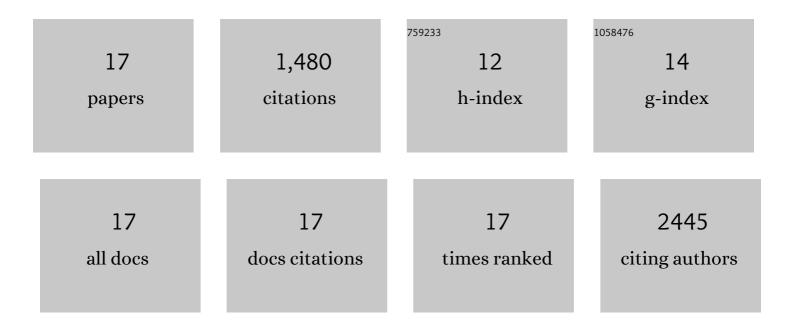
Shu Tao

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Histone H3 trimethylation at lysine 36 guides m6A RNA modification co-transcriptionally. Nature, 2019, 567, 414-419. | 27.8 | 452 |
| 2 | Draft genome of the globally widespread and invasive Argentine ant (<i>Linepithema humile</i>). Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 5673-5678. | 7.1 | 257 |
| 3 | The Genome Sequence of the Leaf-Cutter Ant Atta cephalotes Reveals Insights into Its Obligate Symbiotic Lifestyle. PLoS Genetics, 2011, 7, e1002007. | 3.5 | 231 |
| 4 | Draft genome of the red harvester ant <i>Pogonomyrmex barbatus</i> . Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 5667-5672. | 7.1 | 222 |
| 5 | Prospective Study of Cancer Genetic Variants: Variation in Rate of Reclassification by Ancestry. Journal of the National Cancer Institute, 2018, 110, 1059-1066. | 6.3 | 48 |
| 6 | Inhibition of MDR1 Overcomes Resistance to Brentuximab Vedotin in Hodgkin Lymphoma. Clinical Cancer Research, 2020, 26, 1034-1044. | 7.0 | 48 |
| 7 | Pathogenic and likely pathogenic variants in <i>PALB2</i> , <i>CHEK2</i> , and other known breast cancer susceptibility genes among 1054 <i>BRCA</i> â€negative Hispanics with breast cancer. Cancer, 2019, 125, 2829-2836. | 4.1 | 43 |
| 8 | Genetic Gastric Cancer Susceptibility in the International Clinical Cancer Genomics Community Research Network. Cancer Genetics, 2017, 216-217, 111-119. | 0.4 | 42 |
| 9 | Reprogramming of DNA methylation at NEUROD2-bound sequences during cortical neuron differentiation. Science Advances, 2019, 5, eaax0080. | 10.3 | 32 |
| 10 | Parallel (Randomized) Phase II Evaluation of Tivantinib (ARQ197) and Tivantinib in Combination with Erlotinib in Papillary Renal Cell Carcinoma: SWOG S1107. Kidney Cancer, 2017, 1, 123-132. | 0.4 | 31 |
| 11 | A chymotrypsin-like serine protease cDNA involved in food protein digestion in the common cutworm, Spodoptera litura: Cloning, characterization, developmental and induced expression patterns, and localization. Journal of Insect Physiology, 2010, 56, 788-799. | 2.0 | 30 |
| 12 | The spectrum of genetic variants in hereditary pancreatic cancer includes Fanconi anemia genes. Familial Cancer, 2018, 17, 235-245. | 1.9 | 29 |
| 13 | Repurposing leflunomide for relapsed/refractory multiple myeloma: a phase 1 study. Leukemia and Lymphoma, 2020, 61, 1669-1677. | 1.3 | 13 |
| 14 | The Most Redundant Sequences in Human CpG Island Library Are Derived from Mitochondrial Genome. Genomics, Proteomics and Bioinformatics, 2010, 8, 81-91. | 6.9 | 2 |
| 15 | Exome sequencing of tumor samples from S1107 "Randomized phase II evaluation of tivantinib and tivantinib in combination with erlotinib in patients with papillary renal cell carcinoma (pRCC)― Annals of Oncology, 2017, 28, v323. | 1.2 | 0 |
| 16 | Abstract P4-09-02: Somatic tumor profiles of breast cancer in Latinas. , 2020, , . | | 0 |
| 17 | Abstract B081: Common copy number aberrations in breast cancer in Latinas. , 2020, , . | | 0 |