

# Mercedes Pardo

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4060783/publications.pdf>

Version: 2024-02-01

27  
papers

1,359  
citations

430874

18  
h-index

552781

26  
g-index

32  
all docs

32  
docs citations

32  
times ranked

3025  
citing authors

#	ARTICLE	IF	CITATIONS
1	The ubiquitin-dependent ATPase p97 removes cytotoxic trapped PARP1 from chromatin. <i>Nature Cell Biology</i> , 2022, 24, 62-73.	10.3	66
2	Aneuploidy tolerance caused by BRG1 loss allows chromosome gains and recovery of fitness. <i>Nature Communications</i> , 2022, 13, 1731.	12.8	9
3	CRISPR activation screen in mice identifies novel membrane proteins enhancing pulmonary metastatic colonisation. <i>Communications Biology</i> , 2021, 4, 395.	4.4	12
4	Ubiquitylation of MLKL at lysine 219 positively regulates necroptosis-induced tissue injury and pathogen clearance. <i>Nature Communications</i> , 2021, 12, 3364.	12.8	43
5	HMGB1 Protein Interactions in Prostate and Ovary Cancer Models Reveal Links to RNA Processing and Ribosome Biogenesis through NuRD, THOC and Septin Complexes. <i>Cancers</i> , 2021, 13, 4686.	3.7	4
6	A commercial antibody to the human condensin II subunit NCAPH2 cross-reacts with a SWI/SNF complex component. <i>Wellcome Open Research</i> , 2021, 6, 3.	1.8	2
7	Mutations in FAM50A suggest that Armfield XLID syndrome is a spliceosomopathy. <i>Nature Communications</i> , 2020, 11, 3698.	12.8	38
8	Cyclin B1-Cdk1 facilitates MAD1 release from the nuclear pore to ensure a robust spindle checkpoint. <i>Journal of Cell Biology</i> , 2020, 219, .	5.2	35
9	Landscape of the Plasmodium Interactome Reveals Both Conserved and Species-Specific Functionality. <i>Cell Reports</i> , 2019, 28, 1635-1647.e5.	6.4	49
10	Proteomic navigation using proximity-labeling. <i>Methods</i> , 2019, 164-165, 67-72.	3.8	6
11	The BAF and PRC2 Complex Subunits Dpf2 and Eed Antagonistically Converge on Tbx3 to Control ESC Differentiation. <i>Cell Stem Cell</i> , 2019, 24, 138-152.e8.	11.1	30
12	Nbeal2 interacts with Dock7, Sec16a, and Vac14. <i>Blood</i> , 2018, 131, 1000-1011.	1.4	29
13	UTX-mediated enhancer and chromatin remodeling suppresses myeloid leukemogenesis through noncatalytic inverse regulation of ETS and GATA programs. <i>Nature Genetics</i> , 2018, 50, 883-894.	21.4	117
14	Eros is a novel transmembrane protein that controls the phagocyte respiratory burst and is essential for innate immunity. <i>Journal of Experimental Medicine</i> , 2017, 214, 1111-1128.	8.5	50
15	Genomic Determinants of Protein Abundance Variation in Colorectal Cancer Cells. <i>Cell Reports</i> , 2017, 20, 2201-2214.	6.4	95
16	Myst2/Kat7 histone acetyltransferase interaction proteomics reveals tumour-suppressor Niam as a novel binding partner in embryonic stem cells. <i>Scientific Reports</i> , 2017, 7, 8157.	3.3	12
17	Integrative epigenomics, transcriptomics and proteomics of patient chondrocytes reveal genes and pathways involved in osteoarthritis. <i>Scientific Reports</i> , 2017, 7, 8935.	3.3	90
18	Resolving Affinity Purified Protein Complexes by Blue Native PAGE and Protein Correlation Profiling. <i>Journal of Visualized Experiments</i> , 2017, , .	0.3	8

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19	Activation of the Aryl Hydrocarbon Receptor Interferes with Early Embryonic Development. <i>Stem Cell Reports</i> , 2017, 9, 1377-1386.	4.8	39
20	Multiple short windows of calcium-dependent protein kinase 4 activity coordinate distinct cell cycle events during <i>Plasmodium</i> gametogenesis. <i>ELife</i> , 2017, 6, .	6.0	62
21	Characterization of Two Distinct Nucleosome Remodeling and Deacetylase (NuRD) Complex Assemblies in Embryonic Stem Cells. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 878-891.	3.8	42
22	Mechanisms controlling the temporal degradation of Nek2A and Kif18A by the APC/Câ€“Cdc20 complex. <i>EMBO Journal</i> , 2013, 32, 303-314.	7.8	61
23	Nuclear receptor binding protein 1 regulates intestinal progenitor cell homeostasis and tumour formation. <i>EMBO Journal</i> , 2012, 31, 2486-2497.	7.8	40
24	Assignment of Protein Interactions from Affinity Purification/Mass Spectrometry Data. <i>Journal of Proteome Research</i> , 2012, 11, 1462-1474.	3.7	52
25	An Expanded Oct4 Interaction Network: Implications for Stem Cell Biology, Development, and Disease. <i>Cell Stem Cell</i> , 2010, 6, 382-395.	11.1	338
26	The nuclear rim protein Amo1 is required for proper microtubule cytoskeleton organisation in fission yeast. <i>Journal of Cell Science</i> , 2005, 118, 1705-1714.	2.0	19
27	Identifying and characterising Thrap3, Bclaf1 and Erh interactions using cross-linking mass spectrometry. <i>Wellcome Open Research</i> , 0, 6, 260.	1.8	2