

Genetic variants associated with gastrointestinal sympt

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Precision Oncology: Present Status and Perspectives. <i>Current Clinical Pathology</i> , 2017, , 7-26.	0.0	0
2	Genetic variants associated with Fabry disease progression despite enzyme replacement therapy. <i>Oncotarget</i> , 2017, 8, 107558-107564.	1.8	30
3	Non-specific gastrointestinal features: Could it be Fabry disease?. <i>Digestive and Liver Disease</i> , 2018, 50, 429-437.	0.9	28
4	Pharmacogenomic Profiling of ADME Gene Variants: Current Challenges and Validation Perspectives. <i>High-Throughput</i> , 2018, 7, 40.	4.4	37
5	The Era of PARP inhibitors in ovarian cancer: "Class Action" or not? A systematic review and meta-analysis. <i>Critical Reviews in Oncology/Hematology</i> , 2018, 131, 83-89.	4.4	37
6	The potential role of miRNAs in multiple myeloma therapy. <i>Expert Review of Hematology</i> , 2018, 11, 793-803.	2.2	23
7	Polymorphic Variants in <i>NR113</i> and <i>UGT2B7</i> Predict Taxane Neurotoxicity and Have Prognostic Relevance in Patients With Breast Cancer: A Case-Control Study. <i>Clinical Pharmacology and Therapeutics</i> , 2019, 106, 422-431.	4.7	27
8	An exploratory study by DMET array identifies a germline signature associated with imatinib response in gastrointestinal stromal tumor. <i>Pharmacogenomics Journal</i> , 2019, 19, 390-400.	2.0	20
9	Non-coding RNAs in cancer: platforms and strategies for investigating the genomic "dark matter". <i>Journal of Experimental and Clinical Cancer Research</i> , 2020, 39, 117.	8.6	137
10	DMET™ Genotyping: Tools for Biomarkers Discovery in the Era of Precision Medicine. <i>High-Throughput</i> , 2020, 9, 8.	4.4	11
11	Pharmacogenomics Biomarker Discovery and Validation for Translation in Clinical Practice. <i>Clinical and Translational Science</i> , 2021, 14, 113-119.	3.1	42
12	Pharmacogenetics/Pharmacogenomics of Drug-Metabolizing Enzymes and Transporters. , 2021, , .		1
13	Gastrointestinal Involvement in Anderson-Fabry Disease: A Narrative Review. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 3320.	2.6	8
14	Identifying prognostic markers for multiple myeloma through integration and analysis of MMRF-CoMMpass data. <i>Journal of Computational Science</i> , 2021, 51, 101346.	2.9	6
15	Fabry disease and kidney involvement: starting from childhood to understand the future. <i>Pediatric Nephrology</i> , 2022, 37, 95-103.	1.7	13
16	Vesicular ATP-binding cassette transporters in human disease: relevant aspects of their organization for future drug development. <i>Future Drug Discovery</i> , 2020, 2, .	2.1	8
17	Learning Association Rules for Pharmacogenomic Studies. <i>Lecture Notes in Computer Science</i> , 2018, , 1-15.	1.3	0
18	Ethical Perspectives on Pharmacogenomic Profiling. , 2022, , 3-20.		1

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19	Autoimmune colitis and neutropenia in adjuvant anti-PD-1 therapy for malignant melanoma: efficacy of Vedolizumab, a case report. <i>Therapeutic Advances in Chronic Disease</i> , 2022, 13, 204062232110630.	2.5	3
20	Noncoding RNAs in patients with colorectal cancer. , 2022, , 65-95.		0
21	Risk Alleles for Multiple Myeloma Susceptibility in ADME Genes. <i>Cells</i> , 2022, 11, 189.	4.1	4
22	Using MMRFbiolinks R-Package for Discovering Prognostic Markers in Multiple Myeloma. <i>Methods in Molecular Biology</i> , 2022, 2401, 289-314.	0.9	1
23	Clustering Methods for Microarray Data Sets. <i>Methods in Molecular Biology</i> , 2022, 2401, 249-261.	0.9	0
24	Tools in Pharmacogenomics Biomarker Identification for Cancer Patients. <i>Methods in Molecular Biology</i> , 2022, 2401, 1-12.	0.9	2
25	Microarray Data Analysis Protocol. <i>Methods in Molecular Biology</i> , 2022, 2401, 263-271.	0.9	4
26	Beyond COVID-19 pandemic: Topology-aware optimization of vaccination strategy for minimizing virus spreading. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 2664-2671.	4.1	11
27	A Prognostic and Carboplatin Response Predictive Model in Ovarian Cancer: A Mono-Institutional Retrospective Study Based on Clinics and Pharmacogenomics. <i>Biomedicines</i> , 2022, 10, 1210.	3.2	1
28	A Python Clustering Analysis Protocol of Genes Expression Data Sets. <i>Genes</i> , 2022, 13, 1839.	2.4	3
29	What is precision medicine in oncology?. , 2023, , 1-30.		0
30	SEDEG: An automatic method for preprocessing and selection of seed genes from gene expression data. <i>Informatics in Medicine Unlocked</i> , 2024, 44, 101432.	3.4	0