

Tomohiko Yamamura

List of Publications by Year in descending order

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70
papers

1,255
citations

471509

17
h-index

434195

31
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71
all docs

71
docs citations

71
times ranked

1274
citing authors

#	ARTICLE	IF	CITATIONS
1	A review of clinical characteristics and genetic backgrounds in Alport syndrome. <i>Clinical and Experimental Nephrology</i> , 2019, 23, 158-168.	1.6	135
2	Genetic, Clinical, and Pathologic Backgrounds of Patients with Autosomal Dominant Alport Syndrome. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2016, 11, 1441-1449.	4.5	94
3	Both the rituximab dose and maintenance immunosuppression in steroid-dependent/frequently-relapsing nephrotic syndrome have important effects on outcomes. <i>Kidney International</i> , 2020, 97, 393-401.	5.2	69
4	Differential diagnosis of Bartter syndrome, Gitelman syndrome, and pseudo-Bartter/Gitelman syndrome based on clinical characteristics. <i>Genetics in Medicine</i> , 2016, 18, 180-188.	2.4	67
5	Natural History and Genotype-Phenotype Correlation in Female X-Linked Alport Syndrome. <i>Kidney International Reports</i> , 2017, 2, 850-855.	0.8	62
6	Genotype-phenotype correlations influence the response to angiotensin-targeting drugs in Japanese patients with male X-linked Alport syndrome. <i>Kidney International</i> , 2020, 98, 1605-1614.	5.2	55
7	Comprehensive genetic diagnosis of Japanese patients with severe proteinuria. <i>Scientific Reports</i> , 2020, 10, 270.	3.3	50
8	Clinical and Genetic Characteristics in Patients With Gitelman Syndrome. <i>Kidney International Reports</i> , 2019, 4, 119-125.	0.8	47
9	Development of an exon skipping therapy for X-linked Alport syndrome with truncating variants in COL4A5. <i>Nature Communications</i> , 2020, 11, 2777.	12.8	46
10	Detection of Splicing Abnormalities and Genotype-Phenotype Correlation in X-linked Alport Syndrome. <i>Journal of the American Society of Nephrology: JASN</i> , 2018, 29, 2244-2254.	6.1	43
11	Common risk variants in NPHS1 and TNFSF15 are associated with childhood steroid-sensitive nephrotic syndrome. <i>Kidney International</i> , 2020, 98, 1308-1322.	5.2	39
12	Clinical characteristics of HNF1B-related disorders in a Japanese population. <i>Clinical and Experimental Nephrology</i> , 2019, 23, 1119-1129.	1.6	31
13	Detection of copy number variations by pair analysis using next-generation sequencing data in inherited kidney diseases. <i>Clinical and Experimental Nephrology</i> , 2018, 22, 881-888.	1.6	25
14	Comparison between conventional and comprehensive sequencing approaches for genetic diagnosis of Alport syndrome. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e883.	1.2	25
15	Mycophenolate Mofetil after Rituximab for Childhood-Onset Complicated Frequently-Relapsing or Steroid-Dependent Nephrotic Syndrome. <i>Journal of the American Society of Nephrology: JASN</i> , 2022, 33, 401-419.	6.1	24
16	Characterization of contiguous gene deletions in COL4A6 and COL4A5 in Alport syndrome-diffuse leiomyomatosis. <i>Journal of Human Genetics</i> , 2017, 62, 733-735.	2.3	20
17	Congenital chloride diarrhea needs to be distinguished from Bartter and Gitelman syndrome. <i>Journal of Human Genetics</i> , 2018, 63, 887-892.	2.3	20
18	Trimerization and Genotype-Phenotype Correlation of COL4A5 Mutants in Alport Syndrome. <i>Kidney International Reports</i> , 2020, 5, 718-726.	0.8	18

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19	Inherited salt-losing tubulopathy: An old condition but a new category of tubulopathy. <i>Pediatrics International</i> , 2020, 62, 428-437.	0.5	16
20	Pathogenic evaluation of synonymous <i>COL4A5</i> variants in X-linked Alport syndrome using a minigene assay. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1342.	1.2	16
21	A comparison of splicing assays to detect an intronic variant of the OCRL gene in Lowe syndrome. <i>European Journal of Medical Genetics</i> , 2017, 60, 631-634.	1.3	15
22	Clinical and genetic variability of PAX2-related disorder in the Japanese population. <i>Journal of Human Genetics</i> , 2020, 65, 541-549.	2.3	15
23	Determination of the pathogenicity of known COL4A5 intronic variants by in vitro splicing assay. <i>Scientific Reports</i> , 2019, 9, 12696.	3.3	14
24	Last Nucleotide Substitutions of COL4A5 Exons Cause Aberrant Splicing. <i>Kidney International Reports</i> , 2022, 7, 108-116.	0.8	14
25	Female X-linked Alport syndrome with somatic mosaicism. <i>Clinical and Experimental Nephrology</i> , 2017, 21, 877-883.	1.6	13
26	Examination of the predicted prevalence of Gitelman syndrome by ethnicity based on genome databases. <i>Scientific Reports</i> , 2021, 11, 16099.	3.3	13
27	Genetic background, recent advances in molecular biology, and development of novel therapy in Alport syndrome. <i>Kidney Research and Clinical Practice</i> , 2020, 39, 402-413.	2.2	13
28	Cryptic exon activation in SLC12A3 in Gitelman syndrome. <i>Journal of Human Genetics</i> , 2017, 62, 335-337.	2.3	12
29	The utility of urinary CD80 as a diagnostic marker in patients with renal diseases. <i>Scientific Reports</i> , 2018, 8, 17322.	3.3	12
30	Molecular assay for an intronic variant in NUP93 that causes steroid resistant nephrotic syndrome. <i>Journal of Human Genetics</i> , 2019, 64, 673-679.	2.3	12
31	Clinical spectrum of male patients with OFD1 mutations. <i>Journal of Human Genetics</i> , 2019, 64, 3-9.	2.3	12
32	Systematic Review of Genotype-Phenotype Correlations in Frasier Syndrome. <i>Kidney International Reports</i> , 2021, 6, 2585-2593.	0.8	12
33	Clinical characteristics and long-term outcome of diarrhea-associated hemolytic uremic syndrome: a single center experience. <i>Clinical and Experimental Nephrology</i> , 2017, 21, 889-894.	1.6	10
34	An in vitro splicing assay reveals the pathogenicity of a novel intronic variant in ATP6V0A4 for autosomal recessive distal renal tubular acidosis. <i>BMC Nephrology</i> , 2017, 18, 353.	1.8	10
35	Development of ultra-deep targeted RNA sequencing for analyzing X-chromosome inactivation in female Dent disease. <i>Journal of Human Genetics</i> , 2018, 63, 589-595.	2.3	10
36	The Contribution of COL4A5 Splicing Variants to the Pathogenesis of X-Linked Alport Syndrome. <i>Frontiers in Medicine</i> , 2022, 9, 841391.	2.6	10

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37	Rare renal ciliopathies in non-consanguineous families that were identified by targeted resequencing. <i>Clinical and Experimental Nephrology</i> , 2017, 21, 136-142.	1.6	9
38	Functional analysis of suspected splicing variants in CLCN5 gene in Dent disease 1. <i>Clinical and Experimental Nephrology</i> , 2020, 24, 606-612.	1.6	9
39	X-chromosome inactivation patterns in females with Fabry disease examined by both ultra-deep RNA sequencing and methylation-dependent assay. <i>Clinical and Experimental Nephrology</i> , 2021, 25, 1224-1230.	1.6	9
40	Pathogenesis of hypokalemia in autosomal dominant hypocalcemia type 1. <i>Clinical and Experimental Nephrology</i> , 2016, 20, 253-257.	1.6	8
41	Diagnostic strategy for inherited hypomagnesemia. <i>Clinical and Experimental Nephrology</i> , 2017, 21, 1003-1010.	1.6	8
42	Pair analysis and custom array CGH can detect a small copy number variation in COQ6 gene. <i>Clinical and Experimental Nephrology</i> , 2019, 23, 669-675.	1.6	8
43	Glomerular galactose-deficient IgA1 expression analysis in pediatric patients with glomerular diseases. <i>Scientific Reports</i> , 2020, 10, 14026.	3.3	8
44	Molecular mechanisms determining severity in patients with Pierson syndrome. <i>Journal of Human Genetics</i> , 2020, 65, 355-362.	2.3	8
45	Functional splicing analysis in an infantile case of atypical hemolytic uremic syndrome caused by digenic mutations in C3 and MCP genes. <i>Journal of Human Genetics</i> , 2018, 63, 755-759.	2.3	7
46	Genotype-Phenotype Correlation in WT1 Exon 8 to 9 Missense Variants. <i>Kidney International Reports</i> , 2021, 6, 2114-2121.	0.8	7
47	Detecting MUC1 Variants in Patients Clinicopathologically Diagnosed With Having Autosomal Dominant Tubulointerstitial Kidney Disease. <i>Kidney International Reports</i> , 2022, 7, 857-866.	0.8	7
48	Changes in the numbers of patients with acute gastroenteritis after voluntary introduction of the rotavirus vaccine in a Japanese children's primary emergency medical center. <i>Environmental Health and Preventive Medicine</i> , 2017, 22, 15.	3.4	5
49	Detection of a Splice Site Variant in a Patient with Glomerulopathy and Fibronectin Deposits. <i>Nephron</i> , 2018, 138, 166-171.	1.8	5
50	Comparison of clinical and genetic characteristics between Dent disease 1 and Dent disease 2. <i>Pediatric Nephrology</i> , 2020, 35, 2319-2326.	1.7	5
51	BCS1L mutations produce Fanconi syndrome with developmental disability. <i>Journal of Human Genetics</i> , 2022, 67, 143-148.	2.3	5
52	Use of renin-angiotensin system inhibitors as initial therapy in children with Henoch-Schönlein purpura nephritis of moderate severity. <i>Pediatric Nephrology</i> , 2022, 37, 1845-1853.	1.7	5
53	Comprehensive genetic analysis using next-generation sequencing for the diagnosis of nephronophthisis-related ciliopathies in the Japanese population. <i>Journal of Human Genetics</i> , 2022, 67, 427-440.	2.3	5
54	A Novel Mutation in a Japanese Family with X-linked Alport Syndrome. <i>Internal Medicine</i> , 2016, 55, 2843-2847.	0.7	4

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55	A birth of bipartite exon by intragenic deletion. <i>Molecular Genetics & Genomic Medicine</i> , 2017, 5, 287-294.	1.2	4
56	TGFBI-associated corneal dystrophy and nephropathy: a novel syndrome?. <i>CEN Case Reports</i> , 2019, 8, 14-17.	0.9	3
57	FAT1 biallelic truncating mutation causes a non-syndromic proteinuria in a child. <i>CEN Case Reports</i> , 2021, 10, 100-105.	0.9	3
58	Usefulness of functional splicing analysis to confirm precise disease pathogenesis in Diamond-Blackfan anemia caused by intronic variants in RPS19. <i>Pediatric Hematology and Oncology</i> , 2021, 38, 515-527.	0.8	3
59	Clinical and histological features in pediatric and adolescent/young adult patients with renal disease: a cross-sectional analysis of the Japan Renal Biopsy Registry (J-RBR). <i>Clinical and Experimental Nephrology</i> , 2021, 25, 1018-1026.	1.6	3
60	Evaluation of suspected autosomal Alport Syndrome synonymous variants. <i>Kidney360</i> , 2022, 3, 10.34067/KID.0005252021.	2.1	3
61	Detecting pathogenic deep intronic variants in <scp>Gitelman</scp> syndrome. <i>American Journal of Medical Genetics, Part A</i> , 0, , .	1.2	3
62	Poststreptococcal acute glomerulonephritis can be a risk factor for accelerating kidney dysfunction in Alport syndrome: a case experience. <i>CEN Case Reports</i> , 2020, 9, 418-422.	0.9	2
63	Clinical features of autosomal recessive polycystic kidney disease in the Japanese population and analysis of splicing in PKHD1 gene for determination of phenotypes. <i>Clinical and Experimental Nephrology</i> , 2022, 26, 140-153.	1.6	2
64	Heterozygous Urinary Abnormalityâ€‘Causing Variants of COL4A3 and COL4A4 Affect Severity of Autosomal Recessive Alport Syndrome. <i>Kidney360</i> , 2020, 1, 936-942.	2.1	2
65	A disease-causing variant of COL4A5 in a Chinese family with Alport syndrome: a case series. <i>BMC Nephrology</i> , 2021, 22, 380.	1.8	2
66	Three Severe Cases of Viral Infections with Post-Kidney Transplantation Successfully Confirmed by Polymerase Chain Reaction and Flow Cytometry. <i>Case Reports in Nephrology and Dialysis</i> , 2018, 8, 198-206.	0.6	1
67	A case with somatic and germline mosaicism in COL4A5 detected by multiplex ligation-dependent probe amplification in X-linked Alport syndrome. <i>CEN Case Reports</i> , 2020, 9, 431-436.	0.9	1
68	Utility of glomerular Gd-IgA1 staining for indistinguishable cases of IgA nephropathy or Alport syndrome. <i>Clinical and Experimental Nephrology</i> , 2021, 25, 779-787.	1.6	1
69	Efficacy of combination therapy for childhood complicated focal IgA nephropathy. <i>Clinical and Experimental Nephrology</i> , 2022, , 1.	1.6	1
70	Identification of novel OCRL isoforms associated with phenotypic differences between Dent disease-2 and Lowe syndrome. <i>Nephrology Dialysis Transplantation</i> , 2021, , .	0.7	0