

# Beatriz Alvarado

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4890973/publications.pdf>

Version: 2024-02-01

8  
papers

3,051  
citations

1162367

8  
h-index

1588620

8  
g-index

10  
all docs

10  
docs citations

10  
times ranked

6162  
citing authors

| # | ARTICLE   | IF   | CITATIONS |
|---|---|------|-----------|
| 1 | Outer Radial Glia-like Cancer Stem Cells Contribute to Heterogeneity of Glioblastoma. <i>Cell Stem Cell</i> , 2020, 26, 48-63.e6.   | 5.2  | 222       |
| 2 | The Phenotypes of Proliferating Glioblastoma Cells Reside on a Single Axis of Variation. <i>Cancer Discovery</i> , 2019, 9, 1708-1719.  | 7.7  | 205       |
| 3 | Establishing Cerebral Organoids as Models of Human-Specific Brain Evolution. <i>Cell</i> , 2019, 176, 743-756.e17.  | 13.5 | 423       |
| 4 | Multimodal Single-Cell Analysis Reveals Physiological Maturation in the Developing Human Neocortex. <i>Neuron</i> , 2019, 102, 143-158.e7.  | 3.8  | 61        |
| 5 | Regulation of cell-type-specific transcriptomes by microRNA networks during human brain development. <i>Nature Neuroscience</i> , 2018, 21, 1784-1792.  | 7.1  | 121       |
| 6 | Spatiotemporal gene expression trajectories reveal developmental hierarchies of the human cortex. <i>Science</i> , 2017, 358, 1318-1323.  | 6.0  | 717       |
| 7 | Single-cell profiling of human gliomas reveals macrophage ontogeny as a basis for regional differences in macrophage activation in the tumor microenvironment. <i>Genome Biology</i> , 2017, 18, 234. | 3.8  | 448       |
| 8 | Low-coverage single-cell mRNA sequencing reveals cellular heterogeneity and activated signaling pathways in developing cerebral cortex. <i>Nature Biotechnology</i> , 2014, 32, 1053-1058.            | 9.4  | 850       |