

# Maria Berdasco

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5626070/publications.pdf>

Version: 2024-02-01

31  
papers

3,731  
citations

304743

22  
h-index

454955

30  
g-index

31  
all docs

31  
docs citations

31  
times ranked

7728  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Changes in the pattern of DNA methylation associate with twin discordance in systemic lupus erythematosus. <i>Genome Research</i> , 2010, 20, 170-179.   | 5.5  | 569       |
| 2  | Aberrant Epigenetic Landscape in Cancer: How Cellular Identity Goes Awry. <i>Developmental Cell</i> , 2010, 19, 698-711.   | 7.0  | 529       |
| 3  | Clinical epigenetics: seizing opportunities for translation. <i>Nature Reviews Genetics</i> , 2019, 20, 109-127.   | 16.3 | 353       |
| 4  | A DNA methylation fingerprint of 1628 human samples. <i>Genome Research</i> , 2012, 22, 407-419.   | 5.5  | 341       |
| 5  | The timeline of epigenetic drug discovery: from reality to dreams. <i>Clinical Epigenetics</i> , 2019, 11, 174.  | 4.1  | 275       |
| 6  | Quantitative comparison of DNA methylation assays for biomarker development and clinical applications. <i>Nature Biotechnology</i> , 2016, 34, 726-737.  | 17.5 | 270       |
| 7  | Epigenetic inactivation of the Sotos overgrowth syndrome gene histone methyltransferase NSD1 in human neuroblastoma and glioma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 21830-21835. | 7.1  | 190       |
| 8  | Temozolomide Resistance in Glioblastoma Cell Lines: Implication of MGMT, MMR, P-Glycoprotein and CD133 Expression. <i>PLoS ONE</i> , 2015, 10, e0140131.   | 2.5  | 144       |
| 9  | Genetic syndromes caused by mutations in epigenetic genes. <i>Human Genetics</i> , 2013, 132, 359-383.   | 3.8  | 141       |
| 10 | Interplay between long non-coding RNAs and epigenetic machinery: emerging targets in cancer?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2018, 373, 20170074.   | 4.0  | 112       |
| 11 | Epigenetic mechanisms during ageing and neurogenesis as novel therapeutic avenues in human brain disorders. <i>Clinical Epigenetics</i> , 2017, 9, 67.   | 4.1  | 108       |
| 12 | DNA methylation in stem cell renewal and multipotency. <i>Stem Cell Research and Therapy</i> , 2011, 2, 42.  | 5.5  | 85        |
| 13 | Hot topics in epigenetic mechanisms of aging: 2011. <i>Aging Cell</i> , 2012, 11, 181-186.   | 6.7  | 80        |
| 14 | A Comprehensive DNA Methylation Profile of Epithelial-to-Mesenchymal Transition. <i>Cancer Research</i> , 2014, 74, 5608-5619.   | 0.9  | 69        |
| 15 | S-adenosylmethionine Levels Regulate the Schwann Cell DNA Methylome. <i>Neuron</i> , 2014, 81, 1024-1039.  | 8.1  | 67        |
| 16 | Regulation of DNA Methylation Patterns by CK2-Mediated Phosphorylation of Dnmt3a. <i>Cell Reports</i> , 2014, 8, 743-753.  | 6.4  | 66        |
| 17 | Ethical implications of epigenetics in the era of personalized medicine. <i>Clinical Epigenetics</i> , 2022, 14, 44.   | 4.1  | 61        |
| 18 | DNA Methylomes Reveal Biological Networks Involved in Human Eye Development, Functions and Associated Disorders. <i>Scientific Reports</i> , 2017, 7, 11762.   | 3.3  | 44        |

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|----|--|-----|-----------|
| 19 | Towards a more precise therapy in cancer: Exploring epigenetic complexity. <i>Current Opinion in Chemical Biology</i> , 2020, 57, 41-49.   | 6.1 | 38        |
| 20 | DNA Methylation Plasticity of Human Adipose-Derived Stem Cells in Lineage Commitment. <i>American Journal of Pathology</i> , 2012, 181, 2079-2093.   | 3.8 | 36        |
| 21 | Quantification of Global DNA Methylation by Capillary Electrophoresis and Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2009, 507, 23-34.   | 0.9 | 34        |
| 22 | Effect of genetic ancestry on leukocyte global DNA methylation in cancer patients. <i>BMC Cancer</i> , 2015, 15, 434.  | 2.6 | 28        |
| 23 | Methylation regulation of Antiviral host factors, Interferon Stimulated Genes (ISGs) and T-cell responses associated with natural HIV control. <i>PLoS Pathogens</i> , 2020, 16, e1008678.         | 4.7 | 25        |
| 24 | Towards a druggable epitranscriptome: Compounds that target RNA modifications in cancer. <i>British Journal of Pharmacology</i> , 2022, 179, 2868-2889.  | 5.4 | 19        |
| 25 | <i>In vitro</i> and <i>in vivo</i> activity of a new small-molecule inhibitor of HDAC6 in mantle cell lymphoma. <i>Haematologica</i> , 2018, 103, e537-e540.                                       | 3.5 | 15        |
| 26 | DNA methylation events in transcription factors and gene expression changes in colon cancer. <i>Epigenomics</i> , 2020, 12, 1593-1610.   | 2.1 | 13        |
| 27 | Discovery of novel DNA methylation biomarkers for non-invasive sporadic breast cancer detection in the Latino population. <i>Molecular Oncology</i> , 2021, 15, 473-486.                           | 4.6 | 8         |
| 28 | Epigenetic landscape in the kick-and-kill therapeutic vaccine BCN02 clinical trial is associated with antiretroviral treatment interruption (ATI) outcome. <i>EBioMedicine</i> , 2022, 78, 103956. | 6.1 | 5         |
| 29 | Follow-Up Study Confirms the Presence of Gastric Cancer DNA Methylation Hallmarks in High-Risk Precursor Lesions. <i>Cancers</i> , 2021, 13, 2760.   | 3.7 | 4         |
| 30 | Impact of the Epigenetically Regulated Hoxa-5 Gene in Neural Differentiation from Human Adipose-Derived Stem Cells. <i>Biology</i> , 2021, 10, 802.  | 2.8 | 2         |
| 31 | The human epigenome—implications for the understanding of human disease. , 2020, , 139-148.  |     | 0         |