

Nontaya Nakkam

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

523
citations

932766

10
h-index

794141

19
g-index

20
all docs

20
docs citations

20
times ranked

551
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic variants associated with severe cutaneous adverse drug reactions induced by carbamazepine. <i>British Journal of Clinical Pharmacology</i> , 2022, 88, 773-786.	1.1	8
2	Cross-reactivity between vancomycin, teicoplanin, and telavancin in patients with HLA-A*32:01 positive vancomycin-induced DRESS sharing an HLA class II haplotype. <i>Journal of Allergy and Clinical Immunology</i> , 2021, 147, 403-405.	1.5	26
3	Whole genome sequencing identifies genetic variants associated with co-trimoxazole hypersensitivity in Asians. <i>Journal of Allergy and Clinical Immunology</i> , 2021, 147, 1402-1412.	1.5	46
4	Genetic Polymorphisms of Drug-Metabolizing Enzymes Involved in 6-Mercaptopurine-Induced Myelosuppression in Thai Pediatric Acute Lymphoblastic Leukemia Patients. <i>Journal of Pediatric Genetics</i> , 2021, 10, 029-034.	0.3	2
5	Characterization of T-Cell Responses to SMX and SMX-NO in Co-Trimoxazole Hypersensitivity Patients Expressing HLA-B*13:01. <i>Frontiers in Immunology</i> , 2021, 12, 658593.	2.2	14
6	HLA-B*13:01 Is a Predictive Marker of Dapsone-Induced Severe Cutaneous Adverse Reactions in Thai Patients. <i>Frontiers in Immunology</i> , 2021, 12, 661135.	2.2	29
7	Considerations for cross-reactivity between vancomycin and other glycopeptides. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2021, 9, 3233.	2.0	2
8	Meta-Analysis of NUDT15 Genetic Polymorphism on Thiopurine-Induced Myelosuppression in Asian Populations. <i>Frontiers in Pharmacology</i> , 2021, 12, 784712.	1.6	5
9	NUDT15 is a key genetic factor for prediction of hematotoxicity in pediatric patients who received a standard low dosage regimen of 6-mercaptopurine. <i>Drug Metabolism and Pharmacokinetics</i> , 2021, 43, 100436.	1.1	1
10	Genetic Association of Co-trimoxazole-Induced Severe Cutaneous Adverse Reactions Is Phenotype-specific: HLA Class I Genotypes and Haplotypes. <i>Clinical Pharmacology and Therapeutics</i> , 2020, 108, 1078-1089.	2.3	34
11	Prevalence of CYP2C19, CYP3A4 and FMO3 genetic polymorphisms in healthy northeastern Thai volunteers. <i>ScienceAsia</i> , 2020, 46, 397.	0.2	1
12	HLA Alleles and CYP2C9*3 as Predictors of Phenytoin Hypersensitivity in East Asians. <i>Clinical Pharmacology and Therapeutics</i> , 2019, 105, 476-485.	2.3	53
13	The Medication Risk of Stevens-Johnson Syndrome and Toxic Epidermal Necrolysis in Asians: The Major Drug Causality and Comparison With the US FDA Label. <i>Clinical Pharmacology and Therapeutics</i> , 2019, 105, 112-120.	2.3	54
14	HLA Pharmacogenetic Markers of Drug Hypersensitivity in a Thai Population. <i>Frontiers in Genetics</i> , 2018, 9, 277.	1.1	24
15	Risk factors of allopurinol-induced severe cutaneous adverse reactions in a Thai population. <i>Pharmacogenetics and Genomics</i> , 2017, 27, 255-263.	0.7	25
16	Dapsone-induced severe cutaneous adverse drug reactions are strongly linked with HLA-B*13. <i>Pharmacogenetics and Genomics</i> , 2017, 27, 429-437.	0.7	87
17	Comparison between the HLA-B*13:01 Allele and Single-Nucleotide Polymorphisms in Chromosome 6 for Prediction of Allopurinol-Induced Severe Cutaneous Adverse Reactions. <i>Journal of Immunology Research</i> , 2017, 2017, 1-9.	0.9	12
18	Comparative pharmacodynamic effects of two clopidogrel formulations under steady-state conditions in healthy Thai volunteers. <i>International Journal of Clinical Pharmacology and Therapeutics</i> , 2017, 55, 177-185.	0.3	1

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19	Associations between HLA class I and cytochrome P450 2C9 genetic polymorphisms and phenytoin-related severe cutaneous adverse reactions in a Thai population. <i>Pharmacogenetics and Genomics</i> , 2016, 26, 225-234.	0.7	94
20	The impact of genetic polymorphisms of drug metabolizing enzymes on the pharmacodynamics of clopidogrel under steady state conditions. <i>Drug Metabolism and Pharmacokinetics</i> , 2015, 30, 295-304.	1.1	5