

Ruzhica Bogeska

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3384011/publications.pdf>

Version: 2024-02-01

11
papers

760
citations

1307594

7
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

1330
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Platelet GPIb β is a mediator and potential interventional target for NASH and subsequent liver cancer. <i>Nature Medicine</i> , 2019, 25, 641-655. | 30.7 | 259 |
| 2 | Somatic mutation rates scale with lifespan across mammals. <i>Nature</i> , 2022, 604, 517-524. | 27.8 | 211 |
| 3 | Haematopoietic stem cells in perisinusoidal niches are protected from ageing. <i>Nature Cell Biology</i> , 2019, 21, 1309-1320. | 10.3 | 88 |
| 4 | MPN patients harbor recurrent truncating mutations in transcription factor NF-E2. <i>Journal of Experimental Medicine</i> , 2013, 210, 1003-1019. | 8.5 | 69 |
| 5 | A novel murine model of myeloproliferative disorders generated by overexpression of the transcription factor NF-E2. <i>Journal of Experimental Medicine</i> , 2012, 209, 35-50. | 8.5 | 67 |
| 6 | Continuous mitotic activity of primitive hematopoietic stem cells in adult mice. <i>Journal of Experimental Medicine</i> , 2020, 217, . | 8.5 | 25 |
| 7 | The Fanconi anemia pathway is required for efficient repair of stress-induced DNA damage in haematopoietic stem cells. <i>Cell Cycle</i> , 2015, 14, 2734-2742. | 2.6 | 18 |
| 8 | Elevated Nuclear Factor Erythroid-2 Levels Promote Epo-Independent Erythroid Maturation and Recapitulate the Hematopoietic Stem Cell and Common Myeloid Progenitor Expansion Observed in Polycythemia Vera Patients. <i>Stem Cells Translational Medicine</i> , 2013, 2, 112-117. | 3.3 | 7 |
| 9 | Resilience to social stress: is it in the blood?. <i>FEBS Open Bio</i> , 2021, 11, 2675-2677. | 2.3 | 3 |
| 10 | Fast and high-fidelity in situ 3D imaging protocol for stem cells and niche components for mouse organs and tissues. <i>STAR Protocols</i> , 2022, 3, 101483. | 1.2 | 3 |
| 11 | Deconvolution of Hematopoietic Commitment Decisions By Genome-Wide Analysis of Progressive DNA Methylation Changes. <i>Blood</i> , 2019, 134, 1179-1179. | 1.4 | 0 |