William C L Stewart

List of Publications by Year
in descending order

14 papers	136 citations	1684188 5 h-index	1281871 11 g-index
14 all docs	14 docs citations	14 times ranked	305 citing authors

#	Article	IF	Citations
1	FREQMAX provides an alternative approach for determining high‑resolution allele frequency thresholds in carrier screening. Human Mutation, 2020, 41, 2078-2086.	2.5	1
2	A new linear regression-like residual for survival analysis, with application to genome wide association studies of time-to-event data. PLoS ONE, 2020, 15, e0232300.	2.5	7
3	Brd2 haploinsufficiency extends lifespan and healthspan in C57B6/J mice. PLoS ONE, 2020, 15, e0234910.	2.5	6
4	In Response: *ME2* association analysis in adolescent‑onset genetic generalized epilepsy. Epilepsia, 2019, 60, 2001-2002.	5.1	2
5	Replication, reanalysis, and gene expression: *ME2* and genetic generalized epilepsy. Epilepsia, 2019, 60, 539-546.	5.1	7
6	Lung Allocation Score Thresholds Prioritize Survival After Lung Transplantation. Chest, 2019, 156, 64-70.	0.8	22
7	Mathematical modelling identifies the role of adaptive immunity as a key controller of respiratory syncytial virus in cotton rats. Journal of the Royal Society Interface, 2019, 16, 20190389.	3.4	19
8	DNA methylation of the BRD 2 promoter is associated with juvenile myoclonic epilepsy in Caucasians. Epilepsia, 2018, 59, 1011-1019.	5.1	28
9	Mutations in bacterial genes induce unanticipated changes in the relationship between bacterial pathogens in experimental otitis media. Royal Society Open Science, 2018, 5, 180810.	2.4	3
10	Predictive Utility of Lung Allocation Score for Retransplantation Outcomes. Annals of Thoracic Surgery, 2018, 106, 1525-1532.	1.3	5
11	Connecting the dots across time: reconstruction of single-cell signalling trajectories using time-stamped data. Royal Society Open Science, 2017, 4, 170811.	2.4	6
12	In silico modeling identifies CD45 as a regulator of IL-2 synergy in the NKG2D-mediated activation of immature human NK cells. Science Signaling, 2017, 10, .	3.6	23
13	Increasing the power of association studies with affected families, unrelated cases and controls. Frontiers in Genetics, 2013, 4, 200.	2.3	5
14	Obtaining Accurate p Values from a Dense SNP Linkage Scan. Human Heredity, 2012, 74, 12-16.	0.8	2