

# Hsuan-Cheng Huang

## List of Publications by Year in descending order

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Version: 2024-02-01

164  
papers

9,442  
citations

41344

49  
h-index

42399

92  
g-index

168  
all docs

168  
docs citations

168  
times ranked

10762  
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeting protein interaction networks in mitochondrial dynamics for cancer therapy. <i>Drug Discovery Today</i> , 2022, 27, 1077-1087.	6.4	5
2	Drug Repurposing for the Identification of Compounds with Anti-SARS-CoV-2 Capability via Multiple Targets. <i>Pharmaceutics</i> , 2022, 14, 176.	4.5	6
3	Single-cell RNA sequencing uncovers the individual alteration of intestinal mucosal immunocytes in Dusp6 knockout mice. <i>IScience</i> , 2022, 25, 103738.	4.1	1
4	Quantitative phosphoproteomics reveals ectopic ATP synthase on mesenchymal stem cells to promote tumor progression via ERK/c-Fos pathway activation. <i>Molecular and Cellular Proteomics</i> , 2022, 21, 100237.	3.8	6
5	DockCoV2: a drug database against SARS-CoV-2. <i>Nucleic Acids Research</i> , 2021, 49, D1152-D1159.	14.5	42
6	Proteomic Analysis Reveals That Metformin Suppresses PSMD2, STIP1, and CAP1 for Preventing Gastric Cancer AGS Cell Proliferation and Migration. <i>ACS Omega</i> , 2021, 6, 14208-14219.	3.5	5
7	OUP accepted manuscript. <i>Bioinformatics</i> , 2021, , .	4.1	0
8	Identification of cell states using super-enhancer RNA. <i>BMC Genomics</i> , 2021, 22, 787.	2.8	0
9	Identification of a gut microbiota member that ameliorates DSS-induced colitis in intestinal barrier enhanced Dusp6-deficient mice. <i>Cell Reports</i> , 2021, 37, 110016.	6.4	35
10	Modular signature of long non-coding RNA association networks as a prognostic biomarker in lung cancer. <i>BMC Medical Genomics</i> , 2021, 14, 290.	1.5	4
11	Stratification of lncRNA modulation networks in breast cancer. <i>BMC Medical Genomics</i> , 2021, 14, 300.	1.5	1
12	Multimiomics Reveals Ectopic ATP Synthase Blockade Induces Cancer Cell Death via a lncRNA-mediated Phospho-signaling Network. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 1805-1825.	3.8	11
13	Enhancement of the IFN- $\gamma$ -induced host signature informs repurposed drugs for COVID-19. <i>Heliyon</i> , 2020, 6, e05646.	3.2	18
14	Phosphoproteome Analysis Reveals Dynamic Heat Shock Protein 27 Phosphorylation in Tanshinone IIA-Induced Cell Death. <i>Journal of Proteome Research</i> , 2020, 19, 1620-1634.	3.7	8
15	Quantitative Phosphoproteomics Reveals Cell Alignment and Mitochondrial Length Change under Cyclic Stretching in Lung Cells. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4074.	4.1	4
16	ZNF322A-mediated protein phosphorylation induces autophagosome formation through modulation of IRS1-AKT glucose uptake and HSP-elicited UPR in lung cancer. <i>Journal of Biomedical Science</i> , 2020, 27, 75.	7.0	9
17	RNA-Binding Proteomics Reveals MATR3 Interacting with lncRNA SNHG1 To Enhance Neuroblastoma Progression. <i>Journal of Proteome Research</i> , 2019, 18, 406-416.	3.7	21
18	Quantitative Proteomics of Th-MYCN Transgenic Mice Reveals Aurora Kinase Inhibitor Altered Metabolic Pathways and Enhanced ACADM To Suppress Neuroblastoma Progression. <i>Journal of Proteome Research</i> , 2019, 18, 3850-3866.	3.7	14

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19	Perturbational Gene-Expression Signatures for Combinatorial Drug Discovery. <i>IScience</i> , 2019, 15, 291-306.	4.1	27
20	Therapeutic Targeting of Non-oncogene Dependencies in High-risk Neuroblastoma. <i>Clinical Cancer Research</i> , 2019, 25, 4063-4078.	7.0	14
21	Investigating the role of super-enhancer RNAs underlying embryonic stem cell differentiation. <i>BMC Genomics</i> , 2019, 20, 896.	2.8	14
22	Introduction to Cancer Systems Biology. , 2018, , 1-9.		0
23	A Large-Scale Gene Expression Intensity-Based Similarity Metric for Drug Repositioning. <i>IScience</i> , 2018, 7, 40-52.	4.1	11
24	Dynamics of alternative polyadenylation in human preimplantation embryos. <i>Biochemical and Biophysical Research Communications</i> , 2018, 504, 727-733.	2.1	3
25	High-risk, Expression-Based Prognostic Long Noncoding RNA Signature in Neuroblastoma. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky015.	2.9	17
26	MCM2-regulated functional networks in lung cancer by multi-dimensional proteomic approach. <i>Scientific Reports</i> , 2017, 7, 13302.	3.3	37
27	DynaPho: a web platform for inferring the dynamics of time-series phosphoproteomics. <i>Bioinformatics</i> , 2017, 33, 3664-3666.	4.1	9
28	Identification of lncRNA functions in lung cancer based on associated protein-protein interaction modules. <i>Scientific Reports</i> , 2016, 6, 35939.	3.3	18
29	Oncoprotein ZNF322A transcriptionally deregulates alpha-adducin, cyclin D1 and p53 to promote tumor growth and metastasis in lung cancer. <i>Oncogene</i> , 2016, 35, 2357-2369.	5.9	35
30	Co-expression analysis identifies long noncoding RNA <i>SNHG1</i> as a novel predictor for event-free survival in neuroblastoma. <i>Oncotarget</i> , 2016, 7, 58022-58037.	1.8	59
31	Unveiling MYCN regulatory networks in neuroblastoma via integrative analysis of heterogeneous genomics data. <i>Oncotarget</i> , 2016, 7, 36293-36310.	1.8	37
32	Functional Analysis and Characterization of Differential Coexpression Networks. <i>Scientific Reports</i> , 2015, 5, 13295.	3.3	46
33	Temporal Phosphoproteome Dynamics Induced by an ATP Synthase Inhibitor Citreoviridin*. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 3284-3298.	3.8	23
34	Quantitative Proteomics Reveals Middle Infrared Radiation-Interfered Networks in Breast Cancer Cells. <i>Journal of Proteome Research</i> , 2015, 14, 1250-1262.	3.7	14
35	Integrating transcriptomics and proteomics to show that tanshinone IIA suppresses cell growth by blocking glucose metabolism in gastric cancer cells. <i>BMC Genomics</i> , 2015, 16, 41.	2.8	42
36	Circadian systems biology in Metazoa. <i>Briefings in Bioinformatics</i> , 2015, 16, 1008-1024.	6.5	3

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37	Integrating Phosphoproteomics and Bioinformatics to Study Brassinosteroid-Regulated Phosphorylation Dynamics in Arabidopsis. <i>BMC Genomics</i> , 2015, 16, 533.	2.8	52
38	Aryl Hydrocarbon Receptor Downregulates MYCN Expression and Promotes Cell Differentiation of Neuroblastoma. <i>PLoS ONE</i> , 2014, 9, e88795.	2.5	27
39	Combination therapy targeting ectopic ATP synthase and 26S proteasome induces ER stress in breast cancer cells. <i>Cell Death and Disease</i> , 2014, 5, e1540-e1540.	6.3	41
40	Deciphering molecular determinants of chemotherapy in gastrointestinal malignancy using systems biology approaches. <i>Drug Discovery Today</i> , 2014, 19, 1402-1409.	6.4	8
41	Mirin: identifying microRNA regulatory modules in protein-protein interaction networks. <i>Bioinformatics</i> , 2014, 30, 2527-2528.	4.1	6
42	Dissecting the Human Protein-Protein Interaction Network via Phylogenetic Decomposition. <i>Scientific Reports</i> , 2014, 4, 7153.	3.3	30
43	MicroRNA-mediated networks underlie immune response regulation in papillary thyroid carcinoma. <i>Scientific Reports</i> , 2014, 4, 6495.	3.3	25
44	Identification of MicroRNA 395a in 24-Epibrassinolide-Regulated Root Growth of <i>Arabidopsis thaliana</i> Using MicroRNA Arrays. <i>International Journal of Molecular Sciences</i> , 2013, 14, 14270-14286.	4.1	18
45	Quantitative Proteomics Reveals Diverse Roles of miR-148a from Gastric Cancer Progression to Neurological Development. <i>Journal of Proteome Research</i> , 2013, 12, 3993-4004.	3.7	20
46	Anatomical and transcriptional dynamics of maize embryonic leaves during seed germination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 3979-3984.	7.1	74
47	MicroRNA-Regulated Protein-Protein Interaction Networks and Their Functions in Breast Cancer. <i>International Journal of Molecular Sciences</i> , 2013, 14, 11560-11606.	4.1	56
48	Comprehensive methylome analysis of ovarian tumors reveals hedgehog signaling pathway regulators as prognostic DNA methylation biomarkers. <i>Epigenetics</i> , 2013, 8, 624-634.	2.7	51
49	ValidNESs: a database of validated leucine-rich nuclear export signals. <i>Nucleic Acids Research</i> , 2013, 41, D338-D343.	14.5	75
50	Middle Infrared Radiation Induces G2/M Cell Cycle Arrest in A549 Lung Cancer Cells. <i>PLoS ONE</i> , 2013, 8, e54117.	2.5	27
51	Differential MicroRNA Regulation Correlates with Alternative Polyadenylation Pattern between Breast Cancer and Normal Cells. <i>PLoS ONE</i> , 2013, 8, e56958.	2.5	29
52	Quantitative Proteomic Analysis of Human Lung Tumor Xenografts Treated with the Ectopic ATP Synthase Inhibitor Citreoviridin. <i>PLoS ONE</i> , 2013, 8, e70642.	2.5	26
53	Link Clustering Reveals Structural Characteristics and Biological Contexts in Signed Molecular Networks. <i>PLoS ONE</i> , 2013, 8, e67089.	2.5	7
54	Revealing the Anti-Tumor Effect of Artificial miRNA p-27-5p on Human Breast Carcinoma Cell Line T-47D. <i>International Journal of Molecular Sciences</i> , 2012, 13, 6352-6369.	4.1	5

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55	Revealing the Molecular Mechanism of Gastric Cancer Marker Annexin A4 in Cancer Cell Proliferation Using Exon Arrays. PLoS ONE, 2012, 7, e44615.	2.5	34
56	Ectopic ATP Synthase Blockade Suppresses Lung Adenocarcinoma Growth by Activating the Unfolded Protein Response. Cancer Research, 2012, 72, 4696-4706.	0.9	68
57	Profiling Lipid-protein Interactions Using Nonquenched Fluorescent Liposomal Nanovesicles and Proteome Microarrays. Molecular and Cellular Proteomics, 2012, 11, 1177-1190.	3.8	36
58	Methylomic Analysis Identifies Frequent DNA Methylation of Zinc Finger Protein 582 (ZNF582) in Cervical Neoplasms. PLoS ONE, 2012, 7, e41060.	2.5	72
59	Lengthening of 3'UTR increases with morphological complexity in animal evolution. Bioinformatics, 2012, 28, 3178-3181.	4.1	45
60	MicroRNA Regulation in Cellular Networks. , 2012, , 35-46.		0
61	Discovery of biomarkers for gastric cancer: A proteomics approach. Journal of Proteomics, 2012, 75, 3081-3097.	2.4	85
62	Crosstalk between transcription factors and microRNAs in human protein interaction network. BMC Systems Biology, 2012, 6, 18.	3.0	53
63	Helicobacter pylori Disrupts Host Cell Membranes, Initiating a Repair Response and Cell Proliferation. International Journal of Molecular Sciences, 2012, 13, 10176-10192.	4.1	20
64	Phosphoproteomic Analysis of <i>Rhodospseudomonas palustris</i> Reveals the Role of Pyruvate Phosphate Dikinase Phosphorylation in Lipid Production. Journal of Proteome Research, 2012, 11, 5362-5375.	3.7	37
65	Androgen pathway stimulates MicroRNA-216a transcription to suppress the tumor suppressor in lung cancer-1 gene in early hepatocarcinogenesis. Hepatology, 2012, 56, 632-643.	7.3	98
66	Corrigendum to "Silencing of miR-124 induces neuroblastoma SK-N-SH cell differentiation, cell cycle arrest and apoptosis through promoting AHR" [FEBS Lett. 585 (2011) 3582-3586]. FEBS Letters, 2012, 586, 107-107.	2.8	0
67	Identification of 14q32 in human gastric cancer cells and its potency as a diagnostic and prognostic biomarker. Proteomics, 2011, 11, 2423-2439.	2.2	38
68	Revealing the Functions of the Transketolase Enzyme Isoforms in <i>Rhodospseudomonas palustris</i> Using a Systems Biology Approach. PLoS ONE, 2011, 6, e28329.	2.5	10
69	Silencing of miR-124 induces neuroblastoma SK-N-SH cell differentiation, cell cycle arrest and apoptosis through promoting AHR. FEBS Letters, 2011, 585, 3582-3586.	2.8	67
70	Integrative network analysis reveals active microRNAs and their functions in gastric cancer. BMC Systems Biology, 2011, 5, 99.	3.0	78
71	Coregulation of transcription factors and microRNAs in human transcriptional regulatory network. BMC Bioinformatics, 2011, 12, S41.	2.6	84
72	The effect of narrow bandwidth infrared radiation on the growth of <i>Escherichia coli</i> . Applied Physics Letters, 2011, 99, 163704.	3.3	8

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73	Reviewing Ligand-Based Rational Drug Design: The Search for an ATP Synthase Inhibitor. <i>International Journal of Molecular Sciences</i> , 2011, 12, 5304-5318.	4.1	53
74	<i>Ganoderma lucidum</i> Polysaccharides Induce Macrophage-Like Differentiation in Human Leukemia THP-1 Cells via Caspase and p53 Activation. <i>Evidence-based Complementary and Alternative Medicine</i> , 2011, 2011, 1-13.	1.2	39
75	Prediction of Essential Genes by Mining Gene Ontology Semantics. <i>Lecture Notes in Computer Science</i> , 2011, , 49-60.	1.3	0
76	Modularity of <i>Escherichia coli</i> sRNA regulation revealed by sRNA-target and protein network analysis. <i>BMC Bioinformatics</i> , 2010, 11, S11.	2.6	8
77	Dynamic functional modules in co-expressed protein interaction networks of dilated cardiomyopathy. <i>BMC Systems Biology</i> , 2010, 4, 138.	3.0	45
78	Membrane repair against <i>H. pylori</i> promotes cancer cell proliferation. <i>Nature Precedings</i> , 2010, , .	0.1	0
79	Essential Core of Protein-Protein Interaction Network in <i>Escherichia coli</i> . <i>Journal of Proteome Research</i> , 2009, 8, 1925-1931.	3.7	30
80	Predicting essential genes based on network and sequence analysis. <i>Molecular BioSystems</i> , 2009, 5, 1672.	2.9	109
81	Characterization of microRNA-regulated protein-protein interaction network. <i>Proteomics</i> , 2008, 8, 1975-1979.	2.2	105
82	MelInfoText: associated gene methylation and cancer information from text mining. <i>BMC Bioinformatics</i> , 2008, 9, 22.	2.6	34
83	TCMGeneDIT: a database for associated traditional Chinese medicine, gene and disease information using text mining. <i>BMC Complementary and Alternative Medicine</i> , 2008, 8, 58.	3.7	107
84	Comparative analysis of differentially expressed genes in normal and white spot syndrome virus infected <i>Penaeus monodon</i> . <i>BMC Genomics</i> , 2007, 8, 120.	2.8	116
85	<i>Ganoderma lucidum</i> polysaccharides in human monocytic leukemia cells: from gene expression to network construction. <i>BMC Genomics</i> , 2007, 8, 411.	2.8	57
86	An apoptosis-related gene network induced by novel compound-RGD in human breast cancer cells. <i>FEBS Letters</i> , 2007, 581, 3517-3522.	2.8	11
87	Bioinformatics. <i>Methods in Molecular Biology</i> , 2007, 382, 405-416.	0.9	26
88	Quantitative Proteomic and Genomic Profiling Reveals Metastasis-Related Protein Expression Patterns in Gastric Cancer Cells. <i>Journal of Proteome Research</i> , 2006, 5, 2727-2742.	3.7	71
89	Proteomics analysis of a novel compound: Cyclic RGD in breast carcinoma cell line MCF-7. <i>Proteomics</i> , 2006, 6, 2991-3000.	2.2	20
90	Lipopolysaccharide-stimulated responses in rat aortic endothelial cells by a systems biology approach. <i>Proteomics</i> , 2006, 6, 5915-5928.	2.2	17

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91	An agent-based system to discover protein-protein interactions, identify protein complexes and proteins with multiple peptide mass fingerprints. <i>Journal of Computational Chemistry</i> , 2006, 27, 1020-1032.	3.3	3
92	Large-scale identification of protein-protein interaction of <i>Escherichia coli</i> K-12. <i>Genome Research</i> , 2006, 16, 686-691.	5.5	368
93	A new application of microwave technology to proteomics. <i>Proteomics</i> , 2005, 5, 840-842.	2.2	93
94	Measurements of branching fractions and CP asymmetries in $B^0 \rightarrow \pi^0 h$ decays. <i>Physical Review D</i> , 2005, 71, .	4.7	16
95	Evidence for Direct CP Violation in $B^0 \rightarrow K^* \pi^0$ Decays. <i>Physical Review Letters</i> , 2004, 93, 191802.	7.8	129
96	Measurement of the branching fractions for $B^0 \rightarrow \pi^0 K^0$ and $B^0 \rightarrow \pi^0 \bar{K}^0$ . <i>Physical Review D</i> , 2004, 70, .	4.7	5
97	Observation of $B^0 \rightarrow \pi^+ \pi^- K^0$ , $B^0 \rightarrow \pi^+ \pi^- \bar{K}^0$ , and $B^0 \rightarrow \pi^+ \pi^- K^*+$ . <i>Physical Review Letters</i> , 2004, 92, 131801.	7.8	68
98	Observation of Large CP Violation and Evidence for Direct CP Violation in $B^0 \rightarrow \pi^+ \pi^- \pi^0$ Decays. <i>Physical Review Letters</i> , 2004, 93, 021601.	7.8	67
99	Observation of Radiative $B^0 \rightarrow \pi^0 K^0$ Decays. <i>Physical Review Letters</i> , 2004, 92, 051801.	7.8	12
100	Measurements of the $D_{s1}$ Resonance Properties. <i>Physical Review Letters</i> , 2004, 92, 012002.	7.8	127
101	Measurement of the $e^+ e^- \rightarrow D^{(*)} + D^{(*)}$ cross sections. <i>Physical Review D</i> , 2004, 70, .	4.7	5
102	Study of $B^0 \rightarrow D^{*0} \pi^0$ ( $D^{*0} \rightarrow D^{*0} + \pi^0$ ) decays. <i>Physical Review D</i> , 2004, 69, .	4.7	221
103	Observation of the Radiative Decay $D^0 \rightarrow \pi^0 \pi^0$ . <i>Physical Review Letters</i> , 2004, 92, 101803.	7.8	11
104	Observation of $B^0 \rightarrow \pi^0 (3770) K^+$ . <i>Physical Review Letters</i> , 2004, 93, 051803.	7.8	40
105	Improved measurements of the branching fractions for $B^0 \rightarrow K^0 \pi^0$ , $B^0 \rightarrow K^0 \bar{K}^0$ decays. <i>Physical Review D</i> , 2004, 69, .	4.7	40
106	Evidence for $B^0 \rightarrow \pi^0 \pi^0$ . <i>Physical Review Letters</i> , 2004, 93, 131803.	7.8	17
107	Search for $B^0 \rightarrow \pi^0 \pi^0$ decay. <i>Physical Review D</i> , 2004, 69, .	4.7	4
108	Measurement of $ V_{ub} $ Using Inclusive $B^0 \rightarrow Xu$ , $\pi^0$ Decays with a Novel Xu-Reconstruction Method. <i>Physical Review Letters</i> , 2004, 92, 101801.	7.8	32

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109	Upper Bound on the Decay $B^0 \rightarrow \pi^+ \pi^- \pi^0$ from the Belle Detector. Physical Review Letters, 2004, 92, 171802.	7.8	88
110	Search for the Lepton-Flavor-Violating Decay $B^0 \rightarrow \tau^+ \tau^- \pi^0$ at Belle. Physical Review Letters, 2004, 93, 081803.	7.8	13
111	Study of B meson decays to three-body charmless hadronic final states. Physical Review D, 2004, 69, .	4.7	55
112	GeneNetwork: an interactive tool for reconstruction of genetic networks using microarray data. Bioinformatics, 2004, 20, 3691-3693.	4.1	57
113	ProteMiner-SSM: a web server for efficient analysis of similar protein tertiary substructures. Nucleic Acids Research, 2004, 32, W76-W82.	14.5	19
114	Small molecules targeting severe acute respiratory syndrome human coronavirus. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10012-10017.	7.1	458
115	Baryonic B decays at Belle. European Physical Journal C, 2004, 33, s216-s218.	3.9	2
116	Observation of the decays $B^0 \rightarrow \pi^+ \pi^- \pi^0$ . <small>xmlns:xocs="http://www.elsevier.com/xml/xocs/dtd" xmlns:xs="http://www.w3.org/2001/XMLSchema" xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance" xmlns="http://www.elsevier.com/xml/ja/dtd" xmlns:ja="http://www.elsevier.com/xml/ja/dtd" xmlns:mml="http://www.w3.org/1998/Math/MathML" xmlns:tb="http://www.elsevier.com/xml/common/table/dtd" xmlns:sb="http://www.elsevier.com/xml/common/struct-bib/dtd" xmlns:ce="http://www.elsevier.com/x</small>	4.1	28
117	Measurement of $K^+ K^-$ production in two-photon collisions in the resonant-mass region. European Physical Journal C, 2003, 32, 323-336.	3.9	36
118	Study of $B^0 \rightarrow D^+ D^- \pi^0$ decays. Physics Letters, Section B: Nuclear, Elementary Particle and High-Energy Physics, 2003, 553, 159-166.	4.1	24
119	First measurement of the T-violating muon polarization in the decay $K^+ \rightarrow \pi^+ \mu^+ \mu^-$ . Physics Letters, Section B: Nuclear, Elementary Particle and High-Energy Physics, 2003, 562, 166-172.	4.1	8
120	Observation of the Decay $B^0 \rightarrow \pi^+ \pi^- \pi^0$ . Physical Review Letters, 2003, 90, 121802.	7.8	45
121	Observation of the $D_{s1}(2317)$ and $D_{s1}(2457)$ in B Decays. Physical Review Letters, 2003, 91, 262002.	7.8	229
122	Evidence for $B^0 \rightarrow \pi^+ \pi^- K^0$ . Physical Review Letters, 2003, 91, 241802.	7.8	18
123	Measurement of Time-Dependent CP-Violating Asymmetries in $B^0 \rightarrow \pi^+ \pi^- K_S^0$ , $K^+ K^- K_S^0$ , and $\pi^+ \pi^- K_S^0$ Decays. Physical Review Letters, 2003, 91, 261602.	7.8	82
124	Study of time-dependent CP-violating asymmetries in $B^0 \rightarrow \pi^+ \pi^- \pi^0$ decays. Physical Review D, 2003, 67, .	4.7	27
125	Observation of $B^0 \rightarrow D^+ D^- K^0$ and $B^0 \rightarrow D^+ D^- K^*$ Decays. Physical Review Letters, 2003, 90, 141802.	7.8	28
126	Measurement of branching fractions and charge asymmetries for two-body B meson decays with charmonium. Physical Review D, 2003, 67, .	4.7	34



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127	Measurement of Branching Fractions and Polarization in $B \rightarrow \tau K^*$ Decays. Physical Review Letters, 2003, 91, 201801.	7.8	109
128	Evidence for $B \rightarrow \tau \bar{c} \bar{c} \bar{c}$ . Physical Review Letters, 2003, 91, 261801.	7.8	140
129	Evidence for CP-violating asymmetries in $B \rightarrow \tau \bar{c} \bar{c} \bar{c}$ decays and constraints on the CKM angle $\beta_2$ . Physical Review D, 2003, 68, .	4.7	41
130	Observation of a Narrow Charmoniumlike State in Exclusive $B \rightarrow \tau K^* \bar{c} \bar{c} \bar{c} / \bar{c} \bar{c} \bar{c}$ Decays. Physical Review Letters, 2003, 91, 262001.	7.8	1,526
131	Studies of $B \rightarrow B$ mixing properties with inclusive dilepton events. Physical Review D, 2003, 67, .	4.7	30
132	Measurement of the Electroweak Penguin Process $B \rightarrow X s \bar{c} \bar{c} \bar{c}$ . Physical Review Letters, 2003, 90, 021801.	7.8	57
133	Measurement of Branching Fractions for $B \rightarrow \tau c \bar{c} \bar{c}$ Decays. Physical Review Letters, 2003, 90, 071801.	7.8	29
134	Measurement of branching fraction ratios and CP asymmetries in $B \rightarrow \tau D C P K^* \bar{c}$ . Physical Review D, 2003, 68, .	4.7	19
135	Observation of $B \rightarrow \tau \bar{c} \bar{c} \bar{c}$ Decays. Physical Review Letters, 2003, 91, 221801.	7.8	90
136	Observation of $B \rightarrow \tau \bar{c} \bar{c} \bar{c}$ . Physical Review Letters, 2003, 90, 201802.	7.8	61
137	Studies of the Decay $B \rightarrow \tau D C P K^* \bar{c}$ . Physical Review Letters, 2003, 90, 131803.	7.8	8
138	Improved measurement of the partial-rate CP asymmetry in $B \rightarrow \tau K^* \bar{c} \bar{c}$ and $B \rightarrow \tau K^* \bar{c} \bar{c}$ decays. Physical Review D, 2003, 68, .	4.7	11
139	Observation of $B \rightarrow \tau K^* \bar{c} \bar{c}$ . Physical Review Letters, 2003, 91, 261601.	7.8	79
140	Search for $B \rightarrow \tau l \bar{l}$ at the Belle detector. Physical Review D, 2003, 68, .	4.7	28
141	Measurement of the $B \rightarrow B$ mixing rate with $B_0(B_0) \rightarrow D^* \bar{c} \bar{c}$ partial reconstruction. Physical Review D, 2003, 67, .	4.7	11
142	Observation of $B \rightarrow 0 \tau^+ D^*$ Opposite. Physical Review Letters, 2002, 89, 151802.	7.8	90
143	Observation of $D_s + K^*$ and Evidence for $D_s + \bar{c} \bar{c}$ Final States in Neutral B Decays. Physical Review Letters, 2002, 89, 231804.	7.8	40
144	Observation of the $\bar{c}(2S)$ in Exclusive $B \rightarrow \tau K^* K^* \bar{c} \bar{c}$ Decays. Physical Review Letters, 2002, 89, 102001.	7.8	141

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145	Study of CP-Violating Asymmetries in $B^0 \rightarrow \pi^+ \pi^-$ Decays. Physical Review Letters, 2002, 89, 071801.	7.8	21
146	Observation of $\chi_c^2$ Production in B Meson Decay. Physical Review Letters, 2002, 89, 011803.	7.8	12
147	Radiative B Meson Decays into $K_S^0$ and $K_L^0$ Final States. Physical Review Letters, 2002, 89, 231801.	7.8	43
148	Charmless hadronic two-body B meson decays. Physical Review D, 2002, 66, .	4.7	23
149	Search for charmless two-body baryonic decays of B mesons. Physical Review D, 2002, 65, .	4.7	27
150	Measurement of the $B^0 \rightarrow B^0$ Mixing Parameter $\eta'$ using Semileptonic B Decays. Physical Review Letters, 2002, 89, 251803.	7.8	13
151	Observation of mixing-induced CP violation in the neutral B meson system. Physical Review D, 2002, 66, .	4.7	48
152	Improved measurement of mixing-induced CP violation in the neutral B meson system. Physical Review D, 2002, 66, .	4.7	177
153	Observation of $B^0 \rightarrow \pi^+ \pi^- K^0$ . Physical Review Letters, 2002, 88, 181803.	7.8	105
154	Observation of $B^0 \rightarrow \pi^0 K^0$ Decay. Physical Review Letters, 2002, 89, 191801.	7.8	12
155	Study of three-body charmless B decays. Physical Review D, 2002, 65, .	4.7	73
156	Observation of Double $c\bar{c}$ Production in $e^+e^-$ Annihilation at $\sqrt{s} \approx 10.6$ GeV. Physical Review Letters, 2002, 89, 142001.	7.8	268
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