

Yan Guo

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/120818/publications.pdf>

Version: 2024-02-01

256
papers

9,733
citations

36303

51
h-index

56724

83
g-index

266
all docs

266
docs citations

266
times ranked

17266
citing authors

#	ARTICLE	IF	CITATIONS
1	Randomized Trial of First-Line Tyrosine Kinase Inhibitor With or Without Radiotherapy for Synchronous Oligometastatic EGFR-Mutated Non-Small Cell Lung Cancer. <i>Journal of the National Cancer Institute</i> , 2023, 115, 742-748.	6.3	99
2	Comparison of femoral neck system to multiple cancellous screws and dynamic hip screws in the treatment of femoral neck fractures. <i>Injury</i> , 2023, 54, S28-S35.	1.7	7
3	Two-stage-vote ensemble framework based on integration of mutation data and gene interaction network for uncovering driver genes. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	5
4	Non-autophagy Role of Atg5 and NBR1 in Unconventional Secretion of IL-12 Prevents Gut Dysbiosis and Inflammation. <i>Journal of Crohn's and Colitis</i> , 2022, 16, 259-274.	1.3	10
5	Cell cycle exit and neuronal differentiation 1-engineered embryonic neural stem cells enhance neuronal differentiation and neurobehavioral recovery after experimental traumatic brain injury. <i>Neural Regeneration Research</i> , 2022, 17, 130.	3.0	8
6	SBSA: an online service for somatic binding sequence annotation. <i>Nucleic Acids Research</i> , 2022, 50, e4-e4.	14.5	8
7	1-Hydroxypyrene mediates renal fibrosis through aryl hydrocarbon receptor signalling pathway. <i>British Journal of Pharmacology</i> , 2022, 179, 103-124.	5.4	28
8	Characterization and proteomics of chicken seminal plasma extracellular vesicles. <i>Reproduction in Domestic Animals</i> , 2022, 57, 98-110.	1.4	4
9	Comprehensive Analysis of Co-Mutations Identifies Cooperating Mechanisms of Tumorigenesis. <i>Cancers</i> , 2022, 14, 415.	3.7	8
10	Metallo- β -lactamases inhibitor fisetin attenuates meropenem resistance in NDM-1-producing <i>Escherichia coli</i> . <i>European Journal of Medicinal Chemistry</i> , 2022, 231, 114108.	5.5	8
11	FOXL2 regulates the expression of the Col4a1 collagen gene in chicken granulosa cells. <i>Molecular Reproduction and Development</i> , 2022, 89, 95-103.	2.0	5
12	Facile preparation of a cellulose derived carbon/BN composite aerogel for superior electromagnetic wave absorption. <i>Journal of Materials Chemistry C</i> , 2022, 10, 5311-5320.	5.5	20
13	GTQC: Automated Genotyping Array Quality Control and Report. <i>Journal of Genomics</i> , 2022, 10, 39-44.	0.9	1
14	Is it time to reassess variant annotation?. <i>Trends in Genetics</i> , 2022, 38, 521-523.	6.7	5
15	Serum Metabolites Associated with Blood Pressure in Chronic Kidney Disease Patients. <i>Metabolites</i> , 2022, 12, 281.	2.9	1
16	Hierarchical HCF@NC/Co Derived from Hollow Loofah Fiber Anchored with Metal-Organic Frameworks for Highly Efficient Microwave Absorption. <i>ACS Applied Materials & Interfaces</i> , 2022, 14, 2038-2050.	8.0	44
17	A Novel Strategy to Identify Prognosis-Relevant Gene Sets in Cancers. <i>Genes</i> , 2022, 13, 862.	2.4	0
18	Intrarenal 1-methoxypyrene, an aryl hydrocarbon receptor agonist, mediates progressive tubulointerstitial fibrosis in mice. <i>Acta Pharmacologica Sinica</i> , 2022, 43, 2929-2945.	6.1	23

#	ARTICLE	IF	CITATIONS
19	Engineering hierarchical heterostructure material based on metal-organic frameworks and cotton fiber for high-efficient microwave absorber. <i>Nano Research</i> , 2022, 15, 6841-6850.	10.4	59
20	Prediction of Major Histocompatibility Complex Binding with Bilateral and Variable Long Short Term Memory Networks. <i>Biology</i> , 2022, 11, 848.	2.8	1
21	Drug-Target Network Study Reveals the Core Target-Protein Interactions of Various COVID-19 Treatments. <i>Genes</i> , 2022, 13, 1210.	2.4	0
22	Resistance-Confering Mutations on Whole-Genome Sequencing of Fluoroquinolone-resistant and -Susceptible <i>Mycobacterium tuberculosis</i> Isolates: A Proposed Threshold for Identifying Resistance. <i>Clinical Infectious Diseases</i> , 2021, 72, 1910-1918.	5.8	6
23	Deciphering the cellular mechanisms underlying fibrosis-associated diseases and therapeutic avenues. <i>Pharmacological Research</i> , 2021, 163, 105316.	7.1	32
24	A resilient and lightweight bacterial cellulose-derived C/rGO aerogel-based electromagnetic wave absorber integrated with multiple functions. <i>Journal of Materials Chemistry A</i> , 2021, 9, 5566-5577.	10.3	62
25	<i>Polyporus Umbellatus</i> Protects Against Renal Fibrosis by Regulating Intrarenal Fatty Acyl Metabolites. <i>Frontiers in Pharmacology</i> , 2021, 12, 633566.	3.5	8
26	The Dysregulation of Eicosanoids and Bile Acids Correlates with Impaired Kidney Function and Renal Fibrosis in Chronic Renal Failure. <i>Metabolites</i> , 2021, 11, 127.	2.9	10
27	A streamlined solution for processing, elucidating and quality control of cyclobutane pyrimidine dimer sequencing data. <i>Nature Protocols</i> , 2021, 16, 2190-2212.	12.0	3
28	Rewired Pathways and Disrupted Pathway Crosstalk in Schizophrenia Transcriptomes by Multiple Differential Coexpression Methods. <i>Genes</i> , 2021, 12, 665.	2.4	7
29	MetaGSCA: A tool for meta-analysis of gene set differential coexpression. <i>PLoS Computational Biology</i> , 2021, 17, e1008976.	3.2	2
30	A novel maize biochar-based compound fertilizer for immobilizing cadmium and improving soil quality and maize growth. <i>Environmental Pollution</i> , 2021, 277, 116455.	7.5	35
31	Predicting MHC class I binder: existing approaches and a novel recurrent neural network solution. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	6
32	Approaches to Potentiated Neuroprotective Treatment in the Rodent Model of Ischemic Optic Neuropathy. <i>Cells</i> , 2021, 10, 1440.	4.1	7
33	Electrostatic self-assembled NiFe ₂ O ₄ /Ti ₃ C ₂ T _x MXene nanocomposites for efficient electromagnetic wave absorption at ultralow loading level. <i>Advanced Composites and Hybrid Materials</i> , 2021, 4, 602-613.	21.1	97
34	Detecting SARS-CoV-2 and its variant strains with a full genome tiling array. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	4
35	5-HT ₃ Signaling Alters Development of Sacral Neural Crest Derivatives That Innervate the Lower Urinary Tract. <i>International Journal of Molecular Sciences</i> , 2021, 22, 6838.	4.1	5
36	A systematic view of computational methods for identifying driver genes based on somatic mutation data. <i>Briefings in Functional Genomics</i> , 2021, 20, 333-343.	2.7	4

#	ARTICLE	IF	CITATIONS
37	Risk factors for implant failure of intertrochanteric fractures with lateral femoral wall fracture after intramedullary nail fixation. <i>Injury</i> , 2021, 52, 3397-3403.	1.7	13
38	RNA editing affects cis-regulatory elements and predicts adverse cancer survival. <i>Cancer Medicine</i> , 2021, 10, 6114-6127.	2.8	5
39	Tuberculosis infection screening in children with close contact: a hospital-based study. <i>BMC Infectious Diseases</i> , 2021, 21, 815.	2.9	4
40	miR-378-3p Knockdown Recapitulates Many of the Features of Myelodysplastic Syndromes. <i>American Journal of Pathology</i> , 2021, 191, 2009-2022.	3.8	1
41	Longitudinal Assessment of Cytokine Expression and Plasminogen Activation in Hantavirus Cardiopulmonary Syndrome Reveals Immune Regulatory Dysfunction in End-Stage Disease. <i>Viruses</i> , 2021, 13, 1597.	3.3	4
42	Tracing Transmission of Sin Nombre Virus and Discovery of Infection in Multiple Rodent Species. <i>Journal of Virology</i> , 2021, 95, e0153421.	3.4	14
43	Heterogeneity of Circulating Tumor Cell Neoplastic Subpopulations Outlined by Single-Cell Transcriptomics. <i>Cancers</i> , 2021, 13, 4885.	3.7	17
44	Sirt3 Maintains Microvascular Endothelial Adherens Junction Integrity to Alleviate Sepsis-Induced Lung Inflammation by Modulating the Interaction of VE-Cadherin and β -Catenin. <i>Oxidative Medicine and Cellular Longevity</i> , 2021, 2021, 1-18.	4.0	8
45	EditPredict: Prediction of RNA editable sites with convolutional neural network. <i>Genomics</i> , 2021, 113, 3864-3871.	2.9	6
46	Transcriptional regulation of CYP19A1 expression in chickens: ESR1, ESR2 and NR5A2 form a functional network. <i>General and Comparative Endocrinology</i> , 2021, 315, 113939.	1.8	9
47	AMPK-Mediated Metabolic Switching Is High Effective for Phytochemical Levo-Tetrahydropalmatine (l-THP) to Reduce Hepatocellular Carcinoma Tumor Growth. <i>Metabolites</i> , 2021, 11, 811.	2.9	9
48	Conditional transcriptional relationships may serve as cancer prognostic markers. <i>BMC Medical Genomics</i> , 2021, 14, 101.	1.5	5
49	FeCo alloy nanoparticle decorated cellulose based carbon aerogel as a low-cost and efficient electromagnetic microwave absorber. <i>Journal of Materials Chemistry C</i> , 2021, 10, 126-134.	5.5	30
50	Cancer-specific expression quantitative loci are affected by expression dysregulation. <i>Briefings in Bioinformatics</i> , 2020, 21, 338-347.	6.5	4
51	Development and Validation of a MicroRNA Panel to Differentiate Between Patients with Rheumatoid Arthritis or Systemic Lupus Erythematosus and Controls. <i>Journal of Rheumatology</i> , 2020, 47, 188-196.	2.0	33
52	Small molecule inhibitors of epithelial-mesenchymal transition for the treatment of cancer and fibrosis. <i>Medicinal Research Reviews</i> , 2020, 40, 54-78.	10.5	93
53	MutEx: a multifaceted gateway for exploring integrative pan-cancer genomic data. <i>Briefings in Bioinformatics</i> , 2020, 21, 1479-1486.	6.5	12
54	Advancing Pan-cancer Gene Expression Survival Analysis by Inclusion of Non-coding RNA. <i>RNA Biology</i> , 2020, 17, 1666-1673.	3.1	26

#	ARTICLE	IF	CITATIONS
55	The Endogenous Plasma Small RNAome of Rheumatoid Arthritis. <i>ACR Open Rheumatology</i> , 2020, 2, 97-105.	2.1	16
56	Design, synthesis, <i>in vitro</i> and <i>in vivo</i> evaluation of benzylpiperidine-linked 1,3-dimethylbenzimidazolinones as cholinesterase inhibitors against Alzheimer's disease. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2020, 35, 330-343.	5.2	19
57	Allyl isothiocyanate upregulates MRP1 expression through Notch1 signaling in human bronchial epithelial cells. <i>Canadian Journal of Physiology and Pharmacology</i> , 2020, 98, 324-331.	1.4	4
58	Machine learning distilled metabolite biomarkers for early stage renal injury. <i>Metabolomics</i> , 2020, 16, 4.	3.0	18
59	Expression correlation attenuates within and between key signaling pathways in chronic kidney disease. <i>BMC Medical Genomics</i> , 2020, 13, 134.	1.5	9
60	Management of medically inoperable and tyrosine kinase inhibitor-naïve early-stage lung adenocarcinoma with epidermal growth factor receptor mutations: a retrospective multi-institutional analysis. <i>BMC Cancer</i> , 2020, 20, 646.	2.6	3
61	A Method of Ultra-Large-Scale Matrix Inversion Using Block Recursion. <i>Information (Switzerland)</i> , 2020, 11, 523.	2.9	5
62	Alternative Applications of Genotyping Array Data Using Multivariate Methods. <i>Trends in Genetics</i> , 2020, 36, 857-867.	6.7	7
63	Characterization of Corosolic Acid as a KPC-2 Inhibitor That Increases the Susceptibility of KPC-2-Positive Bacteria to Carbapenems. <i>Frontiers in Pharmacology</i> , 2020, 11, 1047.	3.5	7
64	Discovery of a Novel Natural Allosteric Inhibitor That Targets NDM-1 Against <i>Escherichia coli</i> . <i>Frontiers in Pharmacology</i> , 2020, 11, 581001.	3.5	10
65	SMDB: pivotal somatic sequence alterations reprogramming regulatory cascades. <i>NAR Cancer</i> , 2020, 2, zcaa030.	3.1	8
66	Global Autozygosity Is Associated with Cancer Risk, Mutational Signature and Prognosis. <i>Cancers</i> , 2020, 12, 3646.	3.7	1
67	Application of Oleanolic Acid and Its Analogues in Combating Pathogenic Bacteria <i>in Vitro</i> / <i>in Vivo</i> by a Two-Pronged Strategy of β -Lactamases and Hemolysins. <i>ACS Omega</i> , 2020, 5, 11424-11438.	3.5	8
68	LDB1 Enforces Stability on Direct and Indirect Oncoprotein Partners in Leukemia. <i>Molecular and Cellular Biology</i> , 2020, 40, .	2.3	11
69	Non-canonical RNA-DNA differences and other human genomic features are enriched within very short tandem repeats. <i>PLoS Computational Biology</i> , 2020, 16, e1007968.	3.2	4
70	Identification of endogenous ϵ -aminopyrene as a novel mediator of progressive chronic kidney disease via aryl hydrocarbon receptor activation. <i>British Journal of Pharmacology</i> , 2020, 177, 3415-3435.	5.4	50
71	Achieving enhanced electromagnetic shielding and absorption capacity of cellulose-derived carbon aerogels <i>in vitro</i> tuning the carbonization temperature. <i>Journal of Materials Chemistry C</i> , 2020, 8, 5191-5201.	5.5	51
72	A Silylene-Germylene Molecule Containing a Si ⁺ Ge ⁻ Single Bond. <i>Chemistry - A European Journal</i> , 2020, 26, 6122-6125.	3.3	7

#	ARTICLE	IF	CITATIONS
73	AnnoGen: annotating genome-wide pragmatic features. <i>Bioinformatics</i> , 2020, 36, 2899-2901.	4.1	4
74	Hyperactivation of TORC1 Drives Resistance to the Pan-HER Tyrosine Kinase Inhibitor Neratinib in HER2-Mutant Cancers. <i>Cancer Cell</i> , 2020, 37, 183-199.e5.	16.8	33
75	Isoalantolactone restores the sensitivity of gram-negative <i>Enterobacteriaceae</i> carrying MCR-1 to carbapenems. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 2475-2483.	3.6	14
76	Design, Synthesis, and Evaluation of Acetylcholinesterase and Butyrylcholinesterase Dual-Target Inhibitors against Alzheimer's Diseases. <i>Molecules</i> , 2020, 25, 489.	3.8	12
77	Appraising the role of circulating concentrations of micro-nutrients in epithelial ovarian cancer risk: A Mendelian randomization analysis. <i>Scientific Reports</i> , 2020, 10, 7356.	3.3	21
78	The International Conference on Intelligent Biology and Medicine 2019 (ICIBM 2019): computational methods and applications in medical genomics. <i>BMC Medical Genomics</i> , 2020, 13, 47.	1.5	1
79	Innovating Computational Biology and Intelligent Medicine: ICIBM 2019 Special Issue. <i>Genes</i> , 2020, 11, 437.	2.4	0
80	NFATc3 regulation of collagen V expression contributes to cellular immunity to collagen type V and hypoxic pulmonary hypertension. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2020, 319, L968-L980.	2.9	6
81	Prognostic value and co-expression patterns of metabolic pathways in cancers. <i>BMC Genomics</i> , 2020, 21, 860.	2.8	5
82	Accelerating bioinformatics research with International Conference on Intelligent Biology and Medicine 2020. <i>BMC Bioinformatics</i> , 2020, 21, 563.	2.6	2
83	AITC induces MRP1 expression by protecting against CS/CSE-mediated DJ-1 protein degradation via activation of the DJ-1/Nrf2 axis. <i>Korean Journal of Physiology and Pharmacology</i> , 2020, 24, 481-492.	1.2	2
84	Quality and concordance of genotyping array data of 12,064 samples from 5840 cancer patients. <i>Genomics</i> , 2019, 111, 950-957.	2.9	7
85	Combined melatonin and poricoic acid A inhibits renal fibrosis through modulating the interaction of Smad3 and β -catenin pathway in AKI-to-CKD continuum. <i>Therapeutic Advances in Chronic Disease</i> , 2019, 10, 204062231986911.	2.5	38
86	Natural products against renin-angiotensin system for antifibrosis therapy. <i>European Journal of Medicinal Chemistry</i> , 2019, 179, 623-633.	5.5	30
87	Aryl hydrocarbon receptor activation mediates kidney disease and renal cell carcinoma. <i>Journal of Translational Medicine</i> , 2019, 17, 302.	4.4	42
88	Insight into the Etiology of Undifferentiated Soft Tissue Sarcomas from a Novel Mouse Model. <i>Molecular Cancer Research</i> , 2019, 17, 1024-1035.	3.4	8
89	Two-week administration of engineered <i>Escherichia coli</i> establishes persistent resistance to diet-induced obesity even without antibiotic pre-treatment. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 6711-6723.	3.6	10
90	Microbiome metabolomics reveals gut microbiota associated with glycine-conjugated metabolites and polyamine metabolism in chronic kidney disease. <i>Cellular and Molecular Life Sciences</i> , 2019, 76, 4961-4978.	5.4	146

#	ARTICLE	IF	CITATIONS
91	Activated NF- κ B/Nrf2 and Wnt/ β -catenin pathways are associated with lipid metabolism in CKD patients with microalbuminuria and macroalbuminuria. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2019, 1865, 2317-2332.	3.8	42
92	Architectures and accuracy of artificial neural network for disease classification from omics data. <i>BMC Genomics</i> , 2019, 20, 167.	2.8	58
93	Discovery of a potential MCR-1 inhibitor that reverses polymyxin activity against clinical mcr-1-positive Enterobacteriaceae. <i>Journal of Infection</i> , 2019, 78, 364-372.	3.3	51
94	Aberrant FGFR signaling mediates resistance to CDK4/6 inhibitors in ER+ breast cancer. <i>Nature Communications</i> , 2019, 10, 1373.	12.8	252
95	Preparation and <i>in vitro/in vivo</i> evaluation of azilsartan osmotic pump tablets based on the preformulation investigation. <i>Drug Development and Industrial Pharmacy</i> , 2019, 45, 1079-1088.	2.0	8
96	Synthesis, Characterization, and Anticancer Activities Evaluation of Compounds Derived from 3,4-Dihydropyrimidin-2(1H)-one. <i>Molecules</i> , 2019, 24, 891.	3.8	22
97	Genomic Positional Dissection of RNA Editomes in Tumor and Normal Samples. <i>Frontiers in Genetics</i> , 2019, 10, 211.	2.3	19
98	Unilateral ureteral obstruction causes gut microbial dysbiosis and metabolome disorders contributing to tubulointerstitial fibrosis. <i>Experimental and Molecular Medicine</i> , 2019, 51, 1-18.	7.7	90
99	Identification of serum metabolites associating with chronic kidney disease progression and anti-fibrotic effect of 5-methoxytryptophan. <i>Nature Communications</i> , 2019, 10, 1476.	12.8	171
100	Role of Bile Acids and GLP-1 in Mediating the Metabolic Improvements of Bariatric Surgery. <i>Gastroenterology</i> , 2019, 156, 1041-1051.e4.	1.3	118
101	Integrated Analysis of Mouse and Human Gastric Neoplasms Identifies Conserved microRNA Networks in Gastric Carcinogenesis. <i>Gastroenterology</i> , 2019, 156, 1127-1139.e8.	1.3	35
102	Peripheral Blood Mitochondrial DNA Copy Number Obtained From Genome-Wide Genotype Data Is Associated With Neurocognitive Impairment in Persons With Chronic HIV Infection. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2019, 80, e95-e102.	2.1	16
103	Microbiome-metabolome reveals the contribution of gut-kidney axis on kidney disease. <i>Journal of Translational Medicine</i> , 2019, 17, 5.	4.4	233
104	Pharmacokinetics, Tissue Distribution, Plasma Protein Binding Studies of 10-Dehydroxyl-12-Demethoxy-Conophylline, a Novel Anti-Tumor Candidate, in Rats. <i>Molecules</i> , 2019, 24, 283.	3.8	5
105	Four new diarylheptanoids from <i>Rhizoma Zingiberis</i> . <i>Journal of Asian Natural Products Research</i> , 2019, 21, 1-8.	1.4	6
106	Single-nucleotide variants in human RNA: RNA editing and beyond. <i>Briefings in Functional Genomics</i> , 2019, 18, 30-39.	2.7	17
107	The Effects of Different Catalysts, Substituted Aromatic Aldehydes on One-Pot Three-Component Biginelli Reaction. <i>Current Organic Synthesis</i> , 2019, 16, 181-186.	1.3	0
108	Novel inhibitors of the cellular renin-angiotensin system components, poricoic acids, target Smad3 phosphorylation and Wnt/ β -catenin pathway against renal fibrosis. <i>British Journal of Pharmacology</i> , 2018, 175, 2689-2708.	5.4	154

#	ARTICLE	IF	CITATIONS
109	Dynamics of Zebrafish Heart Regeneration Using an HPLC-ESI-MS/MS Approach. <i>Journal of Proteome Research</i> , 2018, 17, 1300-1308.	3.7	17
110	Novel RAS Inhibitors Poricoic Acid ZG and Poricoic Acid ZH Attenuate Renal Fibrosis via a Wnt/ β 2-Catenin Pathway and Targeted Phosphorylation of smad3 Signaling. <i>Journal of Agricultural and Food Chemistry</i> , 2018, 66, 1828-1842.	5.2	115
111	Altered ADAMTS5 Expression and Versican Proteolysis: A Possible Molecular Mechanism in Barlow's Disease. <i>Annals of Thoracic Surgery</i> , 2018, 105, 1144-1151.	1.3	3
112	Preparation of novel marine antifouling polyurethane coating materials. <i>Polymer Bulletin</i> , 2018, 75, 5143-5162.	3.3	18
113	Power and sample size calculations for high-throughput sequencing-based experiments. <i>Briefings in Bioinformatics</i> , 2018, 19, 1247-1255.	6.5	32
114	Strategies for processing and quality control of Illumina genotyping arrays. <i>Briefings in Bioinformatics</i> , 2018, 19, 765-775.	6.5	57
115	Tri-allelic heteroplasmies, DNA-RNA differences and their polynucleotide tract associations in the mitochondrial genome. <i>Genomics</i> , 2018, 110, 211-220.	2.9	2
116	Frequent <i>BRAF</i> mutations suggest a novel oncogenic driver in colonic neuroendocrine carcinoma. <i>Journal of Surgical Oncology</i> , 2018, 117, 284-289.	1.7	21
117	Construction of a composite photocatalyst with significantly enhanced photocatalytic performance through combination of homo-junction with hetero-junction. <i>Catalysis Science and Technology</i> , 2018, 8, 486-498.	4.1	36
118	Genomic Profiling of T-Cell Neoplasms Reveals Frequent <i>JAK1</i> and <i>JAK3</i> Mutations With Clonal Evasion From Targeted Therapies. <i>JCO Precision Oncology</i> , 2018, 2018, 1-16.	3.0	23
119	Bi-stream CNN Down Syndrome screening model based on genotyping array. <i>BMC Medical Genomics</i> , 2018, 11, 105.	1.5	4
120	The Landscape of Small Non-Coding RNAs in Triple-Negative Breast Cancer. <i>Genes</i> , 2018, 9, 29.	2.4	21
121	Study of disease phenotype and its association with prognosis of paediatric inflammatory bowel disease in China. <i>BMC Pediatrics</i> , 2018, 18, 229.	1.7	14
122	RnaSeqSampleSize: real data based sample size estimation for RNA sequencing. <i>BMC Bioinformatics</i> , 2018, 19, 191.	2.6	59
123	Multi-perspective quality control of Illumina RNA sequencing data analysis. <i>Briefings in Functional Genomics</i> , 2017, 16, elw035.	2.7	68
124	Integrated molecular analysis reveals complex interactions between genomic and epigenomic alterations in esophageal adenocarcinomas. <i>Scientific Reports</i> , 2017, 7, 40729.	3.3	20
125	Comprehensive evaluation of extracellular small RNA isolation methods from serum in high throughput sequencing. <i>BMC Genomics</i> , 2017, 18, 50.	2.8	28
126	Improvements and impacts of GRCh38 human reference on high throughput sequencing data analysis. <i>Genomics</i> , 2017, 109, 83-90.	2.9	123

#	ARTICLE	IF	CITATIONS
127	Maternal BMI as a predictor of methylation of obesity-related genes in saliva samples from preschool-age Hispanic children at-risk for obesity. <i>BMC Genomics</i> , 2017, 18, 57.	2.8	38
128	MicroRNAs and tRNA-derived fragments predict the transformation of myelodysplastic syndromes to acute myeloid leukemia. <i>Leukemia and Lymphoma</i> , 2017, 58, 2144-2155.	1.3	26
129	Comparison of triple-negative breast cancer molecular subtyping using RNA from matched fresh-frozen versus formalin-fixed paraffin-embedded tissue. <i>BMC Cancer</i> , 2017, 17, 241.	2.6	27
130	StrandScript: evaluation of Illumina genotyping array design and strand correction. <i>Bioinformatics</i> , 2017, 33, 2399-2401.	4.1	15
131	Dracorhodin Perochlorate attenuates <i>Staphylococcus aureus</i> USA300 virulence by decreasing $\hat{\iota}$ -toxin expression. <i>World Journal of Microbiology and Biotechnology</i> , 2017, 33, 17.	3.6	10
132	Profiles of Long Noncoding RNAs in Human Naive and Memory T Cells. <i>Journal of Immunology</i> , 2017, 199, 547-558.	0.8	25
133	Hypoplastic Left Heart Syndrome Sequencing Reveals a Novel NOTCH1 Mutation in a Family with Single Ventricle Defects. <i>Pediatric Cardiology</i> , 2017, 38, 1232-1240.	1.3	25
134	Expression Profiling of Clinical Specimens Supports the Existence of Neural Progenitor-Like Stem Cells in Basal Breast Cancers. <i>Clinical Breast Cancer</i> , 2017, 17, 298-306.e7.	2.4	22
135	Estimating relative mitochondrial DNA copy number using high throughput sequencing data. <i>Genomics</i> , 2017, 109, 457-462.	2.9	16
136	Inhibition of Farnesyltransferase Potentiates NOTCH-Targeted Therapy against Glioblastoma Stem Cells. <i>Stem Cell Reports</i> , 2017, 9, 1948-1960.	4.8	20
137	The MAPK Pathway Regulates Intrinsic Resistance to BET Inhibitors in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 2027-2037.	7.0	59
138	Down syndrome prediction/screening model based on deep learning and illumina genotyping array. , 2017, , .		4
139	Recent Advances in High Throughput Sequencing Analysis. <i>International Journal of Genomics</i> , 2017, 2017, 1-1.	1.6	1
140	Current Research on Non-Coding Ribonucleic Acid (RNA). <i>Genes</i> , 2017, 8, 366.	2.4	54
141	The discrepancy among single nucleotide variants detected by DNA and RNA high throughput sequencing data. <i>BMC Genomics</i> , 2017, 18, 690.	2.8	30
142	Targeted sequencing of established and candidate colorectal cancer genes in the Colon Cancer Family Registry Cohort. <i>Oncotarget</i> , 2017, 8, 93450-93463.	1.8	23
143	The International Conference on Intelligent Biology and Medicine (ICIBM) 2016: summary and innovation in genomics. <i>BMC Genomics</i> , 2017, 18, 703.	2.8	4
144	The Utilization of Formalin Fixed-Paraffin-Embedded Specimens in High Throughput Genomic Studies. <i>International Journal of Genomics</i> , 2017, 2017, 1-9.	1.6	59

#	ARTICLE	IF	CITATIONS
145	Targeted Gene Next-Generation Sequencing in Chinese Children with Chronic Pancreatitis and Acute Recurrent Pancreatitis. <i>Journal of Pediatrics</i> , 2017, 191, 158-163.e3.	1.8	32
146	RNA Sequencing of Formalin-Fixed, Paraffin-Embedded Specimens for Gene Expression Quantification and Data Mining. <i>International Journal of Genomics</i> , 2016, 2016, 1-10.	1.6	15
147	BET bromodomain inhibitors suppress EWS-FLI1-dependent transcription and the IGF1 autocrine mechanism in Ewing sarcoma. <i>Oncotarget</i> , 2016, 7, 43504-43517.	1.8	35
148	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	8.4	118
149	Practicability of detecting somatic point mutation from RNA high throughput sequencing data. <i>Genomics</i> , 2016, 107, 163-169.	2.9	30
150	NOTCH1 and SOX10 are Essential for Proliferation and Radiation Resistance of Cancer Stem-“Like Cells in Adenoid Cystic Carcinoma. <i>Clinical Cancer Research</i> , 2016, 22, 2083-2095.	7.0	46
151	Kinesin-5 inhibitor resistance is driven by kinesin-12. <i>Journal of Cell Biology</i> , 2016, 213, 213-227.	5.2	95
152	lncRNA expression in the auditory forebrain during postnatal development. <i>Gene</i> , 2016, 593, 201-216.	2.2	18
153	Practicability of mitochondrial heteroplasmy detection through an Illumina genotyping array. <i>Mitochondrion</i> , 2016, 31, 75-78.	3.4	3
154	Heterozygosity Ratio, a Robust Global Genomic Measure of Autozygosity and Its Association with Height and Disease Risk. <i>Genetics</i> , 2016, 204, 893-904.	2.9	33
155	Analysis of epithelial-mesenchymal transition markers in the histogenesis of hepatic progenitor cell in HBV-related liver diseases. <i>Diagnostic Pathology</i> , 2016, 11, 136.	2.0	5
156	A novel resistance mutation in <i>eccC</i> of the ESX-5 secretion system confers ofloxacin resistance in <i>Mycobacterium tuberculosis</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 2419-2427.	3.0	11
157	A micro-RNA expression signature for human NAFLD progression. <i>Journal of Gastroenterology</i> , 2016, 51, 1022-1030.	5.1	74
158	Antagonistic roles for the ubiquitin ligase Asr1 and the ubiquitin-specific protease Ubp3 in subtelomeric gene silencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1309-1314.	7.1	6
159	Mitochondria single nucleotide variation across six blood cell types. <i>Mitochondrion</i> , 2016, 28, 16-22.	3.4	8
160	Verbascoside Alleviates Pneumococcal Pneumonia by Reducing Pneumolysin Oligomers. <i>Molecular Pharmacology</i> , 2016, 89, 376-387.	2.3	49
161	Mitochondria sequence mapping strategies and practicability of mitochondria variant detection from exome and RNA sequencing data. <i>Briefings in Bioinformatics</i> , 2016, 17, 224-232.	6.5	29
162	BRAF mutations in colonic high-grade neuroendocrine carcinoma. <i>Journal of Clinical Oncology</i> , 2016, 34, 612-612.	1.6	1

#	ARTICLE	IF	CITATIONS
163	Comprehensive mutation screening for 10 genes in Chinese patients suffering very early onset inflammatory bowel disease. <i>World Journal of Gastroenterology</i> , 2016, 22, 5578.	3.3	30
164	Population structure analysis on 2504 individuals across 26 ancestries using bioinformatics approaches. <i>BMC Bioinformatics</i> , 2015, 16, P19.	2.6	2
165	Practicality of identifying mitochondria variants from exome and RNAseq data. <i>BMC Bioinformatics</i> , 2015, 16, P6.	2.6	3
166	Transfer RNA detection by small RNA deep sequencing and disease association with myelodysplastic syndromes. <i>BMC Genomics</i> , 2015, 16, 727.	2.8	42
167	A novel splice site mutation in <i>SMARCAL1</i> results in aberrant exon definition in a child with schimke immunosseous dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2260-2264.	1.2	15
168	Genetic Analysis Identifies DDR2 as a Novel Gene Affecting Bone Mineral Density and Osteoporotic Fractures in Chinese Population. <i>PLoS ONE</i> , 2015, 10, e0117102.	2.5	6
169	Epidemiological and Molecular Characteristics of the PB1-F2 Proteins in H7N9 Influenza Viruses, Jiangsu. <i>BioMed Research International</i> , 2015, 2015, 1-8.	1.9	5
170	Advances in Computational Genomics. <i>BioMed Research International</i> , 2015, 2015, 1-2.	1.9	0
171	RNAseq by Total RNA Library Identifies Additional RNAs Compared to Poly(A) RNA Library. <i>BioMed Research International</i> , 2015, 2015, 1-9.	1.9	34
172	Deletion of the BMP receptor BMPR1a impairs mammary tumor formation and metastasis. <i>Oncotarget</i> , 2015, 6, 22890-22904.	1.8	37
173	MiR-195 suppresses non-small cell lung cancer by targeting CHEK1. <i>Oncotarget</i> , 2015, 6, 9445-9456.	1.8	140
174	(2-Chlorobenzoyloxy)copper catalyzed C-S cross-coupling of di(hetero)aryl disulfides with aryl boronic acids under base-free conditions. <i>RSC Advances</i> , 2015, 5, 45479-45483.	3.6	20
175	Enforced expression of E47 has differential effects on Lmo2-induced T-cell leukemias. <i>Leukemia Research</i> , 2015, 39, 100-109.	0.8	8
176	Bile diversion to the distal small intestine has comparable metabolic benefits to bariatric surgery. <i>Nature Communications</i> , 2015, 6, 7715.	12.8	156
177	Enhancer of Rudimentary Homolog Affects the Replication Stress Response through Regulation of RNA Processing. <i>Molecular and Cellular Biology</i> , 2015, 35, 2979-2990.	2.3	26
178	Expression and functions of long noncoding RNAs during human T helper cell differentiation. <i>Nature Communications</i> , 2015, 6, 6932.	12.8	172
179	Defective structural RNA processing in relapsing-remitting multiple sclerosis. <i>Genome Biology</i> , 2015, 16, 58.	9.6	19
180	Genome measures used for quality control are dependent on gene function and ancestry. <i>Bioinformatics</i> , 2015, 31, 318-323.	4.1	134

#	ARTICLE	IF	CITATIONS
181	Transcriptional maturation of the mouse auditory forebrain. BMC Genomics, 2015, 16, 606.	2.8	25
182	Mining diverse small RNA species in the deep transcriptome. Trends in Biochemical Sciences, 2015, 40, 4-7.	7.5	60
183	Alternative applications for distinct RNA sequencing strategies. Briefings in Bioinformatics, 2015, 16, 629-639.	6.5	35
184	A Cell-Based Systems Biology Assessment of Human Blood to Monitor Immune Responses after Influenza Vaccination. PLoS ONE, 2015, 10, e0118528.	2.5	79
185	Advanced Datamining Using RNAseq Data. Advances in Bioinformatics and Biomedical Engineering Book Series, 2015, , 1-20.	0.4	1
186	Advanced Heat Map and Clustering Analysis Using Heatmap3. BioMed Research International, 2014, 2014, 1-6.	1.9	252
187	MultiRankSeq: Multiperspective Approach for RNAseq Differential Expression Analysis and Quality Control. BioMed Research International, 2014, 2014, 1-8.	1.9	50
188	Detection of internal exon deletion with exon Del. BMC Bioinformatics, 2014, 15, 332.	2.6	12
189	LMO2 induces T-cell leukemia with epigenetic deregulation of CD4. Experimental Hematology, 2014, 42, 581-593.e5.	0.4	8
190	Multi-perspective quality control of Illumina exome sequencing data using QC3. Genomics, 2014, 103, 323-328.	2.9	79
191	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. Nature Genetics, 2014, 46, 533-542.	21.4	212
192	Three-stage quality control strategies for DNA re-sequencing data. Briefings in Bioinformatics, 2014, 15, 879-889.	6.5	141
193	Multistage genome-wide association meta-analyses identified two new loci for bone mineral density. Human Molecular Genetics, 2014, 23, 1923-1933.	2.9	130
194	Illumina human exome genotyping array clustering and quality control. Nature Protocols, 2014, 9, 2643-2662.	12.0	153
195	Incidence, risk factors and clinical outcomes of peripherally inserted central catheter spontaneous dislodgment in oncology patients: A prospective cohort study. International Journal of Nursing Studies, 2014, 51, 955-963.	5.6	32
196	High-throughput sequencing in mitochondrial DNA research. Mitochondrion, 2014, 17, 157-163.	3.4	71
197	RNAseqPS: A Web Tool for Estimating Sample Size and Power for RNAseq Experiment. Cancer Informatics, 2014, 13s6, CIN.S17688.	1.9	28
198	A comparison of microRNA sequencing reproducibility and noise reduction using mirVana and TRIzol isolation methods. International Journal of Computational Biology and Drug Design, 2014, 7, 102.	0.3	11

#	ARTICLE	IF	CITATIONS
199	Statistical strategies for microRNAseq batch effect reduction. <i>Translational Cancer Research</i> , 2014, 3, 260-265.	1.0	19
200	miRNA Predictors of Response to DNA Methyltransferase Inhibitors. <i>Blood</i> , 2014, 124, 1913-1913.	1.4	0
201	Finding the lost treasures in exome sequencing data. <i>Trends in Genetics</i> , 2013, 29, 593-599.	6.7	124
202	Very Low-Level Heteroplasmy mtDNA Variations Are Inherited in Humans. <i>Journal of Genetics and Genomics</i> , 2013, 40, 607-615.	3.9	63
203	Evaluation of read count based RNAseq analysis methods. <i>BMC Genomics</i> , 2013, 14, S2.	2.8	53
204	Gene-gene interaction between <i>RBMS3</i> and <i>ZNF516</i> influences bone mineral density. <i>Journal of Bone and Mineral Research</i> , 2013, 28, 828-837.	2.8	21
205	Copy Number Variation on Chromosome 10q26.3 for Obesity Identified by a Genome-Wide Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, E191-E195.	3.6	19
206	Suggestion of GLYAT gene underlying variation of bone size and body lean mass as revealed by a bivariate genome-wide association study. <i>Human Genetics</i> , 2013, 132, 189-199.	3.8	50
207	An evaluation of allele frequency estimation accuracy using pooled sequencing data. <i>International Journal of Computational Biology and Drug Design</i> , 2013, 6, 279.	0.3	6
208	Comparative Study of Exome Copy Number Variation Estimation Tools Using Array Comparative Genomic Hybridization as Control. <i>BioMed Research International</i> , 2013, 2013, 1-7.	1.9	47
209	MitoSeek: extracting mitochondria information and performing high-throughput mitochondria sequencing analysis. <i>Bioinformatics</i> , 2013, 29, 1210-1211.	4.1	87
210	Meta-analysis identifies a <i>MECOM</i> gene as a novel predisposing factor of osteoporotic fracture. <i>Journal of Medical Genetics</i> , 2013, 50, 212-219.	3.2	30
211	Evaluation of Allele Frequency Estimation Using Pooled Sequencing Data Simulation. <i>Scientific World Journal, The</i> , 2013, 2013, 1-9.	2.1	16
212	Large Scale Comparison of Gene Expression Levels by Microarrays and RNAseq Using TCGA Data. <i>PLoS ONE</i> , 2013, 8, e71462.	2.5	189
213	Mechanistic Phenotypes: An Aggregative Phenotyping Strategy to Identify Disease Mechanisms Using GWAS Data. <i>PLoS ONE</i> , 2013, 8, e81503.	2.5	15
214	The use of next generation sequencing technology to study the effect of radiation therapy on mitochondrial DNA mutation. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2012, 744, 154-160.	1.7	49
215	Exome sequencing generates high quality data in non-target regions. <i>BMC Genomics</i> , 2012, 13, 194.	2.8	130
216	The effect of strand bias in Illumina short-read sequencing data. <i>BMC Genomics</i> , 2012, 13, 666.	2.8	102

#	ARTICLE	IF	CITATIONS
217	Genome-wide association study identifies HMG3 locus for spine bone size variation in Chinese. <i>Human Genetics</i> , 2012, 131, 463-469.	3.8	6
218	Plant growth and architectural modelling and its applications. <i>Annals of Botany</i> , 2011, 107, 723-727.	2.9	51
219	Genetic Association Study of Common Mitochondrial Variants on Body Fat Mass. <i>PLoS ONE</i> , 2011, 6, e21595.	2.5	30
220	Copy Number Variation in CNP267 Region May Be Associated with Hip Bone Size. <i>PLoS ONE</i> , 2011, 6, e22035.	2.5	5
221	The Fat Mass and Obesity Associated Gene, FTO, Is Also Associated with Osteoporosis Phenotypes. <i>PLoS ONE</i> , 2011, 6, e27312.	2.5	38
222	Paranuclear Dot-like Immunostaining for CD99. <i>American Journal of Surgical Pathology</i> , 2011, 35, 799-806.	3.7	74
223	Mitochondria-Wide Association Study of Common Variants in Osteoporosis. <i>Annals of Human Genetics</i> , 2011, 75, 569-574.	0.8	33
224	Assessment of the influence of global dimming on the photosynthetic production of rice based on three-dimensional modeling. <i>Science China Earth Sciences</i> , 2011, 54, 290-297.	5.2	13
225	Pathway-Based Association Analyses Identified TRAIL Pathway for Osteoporotic Fractures. <i>PLoS ONE</i> , 2011, 6, e21835.	2.5	14
226	<i>IL21R</i> and <i>PTH</i> may underlie variation of femoral neck bone mineral density as revealed by a genome-wide association study. <i>Journal of Bone and Mineral Research</i> , 2010, 25, 1042-1048.	2.8	36
227	Genome-wide association study for femoral neck bone geometry. <i>Journal of Bone and Mineral Research</i> , 2010, 25, 320-329.	2.8	43
228	Pathway-based genome-wide association analysis identified the importance of regulation-of-autophagy pathway for ultradistal radius BMD. <i>Journal of Bone and Mineral Research</i> , 2010, 25, 1572-1580.	2.8	103
229	<i>HMGA2</i> Is Confirmed To Be Associated with Human Adult Height. <i>Annals of Human Genetics</i> , 2010, 74, 11-16.	0.8	32
230	Molecular Genetic Studies of Gene Identification for Osteoporosis: The 2009 Update. <i>Endocrine Reviews</i> , 2010, 31, 447-505.	20.1	76
231	Genome-Wide Association Study Identifies <i>ALDH7A1</i> as a Novel Susceptibility Gene for Osteoporosis. <i>PLoS Genetics</i> , 2010, 6, e1000806.	3.5	101
232	Runs of Homozygosity Identify a Recessive Locus 12q21.31 for Human Adult Height. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2010, 95, 3777-3782.	3.6	33
233	The regulation-of-autophagy pathway may influence Chinese stature variation: evidence from elder adults. <i>Journal of Human Genetics</i> , 2010, 55, 441-447.	2.3	20
234	A genome wide association study between copy number variation (CNV) and human height in Chinese population. <i>Journal of Genetics and Genomics</i> , 2010, 37, 779-785.	3.9	12

#	ARTICLE	IF	CITATIONS
235	Calibration of Topological Development in the Procedure of Parametric Identification: Application of the Stochastic GreenLab Model for <i>Pinus sylvestris</i> var. <i>mongolica</i> . , 2009, , .		2
236	Genome-wide association study identifies two novel loci containing <i>FLNB</i> and <i>SBF2</i> genes underlying stature variation. <i>Human Molecular Genetics</i> , 2009, 18, 1661-1669.	2.9	27
237	Genome-wide association study suggested copy number variation may be associated with body mass index in the Chinese population. <i>Journal of Human Genetics</i> , 2009, 54, 199-202.	2.3	78
238	Powerful Bivariate Genome-Wide Association Analyses Suggest the <i>SOX6</i> Gene Influencing Both Obesity and Osteoporosis Phenotypes in Males. <i>PLoS ONE</i> , 2009, 4, e6827.	2.5	87
239	Genome-wide association scan for stature in Chinese: evidence for ethnic specific loci. <i>Human Genetics</i> , 2009, 125, 1-9.	3.8	39
240	Association Analyses of <i>RANKL/RANK/OPG</i> Gene Polymorphisms with Femoral Neck Compression Strength Index Variation in Caucasians. <i>Calcified Tissue International</i> , 2009, 85, 104-112.	3.1	38
241	Genome-wide Association and Follow-Up Replication Studies Identified <i>ADAMTS18</i> and <i>TGFBR3</i> as Bone Mass Candidate Genes in Different Ethnic Groups. <i>American Journal of Human Genetics</i> , 2009, 84, 388-398.	6.2	187
242	Genome-wide Association and Replication Studies Identified <i>TRHR</i> as an Important Gene for Lean Body Mass. <i>American Journal of Human Genetics</i> , 2009, 84, 418-423.	6.2	103
243	Simulating the Distribution of R/FR in Maize Canopies with Monte Carlo Ray Tracing Approach. , 2009, , .		31
244	Assessment of the Effects of Leaf Angle Combinations on Potential Photosynthesis Capacity of Rice with 3-D Models Using High Performance Computing. , 2009, , .		3
245	Whole Genome Distribution and Ethnic Differentiation of Copy Number Variation in Caucasian and Asian Populations. <i>PLoS ONE</i> , 2009, 4, e7958.	2.5	51
246	Sex-Specific Association of the Glucocorticoid Receptor Gene With Extreme BMD. <i>Journal of Bone and Mineral Research</i> , 2008, 23, 247-252.	2.8	15
247	Chromosomal regions 22q13 and 3p25 may harbor quantitative trait loci influencing both age at menarche and bone mineral density. <i>Human Genetics</i> , 2008, 123, 419-427.	3.8	19
248	Genome-wide Copy-Number-Variation Study Identified a Susceptibility Gene, <i>UGT2B17</i> , for Osteoporosis. <i>American Journal of Human Genetics</i> , 2008, 83, 663-674.	6.2	209
249	Comprehensive association analyses of <i>IGF1</i> , <i>ESR2</i> , and <i>CYP17</i> genes with adult height in Caucasians. <i>European Journal of Human Genetics</i> , 2008, 16, 1380-1387.	2.8	13
250	Polymorphisms in the estrogen receptor genes are associated with hip fractures in Chinese. <i>Bone</i> , 2008, 43, 910-914.	2.9	23
251	Comparison of architecture among different cultivars of hybrid rice using a spatial light model based on 3-D digitising. <i>Functional Plant Biology</i> , 2008, 35, 900.	2.1	73
252	Bivariate genome linkage analysis suggests pleiotropic effects on chromosomes 20p and 3p for body fat mass and lean mass. <i>Genetical Research</i> , 2008, 90, 259-268.	0.9	6

#	ARTICLE	IF	CITATIONS
253	Modelling three-dimensional architecture of pine tree (<i>Pinus sylvestris</i> Linn.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 747 903-909.	1.6	4
254	Plant Modeling and Its Applications to Agriculture. , 2006, , .		5
255	Parameter Optimization and Field Validation of the Functional-Structural Model GREENLAB for Maize. Annals of Botany, 2006, 97, 217-230.	2.9	177
256	Hyperactivation of Torc1 Drives Resistance to the Pan-Her Tyrosine Kinase Inhibitor Neratinib in Her2-Mutant Cancers. SSRN Electronic Journal, 0, , .	0.4	0