

Kye-Hwa Lee

List of Publications by Year in descending order

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33
papers

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1040056

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#	ARTICLE	IF	CITATIONS
1	Flat Pattern Peaks of Tacrolimus Absorption and Associated Pharmacogenomic Variants in Kidney Transplantation Recipients. <i>Journal of Korean Medical Science</i> , 2022, 37, e33.	2.5	3
2	Reducing severe cutaneous adverse and type B adverse drug reactions using pre-€stored human leukocyte antigen genotypes. <i>Clinical and Translational Allergy</i> , 2022, 12, e12098.	3.2	4
3	ANNO: A General Annotation Tool for Bilingual Clinical Note Information Extraction. <i>Healthcare Informatics Research</i> , 2022, 28, 89-94.	1.9	0
4	Effects of Patient-Generated Health Data: Comparison of Two Versions of Long-Term Mobile Personal Health Record Usage Logs. <i>Healthcare (Switzerland)</i> , 2022, 10, 53.	2.0	3
5	Digital Health Profile of South Korea: A Cross Sectional Study. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 6329.	2.6	2
6	In Silico Inference of Synthetic Cytotoxic Interactions from Paclitaxel Responses. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1097.	4.1	0
7	Different Seasonal Variations of Potassium in Hemodialysis Patients with High Longitudinal Potassium Levels: A Multicenter Cohort Study Using DialysisNet. <i>Yonsei Medical Journal</i> , 2021, 62, 315.	2.2	5
8	Filtered BERT: Similarity Filter-Based Augmentation with Bidirectional Transfer Learning for Protected Health Information Prediction in Clinical Documents. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 3668.	2.5	7
9	Interplay between IL6 and CRIM1 in thiopurine intolerance due to hematological toxicity in leukemic patients with wild-type NUDT15 and TPMT. <i>Scientific Reports</i> , 2021, 11, 9676.	3.3	1
10	Development and Validation of Targeted Gene Sequencing Panel Based Companion Diagnostic for Korean Patients with Solid Tumors. <i>Cancers</i> , 2021, 13, 5112.	3.7	0
11	Smart Decentralization of Personal Health Records with Physician Apps and Helper Agents on Blockchain: Platform Design and Implementation Study. <i>JMIR Medical Informatics</i> , 2021, 9, e26230.	2.6	6
12	Real-world treatment patterns of renal anemia in hemodialysis patients. <i>Medicine (United States)</i> , 2020, 99, e18749.	1.0	4
13	Identification of synthetic chemosensitivity genes paired with BRAF for BRAF/MAPK inhibitors. <i>Scientific Reports</i> , 2020, 10, 20001.	3.3	2
14	Immune expression signatures as candidate prognostic biomarkers of age and gender survival differences in cutaneous melanoma. <i>Scientific Reports</i> , 2020, 10, 12322.	3.3	6
15	Extracting Structured Genotype Information from Free-Text HLA Reports Using a Rule-Based Approach. <i>Journal of Korean Medical Science</i> , 2020, 35, e78.	2.5	6
16	High-€Throughput Algorithm for Discovering New Drug Indications by Utilizing Large-€Scale Electronic Medical Record Data. <i>Clinical Pharmacology and Therapeutics</i> , 2020, 108, 1299-1307.	4.7	7
17	Identifying genetic variants associated with ritodrine-induced pulmonary edema. <i>PLoS ONE</i> , 2020, 15, e0241215.	2.5	4
18	Clinical MetaData ontology: a simple classification scheme for data elements of clinical data based on semantics. <i>BMC Medical Informatics and Decision Making</i> , 2019, 19, 166.	3.0	7

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19	Safety and Efficacy of Ultrasound-Guided High-Intensity Focused Ultrasound Treatment for Uterine Fibroids and Adenomyosis. <i>Ultrasound in Medicine and Biology</i> , 2019, 45, 3214-3221.	1.5	38
20	Personal genome testing on physicians improves attitudes on pharmacogenomic approaches. <i>PLoS ONE</i> , 2019, 14, e0213860.	2.5	7
21	Identifying genetic variants underlying medication-induced osteonecrosis of the jaw in cancer and osteoporosis: a case control study. <i>Journal of Translational Medicine</i> , 2019, 17, 381.	4.4	24
22	Development of a Controlled Vocabulary-Based Adverse Drug Reaction Signal Dictionary for Multicenter Electronic Health Record-Based Pharmacovigilance. <i>Drug Safety</i> , 2019, 42, 657-670.	3.2	14
23	South Korean geriatrics on Beers Criteria medications at risk of adverse drug events. <i>PLoS ONE</i> , 2018, 13, e0191376.	2.5	13
24	Evolution of Translational Bioinformatics: lessons learned from TBC 2016. <i>BMC Medical Genomics</i> , 2017, 10, 32.	1.5	0
25	Standard-based comprehensive detection of adverse drug reaction signals from nursing statements and laboratory results in electronic health records. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2017, 24, 697-708.	4.4	20
26	Genome Sequence Variability Predicts Drug Precautions and Withdrawals from the Market. <i>PLoS ONE</i> , 2016, 11, e0162135.	2.5	17
27	Role of BRAF and RAS Mutations in Extrathyroidal Extension in Papillary Thyroid Cancer. <i>Cancer Genomics and Proteomics</i> , 2016, 13, 171-81.	2.0	10
28	Current Clinical Status of Telehealth in Korea: Categories, Scientific Basis, and Obstacles. <i>Healthcare Informatics Research</i> , 2015, 21, 244.	1.9	17
29	Correlation between Drug Market Withdrawals and Socioeconomic, Health, and Welfare Indicators Worldwide. <i>Journal of Korean Medical Science</i> , 2015, 30, 1567.	2.5	2
30	Using mobile phones in healthcare management for the elderly. <i>Maturitas</i> , 2014, 79, 381-388.	2.4	43
31	Annotation of Genes Having Candidate Somatic Mutations in Acute Myeloid Leukemia with Whole-Exome Sequencing Using Concept Lattice Analysis. <i>Genomics and Informatics</i> , 2013, 11, 38.	0.8	2
32	Clinical Characteristics of Patients with Medically Unexplained Chronic Widespread Pain: A Primary Care Center Study. <i>Korean Journal of Family Medicine</i> , 2011, 32, 277.	1.2	3
33	Patterns of Complementary and Alternative Medicine Therapies in Patients with Chronic Fatigue or Pain. <i>Korean Journal of Family Medicine</i> , 2009, 30, 182.	1.2	9