

# Szymon J Ciesielski

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6722190/publications.pdf>

Version: 2024-02-01

9  
papers

264  
citations

1306789

7  
h-index

1473754

9  
g-index

9  
all docs

9  
docs citations

9  
times ranked

407  
citing authors

| # | ARTICLE   | IF  | CITATIONS |
|---|---|-----|-----------|
| 1 | Congenital sideroblastic anemia due to mutations in the mitochondrial HSP70 homologue HSPA9. <i>Blood</i> , 2015, 126, 2734-2738.   | 0.6 | 78        |
| 2 | Roles of Intramolecular and Intermolecular Interactions in Functional Regulation of the Hsp70 J-protein Co-Chaperone Sis1. <i>Journal of Molecular Biology</i> , 2015, 427, 1632-1643.  | 2.0 | 46        |
| 3 | Overlapping Binding Sites of the Frataxin Homologue Assembly Factor and the Heat Shock Protein 70 Transfer Factor on the Isu Iron-Sulfur Cluster Scaffold Protein. <i>Journal of Biological Chemistry</i> , 2014, 289, 30268-30278. | 1.6 | 38        |
| 4 | Broadening the functionality of a J-protein/Hsp70 molecular chaperone system. <i>PLoS Genetics</i> , 2017, 13, e1007084.  | 1.5 | 30        |
| 5 | Protection of scaffold protein Isu from degradation by the Lon protease Pim1 as a component of Fe-S cluster biogenesis regulation. <i>Molecular Biology of the Cell</i> , 2016, 27, 1060-1068.                                      | 0.9 | 22        |
| 6 | Iron-Sulfur Cluster Biogenesis Chaperones: Evidence for Emergence of Mutational Robustness of a Highly Specific Protein-Protein Interaction. <i>Molecular Biology and Evolution</i> , 2016, 33, 643-656.                            | 3.5 | 19        |
| 7 | Two-step mechanism of J-domain action in driving Hsp70 function. <i>PLoS Computational Biology</i> , 2020, 16, e1007913.  | 1.5 | 18        |
| 8 | Structure and evolution of the 4-helix bundle domain of Zuotin, a J-domain protein co-chaperone of Hsp70. <i>PLoS ONE</i> , 2019, 14, e0217098.   | 1.1 | 8         |
| 9 | Posttranslational control of the scaffold for Fe-S cluster biogenesis as a compensatory regulatory mechanism. <i>Current Genetics</i> , 2017, 63, 51-56.  | 0.8 | 5         |