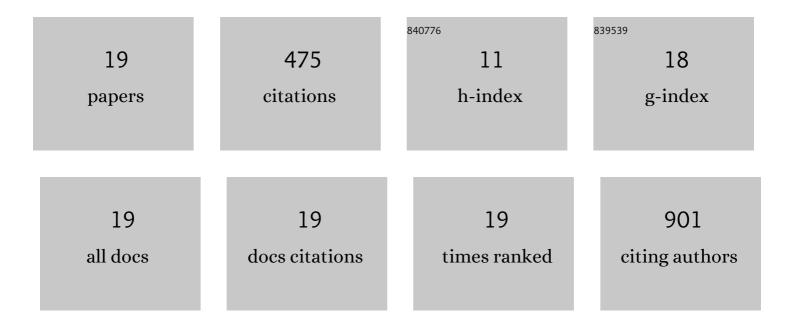
## Jian Wang

List of Publications by Year in descending order

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ΙΙΔΝ ΜΑΝΟ

| #  | Article  | lF   | CITATIONS |
|----|--|------|-----------|
| 1  | An Approach to Analyze Longitudinal Zero-Inflated Microbiome Count Data Using Two-Stage Mixed<br>Effects Models. Statistics in Biosciences, 2021, 13, 267-290.   | 1.2  | 3         |
| 2  | Regulation of tumor immune suppression and cancer cell survival by CXCL1/2 elevation in glioblastoma multiforme. Science Advances, 2021, 7, .  | 10.3 | 54        |
| 3  | A Bayesian hierarchical monitoring design for phase II cancer clinical trials: Incorporating information on response duration into monitoring rules. Statistics in Medicine, 2021, 40, 4629-4639.  | 1.6  | 0         |
| 4  | Mediation model with a categorical exposure and a censored mediator with application to a genetic study. PLoS ONE, 2021, 16, e0257628.   | 2.5  | 2         |
| 5  | Oral microbiome and onset of oral mucositis in patients with squamous cell carcinoma of the head and neck. Cancer, 2020, 126, 5124-5136.   | 4.1  | 30        |
| 6  | Longitudinal associations of family functioning with body mass index in Mexican-origin adolescents living in the U.S Preventive Medicine, 2019, 118, 309-316.  | 3.4  | 4         |
| 7  | FGL2 promotes tumor progression in the CNS by suppressing CD103+ dendritic cell differentiation.<br>Nature Communications, 2019, 10, 448.  | 12.8 | 65        |
| 8  | Mediation analysis in a caseâ€control study when the mediator is a censored variable. Statistics in<br>Medicine, 2019, 38, 1213-1229.  | 1.6  | 5         |
| 9  | Estimation of indirect effect when the mediator is a censored variable. Statistical Methods in Medical Research, 2018, 27, 3010-3025.  | 1.5  | 6         |
| 10 | Processing and Analyzing Human Microbiome Data. Methods in Molecular Biology, 2017, 1666, 649-677.   | 0.9  | 4         |
| 11 | Identifying novel genes and biological processes relevant to the development of cancer<br>therapy-induced mucositis: An informative gene network analysis. PLoS ONE, 2017, 12, e0180396.   | 2.5  | 27        |
| 12 | Genome-wide association study suggests common variants within RP11-634B7.4 gene influencing severe pre-treatment pain in head and neck cancer patients. Scientific Reports, 2016, 6, 34206.  | 3.3  | 12        |
| 13 | MAPK1/ERK2 as novel target genes for pain in head and neck cancer patients. BMC Genetics, 2016, 17, 40.  | 2.7  | 25        |
| 14 | Gene network analysis shows immune-signaling and ERK1/2 as novel genetic markers for multiple addiction phenotypes: alcohol, smoking and opioid addiction. BMC Systems Biology, 2015, 9, 25.   | 3.0  | 43        |
| 15 | Genetic Variations in Interleukin-8 and Interleukin-10 Are Associated With Pain, Depressed Mood, and<br>Fatigue in Lung Cancer Patients. Journal of Pain and Symptom Management, 2013, 46, 161-172.                                      | 1.2  | 57        |
| 16 | Method for Evaluating Multiple Mediators: Mediating Effects of Smoking and COPD on the<br>Association between the CHRNA5-A3 Variant and Lung Cancer Risk. PLoS ONE, 2012, 7, e47705.   | 2.5  | 23        |
| 17 | Estimation of odds ratios of genetic variants for the secondary phenotypes associated with primary diseases. Genetic Epidemiology, 2011, 35, 190-200.  | 1.3  | 37        |
| 18 | Power and type I error results for a bias-correction approach recently shown to provide accurate odds ratios of genetic variants for the secondary phenotypes associated with primary diseases. Genetic Epidemiology, 2011, 35, 739-743. | 1.3  | 11        |

| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Mediating effects of smoking and chronic obstructive pulmonary disease on the relation between the CHRNA5â€A3 genetic locus and lung cancer risk. Cancer, 2010, 116, 3458-3462. | 4.1 | 67        |