

Ana Vivancos

List of Publications by Year in descending order

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Version: 2024-02-01

121
papers

6,774
citations

94381

37
h-index

62565

80
g-index

122
all docs

122
docs citations

122
times ranked

13392
citing authors

#	ARTICLE	IF	CITATIONS
1	Acquired EGFR C797S mutation mediates resistance to AZD9291 in non-small cell lung cancer harboring EGFR T790M. <i>Nature Medicine</i> , 2015, 21, 560-562.	15.2	1,280
2	Cerebrospinal fluid-derived circulating tumour DNA better represents the genomic alterations of brain tumours than plasma. <i>Nature Communications</i> , 2015, 6, 8839.	5.8	605
3	Cancer Genome Interpreter annotates the biological and clinical relevance of tumor alterations. <i>Genome Medicine</i> , 2018, 10, 25.	3.6	366
4	β -catenin confers resistance to PI3K and AKT inhibitors and subverts FOXO3a to promote metastasis in colon cancer. <i>Nature Medicine</i> , 2012, 18, 892-901.	15.2	336
5	Immune-Related Gene Expression Profiling After PD-1 Blockade in Non-Small Cell Lung Carcinoma, Head and Neck Squamous Cell Carcinoma, and Melanoma. <i>Cancer Research</i> , 2017, 77, 3540-3550.	0.4	327
6	RAD51 foci as a functional biomarker of homologous recombination repair and PARP inhibitor resistance in germline BRCA-mutated breast cancer. <i>Annals of Oncology</i> , 2018, 29, 1203-1210.	0.6	280
7	Eph-ephrin-B interactions suppress colorectal cancer progression by compartmentalizing tumor cells. <i>Nature Genetics</i> , 2007, 39, 1376-1383.	9.4	242
8	A cysteine-sulfinic acid in peroxiredoxin regulates H ₂ O ₂ -sensing by the antioxidant Pap1 pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 8875-8880.	3.3	230
9	Circulating tumour cells and cell-free DNA as tools for managing breast cancer. <i>Nature Reviews Clinical Oncology</i> , 2013, 10, 377-389.	12.5	164
10	Concordance of blood- and tumor-based detection of RAS mutations to guide anti-EGFR therapy in metastatic colorectal cancer. <i>Annals of Oncology</i> , 2017, 28, 1294-1301.	0.6	150
11	Tankyrase Inhibition Blocks Wnt/ β -Catenin Pathway and Reverts Resistance to PI3K and AKT Inhibitors in the Treatment of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2016, 22, 644-656.	3.2	143
12	Single-cell transcriptome conservation in cryopreserved cells and tissues. <i>Genome Biology</i> , 2017, 18, 45.	3.8	134
13	Defining a minimal cell: essentiality of small ORFs and ncRNAs in a genome-reduced bacterium. <i>Molecular Systems Biology</i> , 2015, 11, 780.	3.2	133
14	Molecular Diagnosis of Diffuse Gliomas through Sequencing of Cell-Free Circulating Tumor DNA from Cerebrospinal Fluid. <i>Clinical Cancer Research</i> , 2018, 24, 2812-2819.	3.2	128
15	Lifespan extension by calorie restriction relies on the Sty1 MAP kinase stress pathway. <i>EMBO Journal</i> , 2010, 29, 981-991.	3.5	108
16	Activation of the redox sensor Pap1 by hydrogen peroxide requires modulation of the intracellular oxidant concentration. <i>Molecular Microbiology</i> , 2004, 52, 1427-1435.	1.2	104
17	Oxidative stress in <i>Schizosaccharomyces pombe</i> : different H ₂ O ₂ levels, different response pathways. <i>Molecular Genetics and Genomics</i> , 2006, 276, 495-502.	1.0	98
18	Incorporating BEAMing technology as a liquid biopsy into clinical practice for the management of colorectal cancer patients: an expert taskforce review. <i>Annals of Oncology</i> , 2017, 28, 2943-2949.	0.6	89

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19	A Personalized Preclinical Model to Evaluate the Metastatic Potential of Patient-Derived Colon Cancer Initiating Cells. <i>Clinical Cancer Research</i> , 2013, 19, 6787-6801.	3.2	80
20	TET2 controls chemoresistant slow-cycling cancer cell survival and tumor recurrence. <i>Journal of Clinical Investigation</i> , 2018, 128, 3887-3905.	3.9	79
21	Molecular Profiling of Patients with Colorectal Cancer and Matched Targeted Therapy in Phase I Clinical Trials. <i>Molecular Cancer Therapeutics</i> , 2012, 11, 2062-2071.	1.9	77
22	Resistance to Targeted Therapies in Renal Cancer: The Importance of Changing the Mechanism of Action. <i>Targeted Oncology</i> , 2017, 12, 19-35.	1.7	77
23	Strand-specific deep sequencing of the transcriptome. <i>Genome Research</i> , 2010, 20, 989-999.	2.4	76
24	Molecular prescreening to select patient population in early clinical trials. <i>Nature Reviews Clinical Oncology</i> , 2012, 9, 359-366.	12.5	68
25	Prospective multicenter real-world RAS mutation comparison between OncoBEAM-based liquid biopsy and tissue analysis in metastatic colorectal cancer. <i>British Journal of Cancer</i> , 2018, 119, 1464-1470.	2.9	62
26	DualMET andERBB inhibition overcomes intratumor plasticity in osimertinib-resistant-advanced non-small-cell lung cancer (NSCLC). <i>Annals of Oncology</i> , 2017, 28, 2451-2457.	0.6	58
27	The Glycolytic Metabolite Methylglyoxal Activates Pap1 and Sty1 Stress Responses in <i>Schizosaccharomyces pombe</i> . <i>Journal of Biological Chemistry</i> , 2005, 280, 36708-36713.	1.6	57
28	Support systems to guide clinical decision-making in precision oncology: The Cancer Core Europe Molecular Tumor Board Portal. <i>Nature Medicine</i> , 2020, 26, 992-994.	15.2	56
29	The Peroxiredoxin Tpx1 Is Essential as a H ₂ O ₂ Scavenger during Aerobic Growth in Fission Yeast. <i>Molecular Biology of the Cell</i> , 2007, 18, 2288-2295.	0.9	54
30	Multigene panel testing beyond BRCA1/2 in breast/ovarian cancer Spanish families and clinical actionability of findings. <i>Journal of Cancer Research and Clinical Oncology</i> , 2018, 144, 2495-2513.	1.2	53
31	Genomic Analyses across Six Cancer Types Identify Basal-like Breast Cancer as a Unique Molecular Entity. <i>Scientific Reports</i> , 2013, 3, 3544.	1.6	45
32	Early evolutionary divergence between papillary and anaplastic thyroid cancers. <i>Annals of Oncology</i> , 2018, 29, 1454-1460.	0.6	44
33	The Molecular Tumor Board Portal supports clinical decisions and automated reporting for precision oncology. <i>Nature Cancer</i> , 2022, 3, 251-261.	5.7	44
34	Clinical Response to a Lapatinib-Based Therapy for a Li-Fraumeni Syndrome Patient with a Novel <i>HER2</i> V659E Mutation. <i>Cancer Discovery</i> , 2013, 3, 1238-1244.	7.7	43
35	Multicenter Phase II Study of Lurbinectedin in <i>BRCA</i> -Mutated and Unselected Metastatic Advanced Breast Cancer and Biomarker Assessment Substudy. <i>Journal of Clinical Oncology</i> , 2018, 36, 3134-3143.	0.8	43
36	Transcription start site associated RNAs in bacteria. <i>Molecular Systems Biology</i> , 2012, 8, 585.	3.2	40

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37	A Mouse Model Uncovers LKB1 as an UVB-Induced DNA Damage Sensor Mediating CDKN1A (p21WAF1/CIP1) Degradation. <i>PLoS Genetics</i> , 2014, 10, e1004721.	1.5	40
38	Impact of circulating tumor DNA mutant allele fraction on prognosis in <i>RAS</i> mutant metastatic colorectal cancer. <i>Molecular Oncology</i> , 2019, 13, 1827-1835.	2.1	40
39	Combined Analysis of Concordance between Liquid and Tumor Tissue Biopsies for <i>RAS</i> Mutations in Colorectal Cancer with a Single Metastasis Site: The METABEAM Study. <i>Clinical Cancer Research</i> , 2021, 27, 2515-2522.	3.2	39
40	Reprogramming activity of NANOGP8, a NANOG family member widely expressed in cancer. <i>Oncogene</i> , 2014, 33, 2513-2519.	2.6	37
41	<i>Schizosaccharomyces pombe</i> Cells Lacking the Ran-binding Protein Hba1 Show a Multidrug Resistance Phenotype Due to Constitutive Nuclear Accumulation of Pap1. <i>Journal of Biological Chemistry</i> , 2003, 278, 40565-40572.	1.6	36
42	Comparison of the Clinical Sensitivity of the Idylla Platform and the OncoBEAM RAS CRC Assay for KRAS Mutation Detection in Liquid Biopsy Samples. <i>Scientific Reports</i> , 2019, 9, 8976.	1.6	34
43	Microarray and deep sequencing cross-platform analysis of the mirRNome and isomiR variation in response to epidermal growth factor. <i>BMC Genomics</i> , 2013, 14, 371.	1.2	33
44	Clinical value of next generation sequencing of plasma cell-free DNA in gastrointestinal stromal tumors. <i>BMC Cancer</i> , 2020, 20, 99.	1.1	31
45	Epigenetic <i>EGFR</i> Gene Repression Confers Sensitivity to Therapeutic BRAFV600E Blockade in Colon Neuroendocrine Carcinomas. <i>Clinical Cancer Research</i> , 2020, 26, 902-909.	3.2	29
46	Analysis of mutant allele fractions in driver genes in colorectal cancer – biological and clinical insights. <i>Molecular Oncology</i> , 2017, 11, 1263-1272.	2.1	26
47	Regional and subtype-dependent miRNA signatures in sporadic Creutzfeldt-Jakob disease are accompanied by alterations in miRNA silencing machinery and biogenesis. <i>PLoS Pathogens</i> , 2018, 14, e1006802.	2.1	26
48	An evaluation of the challenges to developing tumor BRCA1 and BRCA2 testing methodologies for clinical practice. <i>Human Mutation</i> , 2018, 39, 394-405.	1.1	24
49	Molecular profiling of long-term responders to immune checkpoint inhibitors in advanced non-small cell lung cancer. <i>Molecular Oncology</i> , 2021, 15, 887-900.	2.1	24
50	New approach to cancer therapy based on a molecularly defined cancer classification. <i>Ca-A Cancer Journal for Clinicians</i> , 2014, 64, 70-74.	157.7	22
51	C-terminal truncation of the peroxiredoxin Tpx1 decreases its sensitivity for hydrogen peroxide without compromising its role in signal transduction. <i>Genes To Cells</i> , 2008, 13, 171-179.	0.5	19
52	Multiple platform assessment of the EGF dependent transcriptome by microarray and deep tag sequencing analysis. <i>BMC Genomics</i> , 2011, 12, 326.	1.2	19
53	Pancreatic cancer heterogeneity and response to Mek inhibition. <i>Oncogene</i> , 2017, 36, 5639-5647.	2.6	19
54	MEK plus PI3K/mTORC1/2 Therapeutic Efficacy Is Impacted by <i>TP53</i> Mutation in Preclinical Models of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 5499-5510.	3.2	18

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55	Genetic evolution of nevus of Ota reveals clonal heterogeneity acquiring <i>BAP1</i> and <i>TP53</i> mutations. <i>Pigment Cell and Melanoma Research</i> , 2016, 29, 247-253.	1.5	18
56	Identification of somatic gene mutations in penile squamous cell carcinoma. <i>Genes Chromosomes and Cancer</i> , 2015, 54, 629-637.	1.5	17
57	Limitations in predicting PAM50 intrinsic subtype and risk of relapse score with Ki67 in estrogen receptor-positive HER2-negative breast cancer. <i>Oncotarget</i> , 2017, 8, 21930-21937.	0.8	17
58	Partial short-read sequencing of a highly inbred Iberian pig and genomics inference thereof. <i>Heredity</i> , 2011, 107, 256-264.	1.2	16
59	Activity of HSP90 Inhibitor in a Metastatic Lung Cancer Patient With a Germline BRCA1 Mutation. <i>Journal of the National Cancer Institute</i> , 2018, 110, 914-917.	3.0	16
60	Genetic profile of <i>GNAQ</i> -mutated blue melanocytic neoplasms reveals mutations in genes linked to genomic instability and the PI3K pathway. <i>Oncotarget</i> , 2016, 7, 28086-28095.	0.8	16
61	The homeoprotein SIX1 controls cellular senescence through the regulation of p16INK4A and differentiation-related genes. <i>Oncogene</i> , 2016, 35, 3485-3494.	2.6	15
62	Targeted multiplex proteomics for molecular prescreening and biomarker discovery in metastatic colorectal cancer. <i>Scientific Reports</i> , 2019, 9, 13568.	1.6	14
63	Biomarker Analysis of the Phase III NALA Study of Neratinib + Capecitabine versus Lapatinib + Capecitabine in Patients with Previously Treated Metastatic Breast Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 5818-5827.	3.2	14
64	Spatiotemporal Control of Forkhead Binding to DNA Regulates the Meiotic Gene Expression Program. <i>Cell Reports</i> , 2016, 14, 885-895.	2.9	12
65	High <i>FGFR1</i> mRNA Expression Levels Correlate with Response to Selective FGFR Inhibitors in Breast Cancer. <i>Clinical Cancer Research</i> , 2022, 28, 137-149.	3.2	12
66	Determination of somatic oncogenic mutations linked to target-based therapies using MassARRAY technology. <i>Oncotarget</i> , 2016, 7, 22543-22555.	0.8	11
67	Evolving Landscape of Molecular Prescreening Strategies for Oncology Early Clinical Trials. <i>JCO Precision Oncology</i> , 2020, 4, 505-513.	1.5	10
68	Genomic Biomarkers and Genome-Wide Loss-of-Heterozygosity Scores in Metastatic Prostate Cancer Following Progression on Androgen-Targeting Therapies. <i>JCO Precision Oncology</i> , 2022, , .	1.5	10
69	Genomic heterogeneity and efficacy of PI3K pathway inhibitors in patients with gynaecological cancer. <i>ESMO Open</i> , 2019, 4, e000444.	2.0	8
70	Identification of Expression Profiles Defining Distinct Prognostic Subsets of Radioactive-Iodine Refractory Differentiated Thyroid Cancer from the DECISION Trial. <i>Molecular Cancer Therapeutics</i> , 2020, 19, 312-317.	1.9	8
71	Long duration of immunotherapy in a <i>STK11</i> mutated/ <i>KRAS</i> wild-type non-small cell lung cancer patient. <i>Pulmonology</i> , 2020, 26, 49-50.	1.0	8
72	Clinicopathological and Molecular Characterization of Metastatic Gastrointestinal Stromal Tumors with Prolonged Benefit to Frontline Imatinib. <i>Oncologist</i> , 2019, 24, 680-687.	1.9	7

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73	Concordance of genomic alterations (GA) in synchronous tumor biopsies (tBx) and circulating tumor (ct) DNA from metastatic breast cancer (MBC) patients (pts).. Journal of Clinical Oncology, 2018, 36, 1073-1073.	0.8	7
74	687 Beta-catenin Confers Resistance to PI3K and AKT Inhibitors and Subverts FOXO3a to Promote Metastasis in Colon Cancer. European Journal of Cancer, 2012, 48, S162.	1.3	5
75	Circulating cell-free DNA as predictor of treatment failure after neoadjuvant chemoradiotherapy before surgery in patients with locally advanced rectal cancer: is it ready for primetime?. Annals of Oncology, 2018, 29, 532-534.	0.6	5
76	Case Report: A Case Study Documenting the Activity of Atezolizumab in a PD-L1-Negative Triple-Negative Breast Cancer. Frontiers in Oncology, 2021, 11, 710596.	1.3	5
77	High levels of chromosomal aberrations negatively associate with benefit to checkpoint inhibition in NSCLC. , 2022, 10, e004197.		5
78	Genetic evolution to tyrosine kinase inhibitory therapy in patients with EGFR-mutated non-small-cell lung cancer. British Journal of Cancer, 2021, 125, 1561-1569.	2.9	4
79	Comprehensive profiling of biliary tract cancers (BTC) to reveal molecular heterogeneity with implications for matched targeted therapies (MTT).. Journal of Clinical Oncology, 2016, 34, 4085-4085.	0.8	3
80	Clonality of PIK3CA mutations (mut) and efficacy of PI3K/AKT/mTOR inhibitors (PAMi) in patients (pts) with metastatic breast cancer (MBC).. Journal of Clinical Oncology, 2016, 34, 528-528.	0.8	3
81	First Nationwide Molecular Screening Program in Spain for Patients With Advanced Breast Cancer: Results From the AGATA SOLTI-1301 Study. Frontiers in Oncology, 2021, 11, 744112.	1.3	3
82	Genetic Profiles of Squamous Cell Carcinomas Associated with Recessive Dystrophic Epidermolysis Bullosa Unveil NOTCH and TP53 Mutations and an Increased MYC Expression. Journal of Investigative Dermatology, 2018, 138, 1423-1427.	0.3	2
83	Matching degree between PI3K/AKT/mTOR (PAM) pathway mutations (mut) and therapy (ttx) as predictor of clinical benefit (ClinBen) in early trials.. Journal of Clinical Oncology, 2016, 34, 2572-2572.	0.8	2
84	Prognostic impact of primary tumor site location in metastatic colorectal cancer (mCRC).. Journal of Clinical Oncology, 2016, 34, 578-578.	0.8	2
85	Abstract 930: Analysis of cell-free tumor DNA in cerebrospinal fluid to characterize and monitor the genetic alterations of brain tumors. Cancer Research, 2015, 75, 930-930.	0.4	2
86	Molecular, clinical and prognostic characterization of double KRAS/PIK3CA (dKP) mutated metastatic colorectal cancer (mCRC). Annals of Oncology, 2016, 27, vi185.	0.6	1
87	Molecular screening programmes for precision medicine: lessons learned from personalized medicine trials. Expert Review of Precision Medicine and Drug Development, 2016, 1, 419-430.	0.4	1
88	RNF43- and NOTCH1-Mutated Chemotherapy and Anti-EGFR-Refractory Colorectal Cancer: Should Clonality Guide Target Prioritization With Investigational Therapies?. JCO Precision Oncology, 2019, 3, 1-3.	1.5	1
89	Outcome evolution of matched molecular targeted agents (MTAs) in metastatic colorectal cancer (CRC) patients (pts): VHIO experience.. Journal of Clinical Oncology, 2015, 33, 3602-3602.	0.8	1
90	Impact of molecular prescreening for genomically-guided trials in head and neck cancer (HNC).. Journal of Clinical Oncology, 2016, 34, 6030-6030.	0.8	1

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91	Survival determinants with matched targeted therapies in BRAF mutant metastatic colorectal cancer (mCRC).. Journal of Clinical Oncology, 2016, 34, 649-649.	0.8	1
92	Analysis of <i>RSPO</i> gene expression in solid tumors.. Journal of Clinical Oncology, 2016, 34, e23235-e23235.	0.8	1
93	RNAseq analysis of the sorafenib phase III DECISION trial in differentiated thyroid cancer (DTC): Correlation with clinical outcome.. Journal of Clinical Oncology, 2017, 35, 6083-6083.	0.8	1
94	2399 Impact of KRAS mutations on clinical outcomes in advanced refractory pancreatic cancer patients. European Journal of Cancer, 2015, 51, S470.	1.3	0
95	Mutational load by targeted next generation sequencing (NGS) panels as potential biomarker of response to checkpoint inhibitors. Annals of Oncology, 2015, 26, viii5.	0.6	0
96	AGATA molecular screening program: implementing precision medicine in patients with advanced breast cancer in Spain. Annals of Oncology, 2015, 26, ii21.	0.6	0
97	Prognostic impact of KRAS mutation in metastatic (met) pancreatic cancer patients (pts). Annals of Oncology, 2016, 27, vi232.	0.6	0
98	Multiple primary cancers (MPC) in a series of lung cancer (LC) patient: Incidence and outcome. Annals of Oncology, 2018, 29, viii641.	0.6	0
99	Evaluation of the sensitivity of RAS mutation detection of the Idylla platform in comparison to the OncoBEAM RAS CRC assay. Annals of Oncology, 2018, 29, viii184.	0.6	0
100	Primary results of the first nationwide molecular screening program in Spain for patients with advanced breast cancer (AGATA SOLTI-1301 study). Annals of Oncology, 2018, 29, viii90.	0.6	0
101	MYC Copy Number Detection in Clinical Samples Using a Digital DNA-Hybridization and Detection Method. Methods in Molecular Biology, 2021, 2318, 321-336.	0.4	0
102	Prognostic implications of phosphatidylinositol 3-kinase (PI3K) pathway alterations in metastatic triple-negative breast cancer (mTNBC).. Journal of Clinical Oncology, 2011, 29, 1081-1081.	0.8	0
103	Correlation between molecular features (MFs) and clinical results in endometrial cancer (EC): A single-institution experience.. Journal of Clinical Oncology, 2011, 29, e15572-e15572.	0.8	0
104	Abstract 683: Evaluation of PTEN and PIK3CA status in breast cancer for patient selection: Cross-validation between institutions. , 2012, , .		0
105	Abstract C114: Clinical response to a lapatinib-based therapy of a Li-Fraumeni Syndrome patient with a novel HER2-V659E mutation.. , 2013, , .		0
106	Coexisting KRAS and PIK3CA exon 20 mutations as a potential poor-prognosis factor in metastatic colorectal cancer (mCRC).. Journal of Clinical Oncology, 2014, 32, 3591-3591.	0.8	0
107	Incorporation of FGFR1 and FGFR2 amplification status determination in routine molecular prescreening for targeted therapies.. Journal of Clinical Oncology, 2014, 32, 11105-11105.	0.8	0
108	Clinical and molecular characterization of refractory BRAF mutant metastatic colorectal carcinoma (mCRC): Vall d'Hebron Institute of Oncology phase I program cohort.. Journal of Clinical Oncology, 2015, 33, 587-587.	0.8	0

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109	Early drug development in advanced gynecologic cancer based on genetic tumor profiling.. Journal of Clinical Oncology, 2015, 33, 5562-5562.	0.8	0
110	Knocking on molecular alterations in advanced gastric cancer (AGC).. Journal of Clinical Oncology, 2015, 33, 4063-4063.	0.8	0
111	Abstract LB-123: Analysis of cell-free plasma DNA (cfDNA) identifies 3 molecular subtypes of acquired resistance to AZD9291, a novel EGFR tyrosine kinase inhibitor (TKI), in patients (pts) with advanced lung cancer. , 2015, , .		0
112	Molecular prescreening (MP) to treat patients (pts) with advanced pancreatic cancer (PC) in early clinical trials.. Journal of Clinical Oncology, 2016, 34, 272-272.	0.8	0
113	Abstract P2-08-13: Integrating multiplex and next generation sequencing (NGS) platforms in routine molecular profiling of metastatic breast cancer (MBC) patients (pts): Trends for enrollment in genotype-directed clinical trials (GDTs). , 2016, , .		0
114	Outcome of RAS mutated lung adenocarcinoma (ADC) patients (pts) on standard chemotherapy (chemo) and immune checkpoint inhibitors (immuneCI).. Journal of Clinical Oncology, 2016, 34, e20597-e20597.	0.8	0
115	Association of response to programmed death 1 receptor or ligand (PD1/PDL1) blockade with immune-related gene expression profiling across three cancer-types.. Journal of Clinical Oncology, 2016, 34, 3038-3038.	0.8	0
116	Clonality patterns of driver mutations (mut) to reveal spatial-temporal genomic heterogeneity in colorectal cancer (CRC).. Journal of Clinical Oncology, 2016, 34, 3509-3509.	0.8	0
117	Outcome of <i>KRAS</i> mutated (m) non-small cell lung cancer (NSCLC) patients (pts) treated with immune checkpoint inhibitors (immuneCI).. Journal of Clinical Oncology, 2017, 35, e20526-e20526.	0.8	0
118	Molecular sequencing and gene fusion detection in non-small cell lung cancer (NSCLC) patients: Impact of co-existing alterations.. Journal of Clinical Oncology, 2017, 35, e23103-e23103.	0.8	0
119	Abstract B075: Evolving molecular prescreening program to identify genomic alterations in the NOTCH pathway. , 2018, , .		0
120	Abstract 2953: Adapting a molecular prescreening program to detect notch pathway alterations in the context of early drug development. , 2020, , .		0
121	Abstract 2488: Characterization of gene fusions in paired primary and metastatic samples of breast cancer in the AURORA molecular screening program. , 2020, , .		0