## Qian Wang

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/241009/publications.pdf

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| 27       | 1,880          | 17 h-index   | 27             |
|----------|----------------|--------------|----------------|
| papers   | citations      |              | g-index        |
| 31       | 31             | 31           | 3004           |
| all docs | docs citations | times ranked | citing authors |

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Integrative metabolomicsâ€genomics approach reveals key metabolic pathways and regulators of Alzheimer's disease. Alzheimer's and Dementia, 2022, 18, 1260-1278.                                       | 0.8  | 57        |
| 2  | The landscape of human tissue and cell type specific expression and co-regulation of senescence genes. Molecular Neurodegeneration, 2022, $17, 5$ .  | 10.8 | 34        |
| 3  | Enhancing autophagy maturation with CCZ1-MON1A complex alleviates neuropathology and memory defects in Alzheimer disease models. Theranostics, 2022, 12, 1738-1755.                                    | 10.0 | 13        |
| 4  | Guidelines for bioinformatics of single-cell sequencing data analysis in Alzheimer's disease: review, recommendation, implementation and application. Molecular Neurodegeneration, 2022, 17, 17.       | 10.8 | 40        |
| 5  | Whole genome sequencing–based copy number variations reveal novel pathways and targets in Alzheimer's disease. Alzheimer's and Dementia, 2022, 18, 1846-1867.  | 0.8  | 13        |
| 6  | Microglia Share the Burden. Neuroscience Bulletin, 2022, 38, 695-698.  | 2.9  | 3         |
| 7  | Transformative Network Modeling of Multi-omics Data Reveals Detailed Circuits, Key Regulators, and Potential Therapeutics for Alzheimer's Disease. Neuron, 2021, 109, 257-272.e14.                     | 8.1  | 108       |
| 8  | Molecular subtyping of Alzheimer's disease using RNA sequencing data reveals novel mechanisms and targets. Science Advances, 2021, 7, .  | 10.3 | 137       |
| 9  | Association of neurogranin gene expression with Alzheimer's disease pathology in the perirhinal cortex. Alzheimer's and Dementia: Translational Research and Clinical Interventions, 2021, 7, e12162.  | 3.7  | 6         |
| 10 | Disentangling the Molecular Pathways of Parkinson's Disease using Multiscale Network Modeling.<br>Trends in Neurosciences, 2021, 44, 182-188.  | 8.6  | 3         |
| 11 | ALS-FTLD-linked mutations of SQSTM1/p62 disrupt selective autophagy and NFE2L2/NRF2 anti-oxidative stress pathway. Autophagy, 2020, 16, 917-931.   | 9.1  | 118       |
| 12 | hUC-MSCs ameliorated CUMS-induced depression by modulating complement C3 signaling-mediated microglial polarization during astrocyte-microglia crosstalk. Brain Research Bulletin, 2020, 163, 109-119. | 3.0  | 23        |
| 13 | Sexâ€specific peripheral and central responses to stressâ€induced depression and treatment in a mouse model. Journal of Neuroscience Research, 2020, 98, 2541-2553.                                    | 2.9  | 14        |
| 14 | Synj1 haploinsufficiency causes dopamine neuron vulnerability and alpha-synuclein accumulation in mice. Human Molecular Genetics, 2020, 29, 2300-2312.   | 2.9  | 29        |
| 15 | Microglia clear neuron-released $\hat{l}$ ±-synuclein via selective autophagy and prevent neurodegeneration. Nature Communications, 2020, $11,1386$ .  | 12.8 | 279       |
| 16 | Systems modeling of white matter microstructural abnormalities in Alzheimer's disease. NeuroImage: Clinical, 2020, 26, 102203.   | 2.7  | 12        |
| 17 | The landscape of multiscale transcriptomic networks and key regulators in Parkinson's disease.<br>Nature Communications, 2019, 10, 5234.   | 12.8 | 82        |
| 18 | Autophagy protein NRBF2 has reduced expression in Alzheimer's brains and modulates memory and amyloid-beta homeostasis in mice. Molecular Neurodegeneration, 2019, 14, 43.                             | 10.8 | 63        |

| #  | Article   | IF  | CITATION |
|----|---|-----|----------|
| 19 | Complement C3aR Inactivation Attenuates Tau Pathology and Reverses an Immune Network Deregulated in Tauopathy Models and Alzheimer's Disease. Neuron, 2018, 100, 1337-1353.e5.  | 8.1 | 306      |
| 20 | The Mount Sinai cohort of large-scale genomic, transcriptomic and proteomic data in Alzheimer's disease. Scientific Data, 2018, 5, 180185.  | 5.3 | 320      |
| 21 | Parkinson's Disease-Associated LRRK2 Hyperactive Kinase Mutant Disrupts Synaptic Vesicle Trafficking in Ventral Midbrain Neurons. Journal of Neuroscience, 2017, 37, 11366-11376.   | 3.6 | 103      |
| 22 | Characterization of Gonadotrope Secretoproteome Identifies Neurosecretory Protein VGF-derived Peptide Suppression of Follicle-stimulating Hormone Gene Expression. Journal of Biological Chemistry, 2016, 291, 21322-21334.                 | 3.4 | 9        |
| 23 | Homer1 Alternative Splicing Is Regulated by Gonadotropin-Releasing Hormone and Modulates<br>Gonadotropin Gene Expression. Molecular and Cellular Biology, 2014, 34, 1747-1756.  | 2.3 | 6        |
| 24 | Growth Differentiation Factor 9 (GDF9) Forms an Incoherent Feed-forward Loop Modulating Follicle-stimulating Hormone Î <sup>2</sup> -Subunit (FSHÎ <sup>2</sup> ) Gene Expression. Journal of Biological Chemistry, 2014, 289, 16164-16175. | 3.4 | 26       |
| 25 | Outside the box signaling: Secreted factors modulate GnRH receptor-mediated gonadotropin regulation. Molecular and Cellular Endocrinology, 2014, 385, 56-61.  | 3.2 | 22       |
| 26 | Î <sup>2</sup> -Catenin Regulates GnRH-Induced FSHÎ <sup>2</sup> Gene Expression. Molecular Endocrinology, 2013, 27, 224-237.   | 3.7 | 17       |
| 27 | The Hippocampal Proteomic Analysis of Senescence-Accelerated Mouse: Implications of Uchl3 and Mitofilin in Cognitive Disorder and Mitochondria Dysfunction in SAMP8. Neurochemical Research, 2008, 33, 1776-1782.                           | 3.3 | 26       |