

Jean-Philippe CoppÃ©

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2821470/publications.pdf>

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11
papers

7,573
citations

1040056

9
h-index

1474206

9
g-index

11
all docs

11
docs citations

11
times ranked

10668
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | The Senescence-Associated Secretory Phenotype: The Dark Side of Tumor Suppression. Annual Review of Pathology: Mechanisms of Disease, 2010, 5, 99-118. | 22.4 | 3,486 |
| 2 | Senescence-Associated Secretory Phenotypes Reveal Cell-Nonautonomous Functions of Oncogenic RAS and the p53 Tumor Suppressor. PLoS Biology, 2008, 6, e301. | 5.6 | 3,067 |
| 3 | Tumor Suppressor and Aging Biomarker p16INK4a Induces Cellular Senescence without the Associated Inflammatory Secretory Phenotype. Journal of Biological Chemistry, 2011, 286, 36396-36403. | 3.4 | 380 |
| 4 | A Human-Like Senescence-Associated Secretory Phenotype Is Conserved in Mouse Cells Dependent on Physiological Oxygen. PLoS ONE, 2010, 5, e9188. | 2.5 | 356 |
| 5 | Targetable mechanisms driving immunoevasion of persistent senescent cells link chemotherapy-resistant cancer to aging. JCI Insight, 2019, 4, . | 5.0 | 90 |
| 6 | Targeting amphiregulin (AREG) derived from senescent stromal cells diminishes cancer resistance and averts programmed cell death 1 ligand (PD-L1)-mediated immunosuppression. Aging Cell, 2019, 18, e13027. | 6.7 | 79 |
| 7 | A protein interaction landscape of breast cancer. Science, 2021, 374, eabf3066. | 12.6 | 66 |
| 8 | Mapping phospho-catalytic dependencies of therapy-resistant tumours reveals actionable vulnerabilities. Nature Cell Biology, 2019, 21, 778-790. | 10.3 | 24 |
| 9 | An Atlas of the Human Kinome Reveals the Mutational Landscape Underlying Dysregulated Phosphorylation Cascades in Cancer. Cancer Research, 2016, 76, 1733-1745. | 0.9 | 20 |
| 10 | High-Throughput Kinase Activity Mapping (HT-KAM) system: biochemical assay. Protocol Exchange, 0, , . | 0.3 | 3 |
| 11 | High-Throughput Kinase Activity Mapping (HT-KAM) system: analysis of phospho-catalytic profiles. Protocol Exchange, 0, , . | 0.3 | 2 |