

# Jonathan Alexander Heiss

## List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

6

papers

80

citations

4

h-index

8

g-index

8

ext. papers

131

ext. citations

5.6

avg, IF

3.17

L-index

#	Paper	IF	Citations
6	Identifying mislabeled and contaminated DNA methylation microarray data: an extended quality control toolset with examples from GEO. <i>Clinical Epigenetics</i> , <b>2018</b> , 10, 73	7.7	37
5	Improved filtering of DNA methylation microarray data by detection p values and its impact on downstream analyses. <i>Clinical Epigenetics</i> , <b>2019</b> , 11, 15	7.7	19
4	Battle of epigenetic proportions: comparing Illuminals EPIC methylation microarrays and TruSeq targeted bisulfite sequencing. <i>Epigenetics</i> , <b>2020</b> , 15, 174-182	5.7	16
3	Prenatal lead exposure and cord blood DNA methylation in PROGRESS: an epigenome-wide association study. <i>Environmental Epigenetics</i> , <b>2020</b> , 6, dvaa014	2.4	4
2	Saliva cell type DNA methylation reference panel for epidemiological studies in children. <i>Epigenetics</i> , <b>2021</b> , 1-17	5.7	3
1	Maternal haemoglobin levels in pregnancy and child DNA methylation: a study in the pregnancy and childhood epigenetics consortium. <i>Epigenetics</i> , <b>2021</b> , 1-13	5.7	1