Muhammed Murtaza

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Feasibility of circulating tumor DNA analysis in dogs with naturally occurring malignant and benign splenic lesions. Scientific Reports, 2022, 12, 6337.	3.3	3
2	Analysis of recurrently protected genomic regions in cell-free DNA found in urine. Science Translational Medicine, 2021, 13, .	12.4	40
3	Horizons in Veterinary Precision Oncology: Fundamentals of Cancer Genomics and Applications of Liquid Biopsy for the Detection, Characterization, and Management of Cancer in Dogs. Frontiers in Veterinary Science, 2021, 8, 664718.	2.2	21
4	Plasma metagenomic sequencing to detect and quantify bacterial DNA in ICU patients suspected of sepsis. Journal of Trauma and Acute Care Surgery, 2021, Publish Ahead of Print, 988-994.	2.1	2
5	Circulating tumor DNA as an early cancer detection tool. , 2020, 207, 107458.		123
6	Harmonizing Cell-Free DNA Collection and Processing Practices through Evidence-Based Guidance. Clinical Cancer Research, 2020, 26, 3104-3109.	7.0	66
7	Personalized circulating tumor DNA analysis to detect residual disease after neoadjuvant therapy in breast cancer. Science Translational Medicine, 2019, 11, .	12.4	197
8	Identification of Recurrent Activating <i>HER2</i> Mutations in Primary Canine Pulmonary Adenocarcinoma. Clinical Cancer Research, 2019, 25, 5866-5877.	7.0	24
9	The Genomic and Immune Landscapes of Lethal Metastatic Breast Cancer. Cell Reports, 2019, 27, 2690-2708.e10.	6.4	95
10	Circulating Tumor DNA Analysis and Opportunities for Personalized Cancer Medicine. , 2019, , 229-242.		0
11	Dynamics of multiple resistance mechanisms in plasma DNA during EGFRâ€ŧargeted therapies in nonâ€small cell lung cancer. EMBO Molecular Medicine, 2018, 10, .	6.9	61
12	Evaluation of pre-analytical factors affecting plasma DNA analysis. Scientific Reports, 2018, 8, 7375.	3.3	102
13	Detection of copy number aberrations in cholangiocarcinoma using shallow whole genome sequencing of plasma DNA Journal of Clinical Oncology, 2018, 36, 293-293.	1.6	3
14	Influence of a single nucleotide polymorphism (SNP) and DNA hybridization on the drying patterns of micro droplets. Journal of Nanomedicine, 2018, 1, .	0.3	0
15	Capturing tumor heterogeneity and clonal evolution in solid cancers using circulating tumor DNA analysis. , 2017, 174, 22-26.		28
16	Nucleosome mapping in plasma DNA predicts cancer gene expression. Nature Genetics, 2016, 48, 1105-1106.	21.4	19
17	Long-tunneled versus short-tunneled external ventricular drainage: Prospective experience from a developing country. Journal of Innovative Optical Health Sciences, 2016, 11, 114-117.	1.0	14
18	Spatial and Temporal Heterogeneity in High-Grade Serous Ovarian Cancer: A Phylogenetic Analysis. PLoS Medicine, 2015, 12, e1001789.	8.4	314

2

MUHAMMED MURTAZA

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19	Multifocal clonal evolution characterized using circulating tumour DNA in a case of metastatic breast cancer. Nature Communications, 2015, 6, 8760.	12.8	409
20	Ovarian Cancer Cell Line Panel (OCCP): Clinical Importance of In Vitro Morphological Subtypes. PLoS ONE, 2014, 9, e103988.	2.5	319
21	Clinical and pathological impact of <i>VHL, PBRM1, BAP1, SETD2, KDM6A</i> , and <i>JARID1c</i> in clear cell carcinoma. Genes Chromosomes and Cancer, 2014, 53, 38-51.	2.8	107
22	Frequency and Determinants of Intracranial Atherosclerotic Stroke in Urban Pakistan. Journal of Stroke and Cerebrovascular Diseases, 2014, 23, 2174-2182.	1.6	6
23	Ordering of mutations in preinvasive disease stages of esophageal carcinogenesis. Nature Genetics, 2014, 46, 837-843.	21.4	302
24	Analysis of Circulating Tumor DNA to Monitor Metastatic Breast Cancer. New England Journal of Medicine, 2013, 368, 1199-1209.	27.0	1,884
25	Non-invasive analysis of acquired resistance to cancer therapy by sequencing of plasma DNA. Nature, 2013, 497, 108-112.	27.8	1,443
26	Noninvasive Identification and Monitoring of Cancer Mutations by Targeted Deep Sequencing of Plasma DNA. Science Translational Medicine, 2012, 4, 136ra68.	12.4	1,086
27	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. American Journal of Human Genetics, 2012, 90, 1116-1117.	6.2	0
28	Risk Factor Profiles of South Asians with Cerebrovascular Disease. International Journal of Stroke, 2011, 6, 346-348.	5.9	4
29	A genome-wide association study in Europeans and South Asians identifies five new loci for coronary artery disease. Nature Genetics, 2011, 43, 339-344.	21.4	643
30	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. American Journal of Human Genetics, 2011, 88, 6-18.	6.2	122
31	Risk Factors of Stroke in Pakistan: A Dedicated Stroke Clinic Experience. Canadian Journal of Neurological Sciences, 2010, 37, 252-257.	0.5	8
32	Association of the 9p21.3 Locus With Risk of First-Ever Myocardial Infarction in Pakistanis. Arteriosclerosis, Thrombosis, and Vascular Biology, 2010, 30, 1467-1473.	2.4	48
33	Evaluation of therapeutic control in a Pakistani population with hypertension. Journal of Evaluation in Clinical Practice, 2010, 16, 1081-1084.	1.8	7
34	Genetic Determinants of Major Blood Lipids in Pakistanis Compared With Europeans. Circulation: Cardiovascular Genetics, 2010, 3, 348-357.	5.1	25
35	Classification and Clinical Features of Headache Disorders in Pakistan: A Retrospective Review of Clinical Data. PLoS ONE, 2009, 4, e5827.	2.5	31
36	The Karachi intracranial stenosis study (KISS) Protocol: An urban multicenter case-control investigation reporting the clinical, radiologic and biochemical associations of intracranial stenosis in Pakistan. BMC Neurology, 2009, 9, 31.	1.8	7

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37	The burden of stroke and transient ischemic attack in Pakistan: a community-based prevalence study. BMC Neurology, 2009, 9, 58.	1.8	59
38	The Pakistan Risk of Myocardial Infarction Study: a resource for the study of genetic, lifestyle and other determinants of myocardial infarction in South Asia. European Journal of Epidemiology, 2009, 24, 329-338.	5.7	83