Finn Cilius Nielsen

List of Publications by Year
in descending order

148 papers	6,960 citations	76326 40 h-index	66911 78 g-index
154 all docs	154 docs citations	154 times ranked	13456 citing authors

#	Article	IF	Citations
1	MicroRNA-10a Binds the 5â€²UTR of Ribosomal Protein mRNAs and Enhances Their Translation. Molecular Cell, 2008, 30, 460-471.	9.7	1,168
2	Ascertainment Biases in SNP Chips Affect Measures of Population Divergence. Molecular Biology and Evolution, 2010, 27, 2534-2547.	8.9	317
3	Hereditary breast and ovarian cancer: new genes in confined pathways. Nature Reviews Cancer, 2016, 16, 599-612.	28.4	305
4	The transcriptional program of terminal granulocytic differentiation. Blood, 2005, 105, 1785-1796.	1.4	249
5	NMD is essential for hematopoietic stem and progenitor cells and for eliminating by-products of programmed DNA rearrangements. Genes and Development, 2008, 22, 1381-1396.	5.9	231
6	Mutational spectrum in a worldwide study of 29,700 families withBRCA1orBRCA2mutations. Human Mutation, 2018, 39, 593-620.	2.5	224
7	A Library of 7TM Receptor C-terminal Tails. Journal of Biological Chemistry, 2004, 279, 54291-54303.	3.4	147
8	H19 RNA Binds Four Molecules of Insulin-like Growth Factor II mRNA-binding Protein. Journal of Biological Chemistry, 2000, 275, 29562-29569.	3.4	142
9	Quantitative miRNA expression analysis: Comparing microarrays with next-generation sequencing. Rna, 2009, 15, 2028-2034.	3.5	142
10	Papillon-LefÃ¨vre syndrome patient reveals species-dependent requirements for neutrophil defenses. Journal of Clinical Investigation, 2014, 124, 4539-4548.	8.2	141
11	Helicobacter pylori Infection Induces Genetic Instability of Nuclear and Mitochondrial DNA in Gastric Cells. Clinical Cancer Research, 2009, 15, 2995-3002.	7.0	123
12	Characterization of osteoclasts derived from CD14+ monocytes isolated from peripheral blood. Journal of Bone and Mineral Metabolism, 2006, 25, 36-45.	2.7	122
13	Weaning Triggers a Maturation Step of Pancreatic Î² Cells. Developmental Cell, 2015, 32, 535-545.	7.0	120
14	Continuing rise in oropharyngeal cancer in a high HPV prevalence area: A Danish population-based study from 2011 to 2014. European Journal of Cancer, 2017, 70, 75-82.	2.8	115
15	Sequential dimerization of human zipcode-binding protein IMP1 on RNA: a cooperative mechanism providing RNP stability. Nucleic Acids Research, 2004, 32, 4368-4376.	14.5	99
16	Relatedness mapping and tracts of relatedness for genomeâ€wide data in the presence of linkage disequilibrium. Genetic Epidemiology, 2009, 33, 266-274.	1.3	99
17	MicroRNAs Show Mutually Exclusive Expression Patterns in the Brain of Adult Male Rats. PLoS ONE, 2009, 4, e7225.	2.5	94
18	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	12.8	88

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19	The molecular and cellular biology of insulin-like growth factor II. Progress in Growth Factor Research, 1992, 4, 257-290.	1.6	87
20	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. Genetics in Medicine, 2020, 22, 1653-1666.	2.4	82
21	Reduced Frequency of Extracolonic Cancers in Hereditary Nonpolyposis Colorectal Cancer Families with Monoallelic hMLH1Expression. American Journal of Human Genetics, 1997, 61, 129-138.	6.2	79
22	Down-regulation of microRNAs controlling tumourigenic factors in follicular thyroid carcinoma. Journal of Molecular Endocrinology, 2012, 48, 11-23.	2.5	73
23	Characterization of human exonuclease 1 in complex with mismatch repair proteins, subcellular localization and association with PCNA. Oncogene, 2004, 23, 1457-1468.	5.9	68
24	Câ€MYC and IGFâ€II mRNAâ€binding protein (CRDâ€BP/IMPâ€1) in benign and malignant mesenchymal tumors. International Journal of Cancer, 2001, 94, 480-484.	5.1	63
25	High frequency of pathogenic germline variants within homologous recombination repair in patients with advanced cancer. Npj Genomic Medicine, 2019, 4, 13.	3.8	63
26	HNPCC mutations in the human DNA mismatch repair gene hMLH1 influence assembly of hMutLÎ± and hMLH1â€“hEXO1 complexes. Oncogene, 2001, 20, 3590-3595.	5.9	61
27	ADAM12 Alleviates the Skeletal Muscle Pathology in mdx Dystrophic Mice. American Journal of Pathology, 2002, 161, 1535-1540.	3.8	61
28	De novo mutations in familial adenomatous polyposis (FAP). European Journal of Human Genetics, 2002, 10, 631-637.	2.8	61
29	Copenhagen Prospective Personalized Oncology (CoPPO)â€”Clinical Utility of Using Molecular Profiling to Select Patients to Phase I Trials. Clinical Cancer Research, 2019, 25, 1239-1247.	7.0	59
30	Gene Expression Profiling of Placentas Affected by Pre-Eclampsia. Journal of Biomedicine and Biotechnology, 2010, 2010, 1-11.	3.0	56
31	A Transgenic Mouse Marking Live Replicating Cells Reveals InÂVivo Transcriptional Program of Proliferation. Developmental Cell, 2012, 23, 681-690.	7.0	54
32	ADAM12 redistributes and activates MMP-14, resulting in gelatin degradation, reduced apoptosis, and increased tumor growth. Journal of Cell Science, 2013, 126, 4707-20.	2.0	50
33	Application of wholeâ€exome sequencing to direct the specific functional testing and diagnosis of rare inherited bleeding disorders in patients from the Ã–resund Region, Scandinavia. British Journal of Haematology, 2017, 179, 308-322.	2.5	49
34	Brain response to traumatic brain injury in wildâ€type and interleukinâ€6 knockout mice: a microarray analysis. Journal of Neurochemistry, 2005, 92, 417-432.	3.9	48
35	Gestational Protein Restriction in Mice Has Pronounced Effects on Gene Expression in Newborn Offspring's Liver and Skeletal Muscle; Protective Effect of Taurine. Pediatric Research, 2010, 67, 47-53.	2.3	47
36	Novel roles for metallothioneinâ€I + II (MTâ€I + II) in defense responses, neurogenesis, and tissue restoration after traumatic brain injury: Insights from global gene expression profiling in wildâ€type and MTâ€I + II knockout mice. Journal of Neuroscience Research, 2006, 84, 1452-1474.	2.9	45

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37	Integrative analyses reveal novel strategies in HPV11,-16 and -45 early infection. Scientific Reports, 2012, 2, 515.	3.3	45
38	Hereditary non-polyposis colorectal cancer (HNPCC): phenotype-genotype correlation between patients with and without identified mutation. Human Mutation, 2002, 20, 20-27.	2.5	43
39	A maternal low protein diet has pronounced effects on mitochondrial gene expression in offspring liver and skeletal muscle; protective effect of taurine. Journal of Biomedical Science, 2010, 17, S38.	7.0	43
40	MicroRNA Changes in Cerebrospinal Fluid After Subarachnoid Hemorrhage. Stroke, 2017, 48, 2391-2398.	2.0	43
41	Survival and Long-Term Biochemical Cure in Medullary Thyroid Carcinoma in Denmark 1997â€“2014: A Nationwide Study. Thyroid, 2019, 29, 368-377.	4.5	43
42	Microarray-based classification of diffuse large B-cell lymphoma. European Journal of Haematology, 2005, 74, 453-465.	2.2	42
43	Nuclear translocation contributes to regulation of DNA excision repair activities. DNA Repair, 2009, 8, 682-689.	2.8	42
44	MicroRNA-based classifiers for diagnosis of oral cavity squamous cell carcinoma in tissue and plasma. Oral Oncology, 2018, 83, 46-52.	1.5	41
45	Functional analysis of HNPCC-related missense mutations in MSH2. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2008, 645, 44-55.	1.0	40
46	Expression of the genesdualoxidase2,lipocalin 2andregenerating islet-derived 1 alphain Crohn's disease. Scandinavian Journal of Gastroenterology, 2007, 42, 454-463.	1.5	39
47	Identification of factors interacting with hMSH2 in the fetal liver utilizing the yeast two-hybrid system. Mutation Research DNA Repair, 2000, 460, 41-52.	3.7	38
48	Characterization of miRNA Expression in Human Degenerative Lumbar Disks. Connective Tissue Research, 2013, 54, 197-203.	2.3	36
49	A new NFIA:RAF1 fusion activating the MAPK pathway in pilocytic astrocytoma. Cancer Genetics, 2016, 209, 440-444.	0.4	36
50	Deregulated Genes in Sporadic Vestibular Schwannomas. Otology and Neurotology, 2010, 31, 256-266.	1.3	34
51	Effect of astrocyteâ€targeted production of ILâ€6 on traumatic brain injury and its impact on the cortical transcriptome. Developmental Neurobiology, 2008, 68, 195-208.	3.0	33
52	14-3-3 checkpoint regulatory proteins interact specifically with DNA repair protein human exonuclease 1 (hEXO1) via a semi-conserved motif. DNA Repair, 2012, 11, 267-277.	2.8	33
53	Gene expression demonstrates an immunological capacity of the human endolymphatic sac. Laryngoscope, 2015, 125, E269-75.	2.0	33
54	Is thyroidectomy necessary in RET mutations carriers of the familial medullary thyroid carcinoma syndrome?. Cancer, 2000, 89, 863-867.	4.1	32

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55	Prognostic significance of metallothionein in B-cell lymphomas. Blood, 2006, 108, 3514-3519.	1.4	32
56	Nav1.8 channelopathy in mutant mice deficient for myelin protein zero is detrimental to motor axons. Brain, 2011, 134, 585-601.	7.6	32
57	Incidence and prevalence of multiple endocrine neoplasia 2B in Denmark: a nationwide study. Endocrine-Related Cancer, 2017, 24, L39-L42.	3.1	32
58	Incidence and prevalence of sporadic and hereditary MTC in Denmark 1960â€“2014: a nationwide study. Endocrine Connections, 2018, 7, 829-839.	1.9	32
59	Drosophila Imp iCLIP identifies an RNA assemblage coordinating F-actin formation. Genome Biology, 2015, 16, 123.	8.8	31
60	A Comparison of Tools for Copy-Number Variation Detection in Germline Whole Exome and Whole Genome Sequencing Data. Cancers, 2021, 13, 6283.	3.7	31
61	Nuclear localization of human DNA mismatch repair protein exonuclease 1 (hEXO1). Nucleic Acids Research, 2007, 35, 2609-2619.	14.5	30
62	Influence of Smoking on Colonic Gene Expression Profile in Crohn's Disease. PLoS ONE, 2009, 4, e6210.	2.5	30
63	The biphasic expression of IMP/Vg1-RBP is conserved between vertebrates and Drosophila. Mechanisms of Development, 2000, 96, 129-132.	1.7	29
64	Distribution of RET Mutations in Multiple Endocrine Neoplasia 2 in Denmark 1994â€“2014: A Nationwide Study. Thyroid, 2017, 27, 215-223.	4.5	29
65	Molecular signatures of thyroid follicular neoplasia. Endocrine-Related Cancer, 2010, 17, 691-708.	3.1	28
66	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. Npj Breast Cancer, 2019, 5, 38.	5.2	28
67	Gene Expression in the Human Endolymphatic Sac. Otology and Neurotology, 2015, 36, 915-922.	1.3	27
68	Cell-free DNA in newly diagnosed patients with glioblastoma â€“ a clinical prospective feasibility study. Oncotarget, 2019, 10, 4397-4406.	1.8	27
69	IMP3 Expression in Human Ovarian Cancer is Associated With Improved Survival. International Journal of Gynecological Pathology, 2009, 28, 203-210.	1.4	26
70	RNA assemblages orchestrate complex cellular processes. BioEssays, 2016, 38, 674-681.	2.5	23
71	Whole genome sequencing of breast cancer. Apmis, 2019, 127, 303-315.	2.0	23
72	Differential expression of cellular microRNAs in HPV 11, -16, and -45 transfected cells. Biochemical and Biophysical Research Communications, 2011, 412, 20-25.	2.1	22

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73	Deconvolution of autoencoders to learn biological regulatory modules from single cell mRNA sequencing data. BMC Bioinformatics, 2019, 20, 379.	2.6	22
74	Single mRNP Analysis Reveals that Small Cytoplasmic mRNP Granules Represent mRNA Singletons. Cell Reports, 2019, 29, 736-748.e4.	6.4	22
75	Diverging mechanisms for TNFâ€ī± receptors in normal mouse brains and in functional recovery after injury: From gene to behavior. Journal of Neuroscience Research, 2007, 85, 2668-2685.	2.9	21
76	Functional characterization ofMLH1missense variants identified in lynch syndrome patients. Human Mutation, 2012, 33, 1647-1655.	2.5	21
77	Genetic screens to identify pathogenic gene variants in the common cancer predisposition Lynch syndrome. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 9403-9408.	7.1	21
78	Frequency of the hemochromatosis HFE mutations C282Y, H63D, and S65C in blood donors in the Faroe Islands. Annals of Hematology, 2005, 84, 146-149.	1.8	20
79	Polyposis and early cancer in a patient with low penetrant mutations in MSH6 and APC: hereditary colorectal cancer as a polygenic trait. International Journal of Colorectal Disease, 2006, 21, 847-850.	2.2	20
80	Incidence and prevalence of multiple endocrine neoplasia 2A in Denmark 1901–2014: a nationwide study. Clinical Epidemiology, 2018, Volume 10, 1479-1487.	3.0	19
81	High Prevalence of Papillary Thyroid Microcarcinoma in Danish Patients: A Prospective Study of 854 Consecutive Patients with a Cold Thyroid Nodule Undergoing Fine-Needle Aspiration. European Thyroid Journal, 2012, 1, 110-117.	2.4	18
82	Personalized oncology: genomic screening in phase 1. Apmis, 2014, 122, 723-733.	2.0	18
83	Chromothripsis and DNA Repair Disorders. Journal of Clinical Medicine, 2020, 9, 613.	2.4	18
84	Molecular Forms and Regional Distribution of Cholecystokinin in the Central Nervous System. Neuroscience Intelligence Unit, 1995, , 33-56.	0.5	18
85	The BTNL2 A allele variant is frequent in Danish patients with sarcoidosis. Clinical Respiratory Journal, 2011, 5, 105-111.	1.6	17
86	Effect of endogenous hypergastrinemia on gastrin receptor expressing human colon carcinoma transplanted to athymic rats. Gastroenterology, 1995, 109, 1415-1420.	1.3	16
87	Clinical phenotype and gene expression profile in Crohn's disease. American Journal of Physiology - Renal Physiology, 2007, 292, G298-G304.	3.4	16
88	Detection and quantification of microRNA in cerebral microdialysate. Journal of Translational Medicine, 2015, 13, 149.	4.4	16
89	Founder Effect of the RET^{C611Y} Mutation in Multiple Endocrine Neoplasia 2A in Denmark: A Nationwide Study. Thyroid, 2017, 27, 1505-1510.	4.5	16
90	The human endolymphatic sac expresses natriuretic peptides. Laryngoscope, 2017, 127, E201-E208.	2.0	15

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91	Gene expression, signal transduction pathways and functional networks associated with growth of sporadic vestibular schwannomas. Journal of Neuro-Oncology, 2017, 131, 283-292.	2.9	15
92	Circulating tumor DNA as a marker of treatment response in BRAF V600E mutated non-melanoma solid tumors. Oncotarget, 2018, 9, 32570-32579.	1.8	15
93	Frequency of the HFE C282Y and H63D mutations in Danish patients with clinical haemochromatosis initially diagnosed by phenotypic methods. European Journal of Haematology, 2003, 71, 403-407.	2.2	14
94	Whole-exome sequencing and genome-wide methylation analyses identify novel disease associated mutations and methylation patterns in idiopathic hypereosinophilic syndrome. Oncotarget, 2015, 6, 40588-40597.	1.8	14
95	Using microarrayâ€based subtyping methods for breast cancer in the era of highâ€throughput RNA sequencing. Molecular Oncology, 2018, 12, 2136-2146.	4.6	14
96	Deep sequencing of human papillomavirus positive loco-regionally advanced oropharyngeal squamous cell carcinomas reveals novel mutational signature. BMC Cancer, 2018, 18, 640.	2.6	14
97	Elevated miR-9 in Cerebrospinal Fluid Is Associated with Poor Functional Outcome After Subarachnoid Hemorrhage. Translational Stroke Research, 2020, 11, 1243-1252.	4.2	14
98	Major driver mutations are shared between sinonasal intestinal-type adenocarcinoma and the morphologically identical colorectal adenocarcinoma. Journal of Cancer Research and Clinical Oncology, 2021, 147, 1019-1027.	2.5	14
99	The Number of Signaling Pathways Altered by Driver Mutations in Chronic Lymphocytic Leukemia Impacts Disease Outcome. Clinical Cancer Research, 2020, 26, 1507-1515.	7.0	13
100	Clinical implications of intrinsic molecular subtypes of breast cancer for sentinel node status. Scientific Reports, 2021, 11, 2259.	3.3	13
101	Unexpectedly high but still asymptomatic iron overload in a patient with pyruvate kinase deficiency. The Hematology Journal, 2004, 5, 543-545.	1.4	12
102	Differential expression of cellular microRNAs in HPV-11 transfected cells. An analysis by three different array platforms and qRT-PCR. Biochemical and Biophysical Research Communications, 2010, 403, 357-362.	2.1	12
103	Germline RBBP8 variants associated with early-onset breast cancer compromise replication fork stability. Journal of Clinical Investigation, 2020, 130, 4069-4080.	8.2	12
104	Targeting the tumor mutanome for personalized vaccination in a TMB low non-small cell lung cancer. , 2022, 10, e003821.		12
105	The Spectrum of FANCM Protein Truncating Variants in European Breast Cancer Cases. Cancers, 2020, 12, 292.	3.7	11
106	Replication of newly proposed TNM staging system for medullary thyroid carcinoma: a nationwide study. Endocrine Connections, 2019, 8, 1-7.	1.9	11
107	CARD15 Status and Familial Predisposition for Crohn's Disease and Colonic Gene Expression. Digestive Diseases and Sciences, 2007, 52, 1783-1789.	2.3	10
108	The synthetic NCAM-derived peptide, FGL, modulates the transcriptional response to traumatic brain injury. Neuroscience Letters, 2008, 437, 148-153.	2.1	10

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109	Embryonic expression of Drosophila IMP in the developing CNS and PNS. Gene Expression Patterns, 2009, 9, 138-143.	0.8	10
110	Development and validation of a microRNA based diagnostic assay for primary tumor site classification of liver core biopsies. Molecular Oncology, 2015, 9, 68-77.	4.6	10
111	Tumor miRNA expression profile is related to vestibular schwannoma growth rate. Acta Neurochirurgica, 2020, 162, 1187-1195.	1.7	10
112	Microarrays and Crohn's disease: Collecting reliable information. Scandinavian Journal of Gastroenterology, 2005, 40, 369-377.	1.5	9
113	Treatment response and colonic gene expression in patients with Crohn's disease. Scandinavian Journal of Gastroenterology, 2007, 42, 834-840.	1.5	9
114	Detection of copy number alterations in cell-free tumor DNA from plasma. BBA Clinical, 2017, 7, 120-126.	4.1	9
115	Plasma total cell-free DNA is a prognostic biomarker of overall survival in metastatic solid tumour patients. British Journal of Cancer, 2019, 121, 125-130.	6.4	9
116	Progression of motor axon dysfunction and ectopic Nav1.8 expression in a mouse model of Charcot-Marie-Tooth disease 1B. Neurobiology of Disease, 2016, 93, 201-214.	4.4	8
117	Completeness of RET testing in patients with medullary thyroid carcinoma in Denmark 1997–2013: a nationwide study. Clinical Epidemiology, 2019, Volume 11, 93-99.	3.0	7
118	Tumor mutational burden and purity adjustment before and after treatment with temozolomide in 27 paired samples of glioblastoma: a prospective study. Molecular Oncology, 2022, 16, 206-218.	4.6	7
119	Presymptomatic diagnosis using a deletion of a single codon in families with hereditary non-polyposis colorectal cancer. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2005, 570, 89-96.	1.0	6
120	Blau syndromeâ€associated mutations in exon 4 of the caspase activating recruitment domain 15 (CARD 15) gene are not found in ethnic Danes with sarcoidosis. Clinical Respiratory Journal, 2007, 1, 74-79.	1.6	6
121	Modeling tissue contamination to improve molecular identification of the primary tumor site of metastases. Bioinformatics, 2014, 30, 1417-1423.	4.1	5
122	A simple, safe and sensitive method for SARSâ€CoVâ€2 inactivation and RNA extraction for RTâ€qPCR. Apmis, 2021, 129, 393-400.	2.0	5
123	Cytoplasmic mRNPs revisited: Singletons and condensates. BioEssays, 2020, 42, e2000097.	2.5	4
124	Genomic Alterations in Human Papillomavirusâ€“Positive and â€“Negative Conjunctival Squamous Cell Carcinomas. , 2021, 62, 11.		4
125	Neuronal Fibers and Neurotransmitter Receptor Expression in the Human Endolymphatic Sac. Otology and Neurotology, 2017, 38, 765-773.	1.3	3
126	Unveiling mRNP composition by fluorescence correlation and cross-correlation spectroscopy using cell lysates. Nucleic Acids Research, 2021, 49, e119-e119.	14.5	3

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127	A catalog of curated breast cancer genes. Breast Cancer Research and Treatment, 2022, 191, 431-441.	2.5	3
128	Interpretable Autoencoders Trained on Single Cell Sequencing Data Can Transfer Directly to Data from Unseen Tissues. Cells, 2022, 11, 85.	4.1	3
129	Gene expression profiling in patients with polymyalgia rheumatica before and after symptom-abolishing glucocorticoid treatment. BMC Musculoskeletal Disorders, 2017, 18, 341.	1.9	2
130	Amplicon-Based NGS Panels for Actionable Cancer Target Identification in Follicular Cell-Derived Thyroid Neoplasia. Frontiers in Endocrinology, 2020, 11, 146.	3.5	2
131	Two Types of Receptor for Insulin-Like Growth Factors are Expressed on Normal and Malignant Cells from Mammalian Brain. , 1987, , 297-313.		2
132	The Predictive Value of Gene Expression Profiles for Acute Graft-Versus-Host Disease after Hematopoietic Cell Transplantation with Nonmyeloablative Conditioning for Hematological Malignancy.. Blood, 2007, 110, 1079-1079.	1.4	2
133	Intestinal metaplasia is a precursor lesion for sinonasal intestinalâ€type adenocarcinoma: genomic investigation of a case proving this hypothesis. Apmis, 2022, 130, 53-56.	2.0	2
134	MicroRNA-9-3p: a novel predictor of neurological outcome after cardiac arrest. European Heart Journal: Acute Cardiovascular Care, 2022, 11, 609-616.	1.0	2
135	Regulation of the Human Cholecystokinin Gene. Annals of the New York Academy of Sciences, 1994, 713, 321-323.	3.8	1
136	Isolation of RNP Granules. Methods in Molecular Biology, 2011, 703, 265-273.	0.9	1
137	Copenhagen prospective personalized oncology (CoPPO): Sequencing and array-based pipeline for selection of patients to phase 1 studies.. Journal of Clinical Oncology, 2014, 32, 11097-11097.	1.6	1
138	Correlation between HPV status at T and N sites of oropharyngeal squamous cell carcinomas. Acta Oto-Laryngologica, 2017, 137, 1260-1264.	0.9	0
139	Characterization of basal-like subtype in a Danish consecutive primary breast cancer cohort. Acta OncolÃ³gica, 2018, 57, 51-57.	1.8	0
140	GENE-30. INCREASED TUMOR MUTATIONAL LOAD AFTER RADIOTHERAPY AND TEMOZOLOMIDE IN PROGRESSING GLIOBLASTOMA A PROSPECTIVE STUDY. Neuro-Oncology, 2018, 20, vi109-vi110.	1.2	0
141	Different Subtypes of Difffuse Large B-Cell Lymphomas Identified Using Microarrays and Phenotyping.. Blood, 2004, 104, 4293-4293.	1.4	0
142	Characterization of miRNA expression in human degenerative lumbar discs. Connective Tissue Research, 0, , 130305100219006.	2.3	0
143	Molecular profiling of tumour budding to implicate TGF-Î² mediated epithelial-mesenchymal transition as a therapeutic target.. Journal of Clinical Oncology, 2015, 33, 6059-6059.	1.6	0
144	Actionable targets in recurrent bile duct and pancreatic cancer in a prospective cohort of patients evaluated by whole exome sequencing and SNP array analysis.. Journal of Clinical Oncology, 2016, 34, e23256-e23256.	1.6	0

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145	Dynamics of mutant BRAF V600E in free circulating DNA (fcDNA) of non-melanoma cancer patients (pts) in response to treatment with BRAF and MEK/EGFR inhibitors.. Journal of Clinical Oncology, 2016, 34, 11531-11531.	1.6	0
146	Prognostic Impact of Subclonal TP53 Aberrations in Chronic Lymphocytic Leukemia Validated By a Robust Targeted Next Generation Sequencing Assay. Blood, 2016, 128, 4380-4380.	1.4	0
147	Abstract 149: Elevated mir-9 in Cerebrospinal Fluid is Associated With Poor Functional Outcome After Subarachnoid Hemorrhage. Stroke, 2019, 50, .	2.0	0
148	Differences in gene expression despite identical histomorphology in sinonasal intestinal type adenocarcinoma and metastases from colorectal adenocarcinoma. Apmis, 0, , .	2.0	0