
12	Complete Genome Sequences of Three Representative <i>Mycobacterium tuberculosis</i> Beijing Family Strains Belonging to Distinct Genotype Clusters in Hanoi, Vietnam, during 2007 to 2009. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
13	Complete Genome Sequence of a <i>Mycobacterium tuberculosis</i> Strain Belonging to the East African-Indian Family in the Indo-Oceanic Lineage, Isolated in Hanoi, Vietnam. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
14	Spoligotyping and whole-genome sequencing analysis of lineage 1 strains of <i>Mycobacterium tuberculosis</i> in Da Nang, Vietnam. <i>PLoS ONE</i> , 2017, 12, e0186800.	2.5	8
15	Diffuse Panbronchiolitis. <i>Milestones in Drug Therapy</i> , 2017, , 21-38.	0.1	0
16	Circulating granulysin levels in healthcare workers and latent tuberculosis infection estimated using interferon-gamma release assays. <i>BMC Infectious Diseases</i> , 2016, 16, 580.	2.9	7
17	Identification of ITPA on chromosome 20 as a susceptibility gene for young-onset tuberculosis. <i>Human Genome Variation</i> , 2016, 3, 15067.	0.7	13
18	Variants near the HLA complex group 22 gene (HCG22) confer increased susceptibility to late-onset asthma in Japanese populations. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 138, 281-283.e13.	2.9	28

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19	Influence of the polymorphism of the DUSP14 gene on the expression of immune-related genes and development of pulmonary tuberculosis. <i>Genes and Immunity</i> , 2016, 17, 207-212.	4.1	13
20	Dynamics of immune parameters during the treatment of active tuberculosis showing negative interferon gamma response at the time of diagnosis. <i>International Journal of Infectious Diseases</i> , 2015, 40, 39-44.	3.3	11
21	Sublineages of <i>Mycobacterium tuberculosis</i> Beijing genotype strains and unfavorable outcomes of anti-tuberculosis treatment. <i>Tuberculosis</i> , 2015, 95, 336-342.	1.9	17
22	Identification of a Novel Mucin Gene <i>HCG22</i> Associated With Steroid-Induced Ocular Hypertension. , 2015, 56, 2737.		28
23	<i>Mycobacterium tuberculosis</i> strains spreading in Hanoi, Vietnam: Beijing sublineages, genotypes, drug susceptibility patterns, and host factors. <i>Tuberculosis</i> , 2014, 94, 649-656.	1.9	33
24	Association between tuberculosis recurrence and interferon- γ response during treatment. <i>Journal of Infection</i> , 2014, 69, 616-626.	3.3	8
25	Age-dependent association of mannose-binding lectin polymorphisms with the development of pulmonary tuberculosis in Viet Nam. <i>Human Immunology</i> , 2014, 75, 840-846.	2.4	14
26	MxA transcripts with distinct first exons and modulation of gene expression levels by single-nucleotide polymorphisms in human bronchial epithelial cells. <i>Immunogenetics</i> , 2013, 65, 107-114.	2.4	3
27	Differential effects of a common splice site polymorphism on the generation of OAS1 variants in human bronchial epithelial cells. <i>Human Immunology</i> , 2013, 74, 395-401.	2.4	14
28	Clonal expansion of <i>Mycobacterium tuberculosis</i> isolates and coexisting drug resistance in patients newly diagnosed with pulmonary tuberculosis in Hanoi, Vietnam. <i>BMC Research Notes</i> , 2013, 6, 444.	1.4	6
29	Potential Function of Granulysin, Other Related Effector Molecules and Lymphocyte Subsets in Patients with TB and HIV/TB Coinfection. <i>International Journal of Medical Sciences</i> , 2013, 10, 1003-1014.	2.5	19
30	Primary Drug-Resistant Tuberculosis in Hanoi, Viet Nam: Present Status and Risk Factors. <i>PLoS ONE</i> , 2013, 8, e71867.	2.5	28
31	Macrolide Therapy in Chronic Inflammatory Diseases. <i>Mediators of Inflammation</i> , 2012, 2012, 1-2.	3.0	5
32	Diffuse Panbronchiolitis. <i>Clinics in Chest Medicine</i> , 2012, 33, 297-305.	2.1	44
33	Association of SLC11A1 (NRAMP1) polymorphisms with pulmonary <i>Mycobacterium avium</i> complex infection. <i>Human Immunology</i> , 2012, 73, 529-536.	2.4	20
34	Circulating Levels of Adiponectin, Leptin, Fetuin-A and Retinol-Binding Protein in Patients with Tuberculosis: Markers of Metabolism and Inflammation. <i>PLoS ONE</i> , 2012, 7, e38703.	2.5	27
35	Association of IFNGR2 gene polymorphisms with pulmonary tuberculosis among the Vietnamese. <i>Human Genetics</i> , 2012, 131, 675-682.	3.8	24
36	Association of <i>TLR</i> polymorphisms with development of tuberculosis in Indonesian females. <i>Tissue Antigens</i> , 2012, 79, 190-197.	1.0	39

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37	Inter-rater agreement in the assessment of abnormal chest X-ray findings for tuberculosis between two Asian countries. <i>BMC Infectious Diseases</i> , 2012, 12, 31.	2.9	20
38	Association of CD209 polymorphisms with tuberculosis in an Indonesian population. <i>Human Immunology</i> , 2011, 72, 741-745.	2.4	22
39	Analysis of Factors Lowering Sensitivity of Interferon- γ Release Assay for Tuberculosis. <i>PLoS ONE</i> , 2011, 6, e23806.	2.5	58
40	Genetic predisposition to diffuse panbronchiolitis. <i>Respirology</i> , 2011, 16, 581-588.	2.3	37
41	Decreased plasma granulysin and increased interferon-gamma concentrations in patients with newly diagnosed and relapsed tuberculosis. <i>Microbiology and Immunology</i> , 2011, 55, 565-573.	1.4	11
42	Molecular cloning of two novel mucin-like genes in the disease-susceptibility locus for diffuse panbronchiolitis. <i>Human Genetics</i> , 2011, 129, 117-128.	3.8	64
43	Identification of tuberculosis-associated proteins in whole blood supernatant. <i>BMC Infectious Diseases</i> , 2011, 11, 71.	2.9	29
44	A Case of Familial Pulmonary Mycobacterium avium Complex Disease. <i>Internal Medicine</i> , 2010, 49, 949-953.	0.7	3
45	Association analysis of susceptibility candidate region on chromosome 5q31 for tuberculosis. <i>Genes and Immunity</i> , 2010, 11, 416-422.	4.1	20
46	Prevalence and Risk Factors for Tuberculosis Infection among Hospital Workers in Hanoi, Viet Nam. <i>PLoS ONE</i> , 2009, 4, e6798.	2.5	48
47	Identification of <i>MICA</i> as a Susceptibility Gene for Pulmonary <i>Mycobacterium avium</i> Complex Infection. <i>Journal of Infectious Diseases</i> , 2009, 199, 1707-1715.	4.0	23
48	No evidence for association between the interferon regulatory factor 1 (IRF1) gene and clinical tuberculosis. <i>Tuberculosis</i> , 2009, 89, 71-76.	1.9	6
49	Quality assessment of an interferon-gamma release assay for tuberculosis infection in a resource-limited setting. <i>BMC Infectious Diseases</i> , 2009, 9, 66.	2.9	9
50	Genome-wide SNP-based linkage analysis of tuberculosis in Thais. <i>Genes and Immunity</i> , 2009, 10, 77-83.	4.1	49
51	Association of human leukocyte antigen class II alleles with severe acute respiratory syndrome in the Vietnamese population. <i>Human Immunology</i> , 2009, 70, 527-531.	2.4	99
52	Granulocyte/macrophage colony-stimulating factor autoantibodies and myeloid cell immune functions in healthy subjects. <i>Blood</i> , 2009, 113, 2547-2556.	1.4	131
53	Granulocyte/macrophage colony-stimulating factor autoantibodies and myeloid cell immune functions in healthy subjects. <i>Blood</i> , 2009, 113, 2547-2556.	1.4	80
54	HLA-A, -B, -C, -DRB1 and -DQB1 alleles and haplotypes in the Kinh population in Vietnam. <i>Tissue Antigens</i> , 2008, 71, 127-134.	1.0	70

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55	Safe and Convenient Fluorescent Stain Imager for Proteome Analysis. <i>Current Proteomics</i> , 2007, 4, 115-120.	0.3	0
56	Pulmonary Mycobacterium avium complex infection: association with NRAMP1 polymorphisms. <i>European Respiratory Journal</i> , 2007, 30, 90-96.	6.7	38
57	Protein C Deficiency in a Family with Thromboembolism and Identified Gene Mutations. <i>Internal Medicine</i> , 2007, 46, 997-1003.	0.7	12
58	Anti-cytokine autoantibodies are ubiquitous in healthy individuals. <i>FEBS Letters</i> , 2007, 581, 2017-2021.	2.8	91
59	A survey of tuberculosis prevalence in Hanoi, Vietnam. <i>International Journal of Tuberculosis and Lung Disease</i> , 2007, 11, 562-6.	1.2	22
60	Pulmonary Mycobacterium avium complex infection associated with the IVS8-T5 allele of the CFTR gene. <i>International Journal of Tuberculosis and Lung Disease</i> , 2007, 11, 808-13.	1.2	21
61	Epidemiological and clinical features of idiopathic pulmonary alveolar proteinosis in Japan. <i>Respirology</i> , 2006, 11, S55-S60.	2.3	38
62	Identification of an alternative 5' untranslated exon and new polymorphisms of angiotensin-converting enzyme 2 gene: Lack of association with SARS in the Vietnamese population. <i>American Journal of Medical Genetics, Part A</i> , 2005, 136A, 52-57.	1.2	49
63	Variations of the CFTR gene in the Hanoi-Vietnamese. <i>American Journal of Medical Genetics, Part A</i> , 2005, 136A, 249-253.	1.2	13
64	Evaluation of microsatellite markers in association studies: a search for an immune-related susceptibility gene in sarcoidosis. <i>Immunogenetics</i> , 2005, 56, 861-870.	2.4	12
65	Promoter Analysis and Aberrant Expression of the MUC5B Gene in Diffuse Panbronchiolitis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2005, 171, 949-957.	5.6	64
66	Polymorphisms of interferon-inducible genes OAS-1 and MxA associated with SARS in the Vietnamese population. <i>Biochemical and Biophysical Research Communications</i> , 2005, 329, 1234-1239.	2.1	112
67	RAPID AWARENESS AND TRANSMISSION OF SEVERE ACUTE RESPIRATORY SYNDROME IN HANOI FRENCH HOSPITAL, VIETNAM. <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 73, 17-25.	1.4	57
68	Rapid awareness and transmission of severe acute respiratory syndrome in Hanoi French Hospital, Vietnam. <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 73, 17-25.	1.4	39
69	Assessment of synthetic peptides of severe acute respiratory syndrome coronavirus recognized by long-lasting immunity. <i>Tissue Antigens</i> , 2004, 64, 600-607.	1.0	15
70	Genome-wide linkage analysis of type 2 diabetes mellitus reconfirms the susceptibility locus on 11p13-p12 in Japanese. <i>Journal of Human Genetics</i> , 2004, 49, 629-634.	2.3	18
71	ACE1 polymorphism and progression of SARS. <i>Biochemical and Biophysical Research Communications</i> , 2004, 323, 1124-1129.	2.1	85
72	Direct determination of MUC5B promoter haplotypes based on the method of single-strand conformation polymorphism and their statistical estimation. <i>Genomics</i> , 2004, 84, 613-622.	2.9	10

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73	Diffuse Panbronchiolitis. <i>Seminars in Respiratory and Critical Care Medicine</i> , 2003, 24, 607-618.	2.1	18
74	High-affinity autoantibodies specifically eliminate granulocyte-macrophage colony-stimulating factor activity in the lungs of patients with idiopathic pulmonary alveolar proteinosis. <i>Blood</i> , 2003, 103, 1089-1098.	1.4	201
75	Adenoviral E1A modulates inflammatory mediator expression by lung epithelial cells exposed to PM ₁₀ . <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2003, 284, L290-L297.	2.9	19
76	Genotyping of Hepatitis E Virus from Vietnam. <i>Intervirolgy</i> , 2002, 45, 101-104.	2.8	23
77	Genetic Variants of Human β -Defensin-1 and Chronic Obstructive Pulmonary Disease. <i>Biochemical and Biophysical Research Communications</i> , 2002, 291, 17-22.	2.1	104
78	Diffuse Panbronchiolitis. <i>Treatments in Respiratory Medicine</i> , 2002, 1, 119-131.	1.2	88
79	Identification of novel candidate genes in the diffuse panbronchiolitis critical region of the class I human MHC. <i>Immunogenetics</i> , 2002, 54, 301-309.	2.4	30
80	Polymorphism of Human Leukocyte Antigen-E Gene in the Japanese Population with or without Recurrent Abortion. <i>American Journal of Reproductive Immunology</i> , 2001, 45, 168-173.	1.2	23
81	Association of Gc-globulin variation with susceptibility to COPD and diffuse panbronchiolitis. <i>European Respiratory Journal</i> , 2001, 18, 753-757.	6.7	62
82	Overestimated frequency of a possible emphysema-susceptibility allele when microsomal epoxide hydrolase is genotyped by the conventional polymerase chain reaction-based method. <i>Journal of Human Genetics</i> , 2001, 46, 96-98.	2.3	18
83	Human Alveolar Macrophages and Granulocyte-macrophage Colony-stimulating Factor-induced Monocyte-derived Macrophages Are Resistant to H ₂ O ₂ via Their High Basal and Inducible Levels of Catalase Activity. <i>Journal of Biological Chemistry</i> , 2001, 276, 24360-24364.	3.4	44
84	Prolonged survival of a bare lymphocyte syndrome type I patient with diffuse panbronchiolitis treated with erythromycin. <i>Sarcoidosis Vasculitis and Diffuse Lung Diseases</i> , 2001, 18, 312-3.	0.2	11
85	Fine Localization of a Major Disease-Susceptibility Locus for Diffuse Panbronchiolitis. <i>American Journal of Human Genetics</i> , 2000, 66, 501-507.	6.2	72
86	Endotoxin-specific NF- κ B activation in pulmonary epithelial cells harboring adenovirus E1A. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 1999, 277, L523-L532.	2.9	25
87	Contribution of TAP genes to genetic predisposition for diffuse panbronchiolitis. <i>Tissue Antigens</i> , 1999, 53, 366-373.	1.0	11
88	Association of diffuse panbronchiolitis with microsatellite polymorphism of the human interleukin 8 (IL-8) gene. <i>Journal of Human Genetics</i> , 1999, 44, 169-172.	2.3	25
89	Promoter variation of tumour necrosis factor-alpha gene: possible high risk for chronic bronchitis but not diffuse panbronchiolitis. <i>Respiratory Medicine</i> , 1999, 93, 752-753.	2.9	5
90	Splice acceptor site mutation of the transporter associated with antigen processing-1 gene in human bare lymphocyte syndrome. <i>Journal of Clinical Investigation</i> , 1999, 103, 755-758.	8.2	53

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91	Effect of adenovirus E1A on ICAM-1 promoter activity in human alveolar and bronchial epithelial cells. <i>Gene Expression</i> , 1999, 8, 287-97.	1.2	10
92	Contribution of HLA Genes to Genetic Predisposition in Diffuse Panbronchiolitis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 1998, 158, 846-850.	5.6	60
93	Identification of Glucocorticoid- and Adenovirus E1A-regulated Genes in Lung Epithelial Cells by Differential Display. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 1998, 18, 243-254.	2.9	4
94	Adenovirus E1A gene dysregulates ICAM-1 expression in transformed pulmonary epithelial cells.. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 1997, 16, 23-30.	2.9	52
95	A model of latent adenovirus 5 infection in the guinea pig (<i>Cavia porcellus</i>).. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 1996, 14, 225-231.	2.9	36
96	Erythromycin promotes monocyte to macrophage differentiation.. <i>Journal of Antibiotics</i> , 1994, 47, 80-89.	2.0	60
97	Detection of Lymphomatous Involvement of the Lung by Bronchoalveolar Lavage. <i>Chest</i> , 1994, 105, 458-462.	0.8	27
98	Antilymphocytic activity of erythromycin distinct from that of FK506 or cyclosporin A.. <i>Journal of Antibiotics</i> , 1993, 46, 1406-1413.	2.0	55
99	Effects of an immunosuppressant, FK506, on interleukin 1 β production by human macrophages and a macrophage-like cell line, U937. <i>Cellular Immunology</i> , 1991, 132, 285-294.	3.0	37
100	Serum Concentration of Soluble Interleukin-2 Receptor as a Sensitive Parameter of Disease Activity in Sarcoidosis. <i>Chest</i> , 1990, 98, 1125-1129.	0.8	81
101	Phase II Study of UFT in Patients With Advanced Non-Small Cell Lung Cancer. <i>Japanese Journal of Clinical Oncology</i> , 1986, 16, 143-146.	1.3	45
102	A Case Report of a Three-Year Survivor with Advanced Non-Small Cell Lung Cancer Producing Alfa-Fetoprotein. <i>Japanese Journal of Clinical Oncology</i> , 1986, 16, 175-181.	1.3	11