Jennie Lin

List of Publications by Year in descending order

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LENINIE LINI

#	Article	IF	CITATIONS
1	Automated, electronic alerts for acute kidney injury: a single-blind, parallel-group, randomised controlled trial. Lancet, The, 2015, 385, 1966-1974.	6.3	282
2	False-Positive Rate of AKI Using Consensus Creatinine–Based Criteria. Clinical Journal of the American Society of Nephrology: CJASN, 2015, 10, 1723-1731.	2.2	94
3	The long noncoding RNA landscape in hypoxic and inflammatory renal epithelial injury. American Journal of Physiology - Renal Physiology, 2015, 309, F901-F913.	1.3	70
4	Single-cell RNA sequencing uncovers heterogenous transcriptional signatures in macrophages during efferocytosis. Scientific Reports, 2020, 10, 14333.	1.6	48
5	Interrogation of nonconserved human adipose lincRNAs identifies a regulatory role of <i>linc-ADAL</i> in adipocyte metabolism. Science Translational Medicine, 2018, 10, .	5.8	42
6	Deep RNA Sequencing Uncovers a Repertoire of Human Macrophage Long Intergenic Noncoding RNAs Modulated by Macrophage Activation and Associated With Cardiometabolic Diseases. Journal of the American Heart Association, 2017, 6, .	1.6	36
7	Transcriptome-Wide Analysis Reveals Modulation of Human Macrophage Inflammatory Phenotype Through Alternative Splicing. Arteriosclerosis, Thrombosis, and Vascular Biology, 2016, 36, 1434-1447.	1.1	35
8	Plasma Lipoprotein(a) Levels Are Associated with Mild Renal Impairment in Type 2 Diabetics Independent of Albuminuria. PLoS ONE, 2014, 9, e114397.	1.1	28
9	Genome engineering tools for building cellular models of disease. FEBS Journal, 2016, 283, 3222-3231.	2.2	23
10	Profiling APOL1 Nephropathy Risk Variants in Genome-Edited Kidney Organoids with Single-Cell Transcriptomics. Kidney360, 2020, 1, 203-215.	0.9	18
11	From Genotype to Phenotype. Circulation Genomic and Precision Medicine, 2018, 11, .	1.6	16
12	Relation of atherogenic lipoproteins with estimated glomerular filtration rate decline: a longitudinal study. BMC Nephrology, 2015, 16, 130.	0.8	14
13	PennDiff: detecting differential alternative splicing and transcription by RNA sequencing. Bioinformatics, 2018, 34, 2384-2391.	1.8	14
14	Picking Up the Slack: Collaboration Tools to Build Community and Increase Productivity in Nephrology. Seminars in Nephrology, 2020, 40, 298-302.	0.6	5
15	From Genotype to Phenotype: A Primer on the Functional Follow-up of Genome-Wide Association Studies in Cardiovascular Disease. Circulation Genomic and Precision Medicine, 2018, 11, .	1.6	5
16	Is Hemodialysis Bad for the Heart?. Seminars in Dialysis, 2012, 25, 86-87.	0.7	1
17	Outscoring Current Classification Systems for Nephrotic Syndrome. American Journal of Kidney Diseases, 2022, , .	2.1	0