

# Yoshihiro Taguchi

## List of Publications by Year in descending order

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Version: 2024-02-01

150  
papers

2,864  
citations

185998

28  
h-index

233125

45  
g-index

190  
all docs

190  
docs citations

190  
times ranked

2388  
citing authors

#	ARTICLE	IF	CITATIONS
1	Tumor Heterogeneity and Molecular Characteristics of Glioblastoma Revealed by Single-Cell RNA-Seq Data Analysis. <i>Genes</i> , 2022, 13, 428.	1.0	2
2	Novel feature selection method via kernel tensor decomposition for improved multi-omics data analysis. <i>BMC Medical Genomics</i> , 2022, 15, 37.	0.7	4
3	Integrated Analysis of Tissue-Specific Gene Expression in Diabetes by Tensor Decomposition Can Identify Possible Associated Diseases. <i>Genes</i> , 2022, 13, 1097.	1.0	0
4	Application of PCA based unsupervised FE to neurodegenerative diseases. , 2021, , 131-144.		0
5	Identification of genes associated with altered gene expression and m6A profiles during hypoxia using tensor decomposition based unsupervised feature extraction. <i>Scientific Reports</i> , 2021, 11, 8909.	1.6	10
6	Editorial: miRNAs and Neurological Diseases. <i>Frontiers in Neurology</i> , 2021, 12, 662373.	1.1	9
7	Application of Tensor Decomposition to Gene Expression of Infection of Mouse Hepatitis Virus Can Identify Critical Human Genes and Effective Drugs for SARS-CoV-2 Infection. <i>IEEE Journal on Selected Topics in Signal Processing</i> , 2021, 15, 746-758.	7.3	5
8	Mathematical formulation and application of kernel tensor decomposition based unsupervised feature extraction. <i>Knowledge-Based Systems</i> , 2021, 217, 106834.	4.0	4
9	Unsupervised tensor decomposition-based method to extract candidate transcription factors as histone modification bookmarks in post-mitotic transcriptional reactivation. <i>PLoS ONE</i> , 2021, 16, e0251032.	1.1	3
10	Novel method for the prediction of drug-drug Interaction based on gene expression profiles. <i>European Journal of Pharmaceutical Sciences</i> , 2021, 160, 105742.	1.9	4
11	Discriminating the single-cell gene regulatory networks of human pancreatic islets: A novel deep learning application. <i>Computers in Biology and Medicine</i> , 2021, 132, 104257.	3.9	11
12	PCA-based unsupervised feature extraction for gene expression analysis of COVID-19 patients. <i>Scientific Reports</i> , 2021, 11, 17351.	1.6	11
13	Tensor-Decomposition-Based Unsupervised Feature Extraction in Single-Cell Multiomics Data Analysis. <i>Genes</i> , 2021, 12, 1442.	1.0	8
14	In Silico Drug Discovery for COVID-19 Using an Unsupervised Feature Extraction Method. , 2021, , .		0
15	Identification of Transcription Factors, Biological Pathways, and Diseases as Mediated by N6-methyladenosine Using Tensor Decomposition-Based Unsupervised Feature Extraction. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 213.	1.3	2
16	Identification of Enhancers and Promoters in the Genome by Multidimensional Scaling. <i>Genes</i> , 2021, 12, 1671.	1.0	1
17	End-to-End Deep Learning for Detecting Metastatic Breast Cancer in Axillary Lymph Node from Digital Pathology Images. <i>Lecture Notes in Computer Science</i> , 2021, , 343-353.	1.0	0
18	Effects of Collagenâ€™Glycosaminoglycan Mesh on Gene Expression as Determined by Using Principal Component Analysis-Based Unsupervised Feature Extraction. <i>Polymers</i> , 2021, 13, 4117.	2.0	1

#	ARTICLE	IF	CITATIONS
19	Unsupervised Feature Extraction Applied to Bioinformatics. Unsupervised and Semi-supervised Learning, 2020, , .	0.4	48
20	Development of a novel anti-hepatitis B virus agent via Sp1. Scientific Reports, 2020, 10, 47.	1.6	8
21	Identifying suitable tools for variant detection and differential gene expression using RNA-seq data. Genomics, 2020, 112, 2166-2172.	1.3	8
22	Incremental Dilations Using CNN for Brain Tumor Classification. Applied Sciences (Switzerland), 2020, 10, 4915.	1.3	27
23	Identification of miRNA signatures for kidney renal clear cell carcinoma using the tensor-decomposition method. Scientific Reports, 2020, 10, 15149.	1.6	15
24	A new advanced in silico drug discovery method for novel coronavirus (SARS-CoV-2) with tensor decomposition-based unsupervised feature extraction. PLoS ONE, 2020, 15, e0238907.	1.1	38
25	Universal Nature of Drug Treatment Responses in Drug-Tissue-Wide Model-Animal Experiments Using Tensor Decomposition-Based Unsupervised Feature Extraction. Frontiers in Genetics, 2020, 11, 695.	1.1	3
26	Developing a diagnostic method for latent tuberculosis infection using circulating miRNA. Translational Medicine Communications, 2020, 5, .	0.5	5
27	Tensor-Decomposition-Based Unsupervised Feature Extraction Applied to Prostate Cancer Multiomics Data. Genes, 2020, 11, 1493.	1.0	3
28	Comprehensive analysis of liver and blood miRNA in precancerous conditions. Scientific Reports, 2020, 10, 21766.	1.6	11
29	SCGRNs: Novel supervised inference of single-cell gene regulatory networks of complex diseases. Computers in Biology and Medicine, 2020, 118, 103656.	3.9	14
30	Neurological Disorder Drug Discovery from Gene Expression with Tensor Decomposition. Current Pharmaceutical Design, 2020, 25, 4589-4599.	0.9	6
31	TD Based Unsupervised FE. Unsupervised and Semi-supervised Learning, 2020, , 103-116.	0.4	0
32	PCA Based Unsupervised FE. Unsupervised and Semi-supervised Learning, 2020, , 81-102.	0.4	0
33	Application of TD Based Unsupervised FE to Bioinformatics. Unsupervised and Semi-supervised Learning, 2020, , 213-296.	0.4	0
34	Introduction to Linear Algebra. Unsupervised and Semi-supervised Learning, 2020, , 3-22.	0.4	0
35	Applications of PCA Based Unsupervised FE to Bioinformatics. Unsupervised and Semi-supervised Learning, 2020, , 119-211.	0.4	1
36	Exploring the selective vulnerability in Alzheimer disease using tissue specific variant analysis. Genomics, 2019, 111, 936-949.	1.3	19

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37	Comparative Transcriptomics Analysis. , 2019, , 814-818.		4
38	Regulation of Gene Expression. , 2019, , 806-813.		1
39	Multomics Data Analysis Using Tensor Decomposition Based Unsupervised Feature Extraction. Lecture Notes in Computer Science, 2019, , 565-574.	1.0	3
40	Tensor Decomposition-Based Unsupervised Feature Extraction Applied to Single-Cell Gene Expression Analysis. Frontiers in Genetics, 2019, 10, 864.	1.1	13
41	Machine learning algorithms for predicting drugsâ€™tissues relationships. Expert Systems With Applications, 2019, 127, 167-186.	4.4	12
42	Drug candidate identification based on gene expression of treated cells using tensor decomposition-based unsupervised feature extraction for large-scale data. BMC Bioinformatics, 2019, 19, 388.	1.2	34
43	Investigating the energy crisis in Alzheimer disease using transcriptome study. Scientific Reports, 2019, 9, 18509.	1.6	23
44	Tensor Decomposition Based Unsupervised Feature Extraction Applied to Bioinformatics. , 2019, , 159-187.		5
45	Discovery of a Robust Gene Regulatory Network with a Complex Transcription Factor Network on Organ Cancer Cell-line RNA Sequence Data. Chem-Bio Informatics Journal, 2019, 19, 32-55.	0.1	0
46	Collaborative environmental DNA sampling from petal surfaces of flowering cherry <i>Cerasus yedoensis</i> 'Somei-yoshino' across the Japanese archipelago. Journal of Plant Research, 2018, 131, 709-717.		1
47	[Regular Paper] Tensor Decomposition-Based Unsupervised Feature Extraction for Integrated Analysis of TCGA Data on MicroRNA Expression and Promoter Methylation of Genes in Ovarian Cancer. , 2018, , .		14
48	Exploring MicroRNA Biomarkers for Parkinson's Disease from mRNA Expression Profiles. Cells, 2018, 7, 245.	1.8	34
49	Tensor Decomposition-Based Unsupervised Feature Extraction Can Identify the Universal Nature of Sequence-Nonspecific Off-Target Regulation of mRNA Mediated by MicroRNA Transfection. Cells, 2018, 7, 54.	1.8	17
50	Tensor decomposition-based and principal-component-analysis-based unsupervised feature extraction applied to the gene expression and methylation profiles in the brains of social insects with multiple castes. BMC Bioinformatics, 2018, 19, 99.	1.2	37
51	Exploring microRNA Biomarker for Amyotrophic Lateral Sclerosis. International Journal of Molecular Sciences, 2018, 19, 1318.	1.8	44
52	Principal Component Analysis-Based Unsupervised Feature Extraction Applied to Single-Cell Gene Expression Analysis. Lecture Notes in Computer Science, 2018, , 816-826.	1.0	12
53	Principal Components Analysis Based Unsupervised Feature Extraction Applied to Gene Expression Analysis of Blood from Dengue Haemorrhagic Fever Patients. Scientific Reports, 2017, 7, 44016.	1.6	32
54	Microarray analysis of circulating microRNAs in familial Mediterranean fever. Modern Rheumatology, 2017, 27, 1040-1046.	0.9	22

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55	Identification of candidate drugs using tensor-decomposition-based unsupervised feature extraction in integrated analysis of gene expression between diseases and DrugMatrix datasets. <i>Scientific Reports</i> , 2017, 7, 13733.	1.6	40
56	Principal component analysis based unsupervised feature extraction applied to bioinformatics analysis. , 2017, , 153-182.		10
57	An iterative compound screening contest method for identifying target protein inhibitors using the tyrosine-protein kinase Yes. <i>Scientific Reports</i> , 2017, 7, 12038.	1.6	28
58	One-class Differential Expression Analysis using Tensor Decomposition-based Unsupervised Feature Extraction Applied to Integrated Analysis of Multiple Omics Data from 26 Lung Adenocarcinoma Cell Lines. , 2017, , .		20
59	Genetic Association between Amyotrophic Lateral Sclerosis and Cancer. <i>Genes</i> , 2017, 8, 243.	1.0	27
60	Expression of Serum Exosomal and Esophageal MicroRNA in Rat Reflux Esophagitis. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1611.	1.8	6
61	Tensor decomposition-based unsupervised feature extraction identifies candidate genes that induce post-traumatic stress disorder-mediated heart diseases. <i>BMC Medical Genomics</i> , 2017, 10, 67.	0.7	15
62	Identification of Candidate Drugs for Heart Failure Using Tensor Decomposition-Based Unsupervised Feature Extraction Applied to Integrated Analysis of Gene Expression Between Heart Failure and DrugMatrix Datasets. <i>Lecture Notes in Computer Science</i> , 2017, , 517-528.	1.0	10
63	Tensor decomposition-based unsupervised feature extraction applied to matrix products for multi-view data processing. <i>PLoS ONE</i> , 2017, 12, e0183933.	1.1	27
64	Identification of More Feasible MicroRNA-mRNA Interactions within Multiple Cancers Using Principal Component Analysis Based Unsupervised Feature Extraction. <i>International Journal of Molecular Sciences</i> , 2016, 17, 696.	1.8	35
65	Principal component analysis based unsupervised feature extraction applied to publicly available gene expression profiles provides new insights into the mechanisms of action of histone deacetylase inhibitors. <i>Neuroepigenetics</i> , 2016, 8, 1-18.	2.8	23
66	microRNA-mRNA Interaction Identification in Wilms Tumor Using Principal Component Analysis Based Unsupervised Feature Extraction. , 2016, , .		17
67	Comparative Gene Expression Analysis of Mouse and Human Cardiac Maturation. <i>Genomics, Proteomics and Bioinformatics</i> , 2016, 14, 207-215.	3.0	40
68	SFRP1 is a possible candidate for epigenetic therapy in non-small cell lung cancer. <i>BMC Medical Genomics</i> , 2016, 9, 28.	0.7	42
69	Principal component analysis based unsupervised feature extraction applied to budding yeast temporally periodic gene expression. <i>BioData Mining</i> , 2016, 9, 22.	2.2	36
70	MicroRNA expression in hepatocellular carcinoma after the eradication of chronic hepatitis virus C infection using interferon therapy. <i>Hepatology Research</i> , 2016, 46, E26-35.	1.8	15
71	Identification of potential inhibitors based on compound proposal contest: Tyrosine-protein kinase Yes as a target. <i>Scientific Reports</i> , 2015, 5, 17209.	1.6	33
72	<i>In silico</i> Spleen Tyrosine Kinase Inhibitor Screening by ChooseLD. <i>IPSI Transactions on Bioinformatics</i> , 2015, 8, 14-20.	0.2	0

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73	Comprehensive analysis of transcriptome and metabolome analysis in Intrahepatic Cholangiocarcinoma and Hepatocellular Carcinoma. <i>Scientific Reports</i> , 2015, 5, 16294.	1.6	104
74	Identification of aberrant gene expression associated with aberrant promoter methylation in primordial germ cells between E13 and E16 rat F3 generation vinclozolin lineage. <i>BMC Bioinformatics</i> , 2015, 16, S16.	1.2	27
75	Apparent microRNA-Target-specific Histone Modification in Mammalian Spermatogenesis. <i>Evolutionary Bioinformatics</i> , 2015, 11s1, EBO.S21832.	0.6	10
76	Heuristic principal component analysis-based unsupervised feature extraction and its application to gene expression analysis of amyotrophic lateral sclerosis data sets. , 2015, , .		24
77	Exploring effective multiplicity in multichannel functional near-infrared spectroscopy using eigenvalues of correlation matrices. <i>Neurophotonics</i> , 2015, 2, 015002.	1.7	7
78	Principal component analysis-based unsupervised feature extraction applied to in silico drug discovery for posttraumatic stress disorder-mediated heart disease. <i>BMC Bioinformatics</i> , 2015, 16, 139.	1.2	46
79	Heuristic Principal Component Analysis-Based Unsupervised Feature Extraction and Its Application to Bioinformatics. <i>Advances in Bioinformatics and Biomedical Engineering Book Series</i> , 2015, , 138-162.	0.2	23
80	Universal disease biomarker: can a fixed set of blood microRNAs diagnose multiple diseases?. <i>BMC Research Notes</i> , 2014, 7, 581.	0.6	36
81	Bacterial Type III Secretion System Effector Proteins are Distinct between Plant Symbiotic, Plant Pathogenic and Animal Pathogenic Bacteria. <i>IPSI Transactions on Bioinformatics</i> , 2014, 7, 2-15.	0.2	1
82	TINAGL1 and B3GALNT1 are potential therapy target genes to suppress metastasis in non-small cell lung cancer. <i>BMC Genomics</i> , 2014, 15, S2.	1.2	66
83	Genes associated with genotype-specific DNA methylation in squamous cell carcinoma as candidate drug targets. <i>BMC Systems Biology</i> , 2014, 8, S4.	3.0	36
84	Integrative Analysis of Gene Expression and Promoter Methylation during Reprogramming of a Non-Small-Cell Lung Cancer Cell Line Using Principal Component Analysis-Based Unsupervised Feature Extraction. <i>Lecture Notes in Computer Science</i> , 2014, , 445-455.	1.0	20
85	Comparison of Hepatocellular Carcinoma miRNA Expression Profiling as Evaluated by Next Generation Sequencing and Microarray. <i>PLoS ONE</i> , 2014, 9, e106314.	1.1	74
86	MicroRNA-mediated regulation of target genes in several brain regions is correlated to both microRNA-targeting-specific promoter methylation and differential microRNA expression. <i>BioData Mining</i> , 2013, 6, 11.	2.2	15
87	Bioinformatic Screening of Autoimmune Disease Genes and Protein Structure Prediction with FAMS for Drug Discovery. <i>Protein and Peptide Letters</i> , 2013, 21, 828-839.	0.4	36
88	Principal Component Analysis Based Feature Extraction Approach to Identify Circulating microRNA Biomarkers. <i>PLoS ONE</i> , 2013, 8, e66714.	1.1	67
89	Possible miRNA Coregulation of Target Genes in Brain Regions by Both Differential miRNA Expression and miRNA-Targeting-Specific Promoter Methylation. <i>Communications in Computer and Information Science</i> , 2013, , 225-230.	0.4	0
90	MiRaGE: Inference of Gene Expression Regulation via MicroRNA Transfection II. <i>Lecture Notes in Computer Science</i> , 2012, , 129-135.	1.0	2

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91	Comprehensive miRNA Expression Analysis in Peripheral Blood Can Diagnose Liver Disease. PLoS ONE, 2012, 7, e48366.	1.1	149
92	Inference of Target Gene Regulation via miRNAs during Cell Senescence by Using the MiRaGE Server. Communications in Computer and Information Science, 2012, , 441-446.	0.4	7
93	Principal Component Analysis for Bacterial Proteomic Analysis. Lecture Notes in Computer Science, 2012, , 141-152.	1.0	16
94	Inference of Target Gene Regulation via miRNAs during Cell Senescence by Using the MiRaGE Server. , 2012, 3, 301-6.		6
95	The analysis of DNA shuffling by nMDS. , 2011, , .		0
96	Feature extraction for discriminance of symbiotic/parasitic bacterial type III effector protein using principal component analysis. , 2011, , .		0
97	Tissue specific methylation and genotype. , 2011, , .		0
98	Principal component analysis for bacterial proteomic analysis. , 2011, , .		4
99	Gene expression regulation during differentiation from murine ES cells due to microRNA. , 2011, , .		0
100	Protein binding prediction using non-metric multidimensional scaling method. , 2011, , .		1
101	Inference of Gene Regulation via miRNAs During ES Cell Differentiation Using MiRaGE Method. International Journal of Molecular Sciences, 2011, 12, 9265-9276.	1.8	14
102	Inference of Gene Expression Regulation via microRNA Transfection. Lecture Notes in Computer Science, 2010, , 672-679.	1.0	2
103	Gene Ontology term prediction based upon amino acid occurrence. , 2008, , .		0
104	Nonmetric Distances for Barcode of Life. IPSJ Transactions on Bioinformatics, 2008, 1, 35-41.	0.2	0
105	Application of amino acid occurrence for discriminating different folding types of globular proteins. BMC Bioinformatics, 2007, 8, 404.	1.2	58
106	Temperature measurement under convection and segregation in a vibrated bed of powder: A numerical study. Granular Matter, 2006, 8, 27-33.	1.1	0
107	Some implications of renormalization group theoretical ideas to statistics. Physica D: Nonlinear Phenomena, 2005, 205, 207-214.	1.3	3
108	Nonmetric Multidimensional Scaling As a Data-Mining Tool: New Algorithm and New Targets. Advances in Chemical Physics, 2005, , 315-351.	0.3	9

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109	Relational patterns of gene expression via non-metric multidimensional scaling analysis. <i>Bioinformatics</i> , 2005, 21, 730-740.	1.8	193
110	Why Do Physicists Study Barchan Dunes?. <i>JPSJ News and Comments</i> , 2005, 2, 02.	0.2	0
111	New Phases in Optimal Velocity Model. <i>IFAC Postprint Volumes IPPV / International Federation of Automatic Control</i> , 2003, 36, 395-399.	0.4	0
112	A Toy Model of Flying Snake's Glide. <i>Journal of the Physical Society of Japan</i> , 2003, 72, 3002-3005.	0.7	0
113	Verifying Relationship between Height and Spacing, in Barchan Dunes Simulated by the Coupled Map Lattice Model. <i>Journal of the Physical Society of Japan</i> , 2003, 72, 2685-2689.	0.7	3
114	Analysis of Segregation Property of Burden Using 2-Dimensional Discrete Model. <i>Tetsu-To-Hagane/Journal of the Iron and Steel Institute of Japan</i> , 2002, 88, 823-830.	0.1	16
115	Numerical investigation of surface level instability due to a tube in a vibrating bed of powder. <i>Physica A: Statistical Mechanics and Its Applications</i> , 1996, 232, 27-39.	1.2	12
116	FRACTAL LIMIT DISTRIBUTIONS IN RANDOM TRANSPORTS. <i>Fractals</i> , 1996, 04, 257-264.	1.8	2
117	Dynamics of Granular Matter from the Physical Point of View(I).. <i>Journal of the Society of Powder Technology, Japan</i> , 1995, 32, 240-246.	0.0	4
118	Dynamics of Granular Matter from the Physical Point of View. (III).. <i>Journal of the Society of Powder Technology, Japan</i> , 1995, 32, 412-418.	0.0	1
119	Dynamics of Granular Matter*1. <i>Japanese Journal of Applied Physics</i> , 1995, 34, 397-408.	0.8	30
120	Numerical study of granular turbulence and the appearance of the energy spectrum without flow. <i>Physica D: Nonlinear Phenomena</i> , 1995, 80, 61-71.	1.3	3
121	Power Law Velocity Fluctuations Due to Inelastic Collisions in Numerically Simulated Vibrated Bed of Powder. <i>Europhysics Letters</i> , 1995, 30, 499-504.	0.7	49
122	NUMERICAL MODELING OF MORPHOLOGICAL CHANGES OF THE FRACTURE PROPAGATION IN THE QUENCHED GLASS PLATE. <i>Modern Physics Letters B</i> , 1994, 08, 1335-1341.	1.0	4
123	NON-GAUSSIAN DISTRIBUTION IN RANDOM TRANSPORT DYNAMICS. <i>International Journal of Modern Physics B</i> , 1994, 08, 3887-3961.	1.0	13
124	Dynamical modelling of fracture propagation. <i>Materials Science &amp; Engineering A: Structural Materials: Properties, Microstructure and Processing</i> , 1994, 176, 295-298.	2.6	3
125	Numerical modeling of convective motion in granular materials. <i>Advanced Powder Technology</i> , 1994, 5, 297-303.	2.0	4
126	TURBULENT FLOW IN VIBRATED BED OF POWDER: NEW TARGET TO INVESTIGATE TURBULENT FLOW. , 1994, , 636-641.		0



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127	A new mesoscopic scale model for simulating fluid turbulence: the lattice vortex tube model. Physica D: Nonlinear Phenomena, 1993, 69, 366-379.	1.3	7
128	NUMERICAL MODELING OF VIBRATED BEDS. International Journal of Modern Physics B, 1993, 07, 1839-1858.	1.0	15
129	TURBULENT FLOW IN VIBRATED BED OF POWDER: NEW TARGET TO INVESTIGATE TURBULENT FLOW. Fractals, 1993, 01, 1080-1085.	1.8	4
130	$k^{-5/3}$ Power Spectrum in Powderâ€“Turbulent Flow in a Vibrated Bed: Numerical Results. Europhysics Letters, 1993, 24, 203-209.	0.7	28
131	Non-Gaussian distribution in random advection dynamics. Physical Review Letters, 1993, 70, 782-785.	2.9	13
132	Numerical Modeling of Convective Motion in Granular Materials.. Journal of the Society of Powder Technology, Japan, 1993, 30, 173-177.	0.0	12
133	New origin of a convective motion: Elastically induced convection in granular materials. Physical Review Letters, 1992, 69, 1367-1370.	2.9	231
134	Powder turbulence: direct onset of turbulent flow. Journal De Physique II, 1992, 2, 2103-2114.	0.9	67
135	Lorentzian distribution of interacting vortex tubes. Physical Review A, 1990, 41, 2249-2251.	1.0	5
136	Fracture propagation governed by the Laplace equation. Physica A: Statistical Mechanics and Its Applications, 1989, 156, 741-755.	1.2	15
137	Self-avoiding walks on Sierpinski carpets. Journal of Physics A, 1988, 21, 1929-1935.	1.6	8
138	Aggregation of particles which move on deterministic trajectories with fractal dimension two. I. A simple and new model for DLA. Journal of Physics A, 1988, 21, 4235-4240.	1.6	3
139	Noninteger-dimensional hyper-Euclidean lattices on Sierpinski carpets. Journal of Physics A, 1988, 21, 855-857.	1.6	4
140	The Correlation Function of the $\hat{A}_J$ Model on the Finite Square and Simple Cubic Lattices. Progress of Theoretical Physics, 1987, 77, 775-780.	2.0	4
141	Lacunarity and universality. Journal of Physics A, 1987, 20, 6611-6616.	1.6	21
142	Numerical Diagonalization of Quantum Spin Hamiltonians. Progress of Theoretical Physics Supplement, 1986, 87, 247-255.	0.2	16
143	Ground State of Antiferromagnetic Quantum Spin Systems on the Triangular Lattice. Journal of the Physical Society of Japan, 1986, 55, 323-330.	0.7	76
144	Ground State of Quantum Spin Glass with Infinite Range Interactions. Journal of the Physical Society of Japan, 1986, 55, 656-659.	0.7	5

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145	A Numerical Study of Spin-1/2 Alternating Antiferromagnetic Heisenberg Linear Chains. Journal of the Physical Society of Japan, 1986, 55, 1458-1465.	0.7	35
146	Ground-state properties of the Heisenberg antiferromagnet-numerical study. Journal of Magnetism and Magnetic Materials, 1986, 54-57, 1353-1354.	1.0	1
147	Transfer Matrix and Finite-Size Scaling for the Ising Model on Two- and Three-Dimensional Lattices. Progress of Theoretical Physics Supplement, 1986, 87, 23-32.	0.2	8
148	The Spin Wave Theory in Antiferromagnetic Heisenberg Model on Face Centered Cubic Lattice. Journal of the Physical Society of Japan, 1985, 54, 4494-4497.	0.7	39
149	Correlation between miRNA-targeted-gene promoter methylation and miRNA regulation of target genes. F1000Research, 0, 2, 21.	0.8	1
150	Correlation between miRNA-targeted-gene promoter methylation and miRNA regulation of target genes. F1000Research, 0, 2, 21.	0.8	2