

Valsamo Anagnostou

List of Publications by Year in descending order

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Version: 2024-02-01

54
papers

8,883
citations

126907

33
h-index

206112

48
g-index

57
all docs

57
docs citations

57
times ranked

15221
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Neoadjuvant PD-1 Blockade in Resectable Lung Cancer. <i>New England Journal of Medicine</i> , 2018, 378, 1976-1986. | 27.0 | 1,495 |
| 2 | Direct detection of early-stage cancers using circulating tumor DNA. <i>Science Translational Medicine</i> , 2017, 9, . | 12.4 | 808 |
| 3 | Genome-wide cell-free DNA fragmentation in patients with cancer. <i>Nature</i> , 2019, 570, 385-389. | 27.8 | 764 |
| 4 | Evolution of Neoantigen Landscape during Immune Checkpoint Blockade in Nonâ€“Small Cell Lung Cancer. <i>Cancer Discovery</i> , 2017, 7, 264-276. | 9.4 | 706 |
| 5 | Antibody validation. <i>BioTechniques</i> , 2010, 48, 197-209. | 1.8 | 548 |
| 6 | The genomic landscape of response to EGFR blockade in colorectal cancer. <i>Nature</i> , 2015, 526, 263-267. | 27.8 | 398 |
| 7 | Clinical implications of genomic alterations in the tumour and circulation of pancreatic cancer patients. <i>Nature Communications</i> , 2015, 6, 7686. | 12.8 | 393 |
| 8 | Epigenetic Therapy Ties MYC Depletion to Reversing Immune Evasion and Treating Lung Cancer. <i>Cell</i> , 2017, 171, 1284-1300.e21. | 28.9 | 366 |
| 9 | Personalized genomic analyses for cancer mutation discovery and interpretation. <i>Science Translational Medicine</i> , 2015, 7, 283ra53. | 12.4 | 347 |
| 10 | GOLPH3 modulates mTOR signalling and rapamycin sensitivity in cancer. <i>Nature</i> , 2009, 459, 1085-1090. | 27.8 | 311 |
| 11 | Transcriptional programs of neoantigen-specific TIL in anti-PD-1-treated lung cancers. <i>Nature</i> , 2021, 596, 126-132. | 27.8 | 234 |
| 12 | Dynamics of Tumor and Immune Responses during Immune Checkpoint Blockade in Nonâ€“Small Cell Lung Cancer. <i>Cancer Research</i> , 2019, 79, 1214-1225. | 0.9 | 226 |
| 13 | Conserved Interferon- γ Signaling Drives Clinical Response to Immune Checkpoint Blockade Therapy in Melanoma. <i>Cancer Cell</i> , 2020, 38, 500-515.e3. | 16.8 | 203 |
| 14 | Detection and characterization of lung cancer using cell-free DNA fragmentomes. <i>Nature Communications</i> , 2021, 12, 5060. | 12.8 | 161 |
| 15 | White blood cell and cell-free DNA analyses for detection of residual disease in gastric cancer. <i>Nature Communications</i> , 2020, 11, 525. | 12.8 | 158 |
| 16 | Multimodal genomic features predict outcome of immune checkpoint blockade in non-small-cell lung cancer. <i>Nature Cancer</i> , 2020, 1, 99-111. | 13.2 | 141 |
| 17 | Ipilimumab plus nivolumab and DNA-repair defects in AR-V7-expressing metastatic prostate cancer. <i>Oncotarget</i> , 2018, 9, 28561-28571. | 1.8 | 129 |
| 18 | The Mutation-Associated Neoantigen Functional Expansion of Specific T Cells (MANAFEST) Assay: A Sensitive Platform for Monitoring Antitumor Immunity. <i>Cancer Immunology Research</i> , 2018, 6, 888-899. | 3.4 | 118 |

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|----|---|------|-----------|
| 19 | Association of High Tumor Mutation Burden in Non-Small Cell Lung Cancers With Increased Immune Infiltration and Improved Clinical Outcomes of PD-L1 Blockade Across PD-L1 Expression Levels. <i>JAMA Oncology</i> , 2022, 8, 1160. | 7.1 | 117 |
| 20 | Immuno-oncology Trial Endpoints: Capturing Clinically Meaningful Activity. <i>Clinical Cancer Research</i> , 2017, 23, 4959-4969. | 7.0 | 115 |
| 21 | Neoadjuvant nivolumab plus ipilimumab in resectable non-small cell lung cancer. , 2020, 8, e001282. | | 108 |
| 22 | High-Throughput Prediction of MHC Class I and II Neoantigens with MHCnuggets. <i>Cancer Immunology Research</i> , 2020, 8, 396-408. | 3.4 | 103 |
| 23 | The alveolar immune cell landscape is dysregulated in checkpoint inhibitor pneumonitis. <i>Journal of Clinical Investigation</i> , 2019, 129, 4305-4315. | 8.2 | 100 |
| 24 | Compartmental Analysis of T-cell Clonal Dynamics as a Function of Pathologic Response to Neoadjuvant PD-1 Blockade in Resectable Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 1327-1337. | 7.0 | 90 |
| 25 | A machine learning approach for somatic mutation discovery. <i>Science Translational Medicine</i> , 2018, 10, . | 12.4 | 80 |
| 26 | Early Noninvasive Detection of Response to Targeted Therapy in Non-Small Cell Lung Cancer. <i>Cancer Research</i> , 2019, 79, 1204-1213. | 0.9 | 75 |
| 27 | Durvalumab with platinum-pemetrexed for unresectable pleural mesothelioma: survival, genomic and immunologic analyses from the phase 2 PrE0505 trial. <i>Nature Medicine</i> , 2021, 27, 1910-1920. | 30.7 | 62 |
| 28 | Multi-Level Targeting of the Phosphatidylinositol-3-Kinase Pathway in Non-Small Cell Lung Cancer Cells. <i>PLoS ONE</i> , 2012, 7, e31331. | 2.5 | 55 |
| 29 | Preanalytical variables and phosphoepitope expression in FFPE tissue: quantitative epitope assessment after variable cold ischemic time. <i>Laboratory Investigation</i> , 2015, 95, 334-341. | 3.7 | 52 |
| 30 | A tissue quality index: an intrinsic control for measurement of effects of preanalytical variables on FFPE tissue. <i>Laboratory Investigation</i> , 2014, 94, 467-474. | 3.7 | 48 |
| 31 | The status of tumor mutational burden and immunotherapy. <i>Nature Cancer</i> , 2022, 3, 652-656. | 13.2 | 48 |
| 32 | Estrogen receptor co-activator (AIB1) protein expression by automated quantitative analysis (AQUA) in a breast cancer tissue microarray and association with patient outcome. <i>Breast Cancer Research and Treatment</i> , 2009, 115, 77-85. | 2.5 | 45 |
| 33 | Integrative Tumor and Immune Cell Multi-omic Analyses Predict Response to Immune Checkpoint Blockade in Melanoma. <i>Cell Reports Medicine</i> , 2020, 1, 100139. | 6.5 | 45 |
| 34 | Persistent mutant oncogene specific T cells in two patients benefitting from anti-PD-1. , 2019, 7, 40. | | 42 |
| 35 | Measurement of Aldehyde Dehydrogenase 1 Expression Defines a Group with Better Prognosis in Patients with Non-Small Cell Lung Cancer. <i>American Journal of Pathology</i> , 2012, 181, 1436-1442. | 3.8 | 41 |
| 36 | Peripheral blood immune cell dynamics reflect antitumor immune responses and predict clinical response to immunotherapy. , 2022, 10, e004688. | | 34 |

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|----|--|------|-----------|
| 37 | Standardization of Epidermal Growth Factor Receptor (EGFR) Measurement by Quantitative Immunofluorescence and Impact on Antibody-Based Mutation Detection in Non-Small Cell Lung Cancer. <i>American Journal of Pathology</i> , 2011, 179, 580-589. | 3.8 | 21 |
| 38 | Ontogeny of intrinsic innervation in the human kidney. <i>Anatomy and Embryology</i> , 2004, 209, 41-47. | 1.5 | 17 |
| 39 | Visual storytelling enhances knowledge dissemination in biomedical science. <i>Journal of Biomedical Informatics</i> , 2020, 107, 103458. | 4.3 | 14 |
| 40 | Soluble triggering receptor expressed on myeloid cells-1 (sTREM-1) detection in cancer patients: a prognostic marker for lung metastases from solid malignancies. <i>Anticancer Research</i> , 2008, 28, 1411-5. | 1.1 | 11 |
| 41 | Translating noninvasive molecular responses into clinical reality for cancer immunotherapy. <i>Nature Reviews Clinical Oncology</i> , 2021, 18, 65-66. | 27.6 | 9 |
| 42 | Protocol of DREAM3R: Durvalumab with chemotherapy as first-line treatment in advanced pleural mesothelioma—a phase 3 randomised trial. <i>BMJ Open</i> , 2022, 12, e057663. | 1.9 | 9 |
| 43 | Developing a multivariable prognostic model for pancreatic endocrine tumors using the clinical data warehouse resources of a single institution. <i>Applied Clinical Informatics</i> , 2010, 01, 38-49. | 1.7 | 6 |
| 44 | Genetic variation in antigen presentation and cancer immunotherapy. <i>Immunity</i> , 2022, 55, 3-6. | 14.3 | 5 |
| 45 | Multicenter phase II study of neoadjuvant nivolumab or nivolumab plus relatlimab (anti-LAG3) in resectable non-small cell lung carcinoma. <i>Journal of Clinical Oncology</i> , 2022, 40, 321-321. | 1.6 | 5 |
| 46 | An Evaluation of Pretrained BERT Models for Comparing Semantic Similarity Across Unstructured Clinical Trial Texts. <i>Studies in Health Technology and Informatics</i> , 2022, 289, 18-21. | 0.3 | 4 |
| 47 | Epithelioid haemangioendothelioma of the lung presenting with pulmonary nocardiosis. <i>In Vivo</i> , 2007, 21, 1123-6. | 1.3 | 4 |
| 48 | Primary parotid adenocarcinoma metastasis to the spleen with mutation: cytological findings and review of the literature. <i>International Journal of Clinical and Experimental Pathology</i> , 2017, 10, 5999-6005. | 0.5 | 2 |
| 49 | Artificial Intelligence-Assisted Serial Analysis of Clinical Cancer Genomics Data Identifies Changing Treatment Recommendations and Therapeutic Targets. <i>Clinical Cancer Research</i> , 2022, 28, 2361-2372. | 7.0 | 2 |
| 50 | Comprehensive modeling of longitudinal circulating tumor DNA dynamics to predict clinical response to first-line immunotherapy and chemoimmunotherapy in advanced non-small cell lung cancer. <i>Journal of Clinical Oncology</i> , 2020, 38, 9525-9525. | 1.6 | 1 |
| 51 | Immunogenomic features of pathologic response to neoadjuvant immune checkpoint blockade in esophageal cancer. <i>Journal of Clinical Oncology</i> , 2021, 39, 4042-4042. | 1.6 | 0 |
| 52 | Multicenter phase II study of abemaciclib and ramucirumab in metastatic esophageal/gastroesophageal junction carcinoma. <i>Journal of Clinical Oncology</i> , 2022, 40, TPS4169-TPS4169. | 1.6 | 0 |
| 53 | DREAM3R: Durvalumab with chemotherapy as first-line treatment in advanced pleural mesothelioma—a phase 3 randomized trial. <i>Journal of Clinical Oncology</i> , 2022, 40, TPS8599-TPS8599. | 1.6 | 0 |
| 54 | Natural Language Processing Approaches for Retrieval of Clinically Relevant Genomic Information in Cancer. <i>Studies in Health Technology and Informatics</i> , 2022, , . | 0.3 | 0 |