

# Roger A Greenberg

## List of Publications by Citations

**Source:** <https://exaly.com/author-pdf/7816107/roger-a-greenberg-publications-by-citations.pdf>

**Version:** 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

58  
papers

5,267  
citations

31  
h-index

60  
g-index

60  
ext. papers

6,451  
ext. citations

14.2  
avg, IF

5.9  
L-index

| #  | Paper  | IF   | Citations |
|----|--|------|-----------|
| 58 | Mitotic progression following DNA damage enables pattern recognition within micronuclei. <i>Nature</i> , <b>2017</b> , 548, 466-470  | 50.4 | 659       |
| 57 | RAP80 targets BRCA1 to specific ubiquitin structures at DNA damage sites. <i>Science</i> , <b>2007</b> , 316, 1198-2023  | 33.3 | 547       |
| 56 | ATM-dependent chromatin changes silence transcription in cis to DNA double-strand breaks. <i>Cell</i> , <b>2010</b> , 141, 970-81  | 56.2 | 501       |
| 55 | Acetylation limits 53BP1 association with damaged chromatin to promote homologous recombination. <i>Nature Structural and Molecular Biology</i> , <b>2013</b> , 20, 317-25 | 17.6 | 347       |
| 54 | Break-induced telomere synthesis underlies alternative telomere maintenance. <i>Nature</i> , <b>2016</b> , 539, 54-58  | 50.4 | 240       |
| 53 | Multifactorial contributions to an acute DNA damage response by BRCA1/BARD1-containing complexes. <i>Genes and Development</i> , <b>2006</b> , 20, 34-46                   | 12.6 | 238       |
| 52 | Biallelic mutations in BRCA1 cause a new Fanconi anemia subtype. <i>Cancer Discovery</i> , <b>2015</b> , 5, 135-42   | 24.4 | 215       |
| 51 | Interchromosomal homology searches drive directional ALT telomere movement and synapsis. <i>Cell</i> , <b>2014</b> , 159, 108-121  | 56.2 | 215       |
| 50 | DNA Damage Follows Repair Factor Depletion and Portends Genome Variation in Cancer Cells after Pore Migration. <i>Current Biology</i> , <b>2017</b> , 27, 210-223          | 6.3  | 163       |
| 49 | The BRCA1-RAP80 complex regulates DNA repair mechanism utilization by restricting end resection. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 13669-80      | 5.4  | 162       |
| 48 | DNA-damage-induced type I interferon promotes senescence and inhibits stem cell function. <i>Cell Reports</i> , <b>2015</b> , 11, 785-797                                  | 10.6 | 139       |
| 47 | Diverse and Reversion Mutations in Circulating Cell-Free DNA of Therapy-Resistant Breast or Ovarian Cancer. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 6708-6720  | 12.9 | 132       |
| 46 | Mechanosensing by the Lamina Protects against Nuclear Rupture, DNA Damage, and Cell-Cycle Arrest. <i>Developmental Cell</i> , <b>2019</b> , 49, 920-935.e5                 | 10.2 | 129       |
| 45 | ALTERNATIVE Telomere Maintenance and Cancer. <i>Trends in Cancer</i> , <b>2015</b> , 1, 145-156  | 12.5 | 129       |
| 44 | MERIT40 controls BRCA1-Rap80 complex integrity and recruitment to DNA double-strand breaks. <i>Genes and Development</i> , <b>2009</b> , 23, 740-54                        | 12.6 | 117       |
| 43 | Whole-exome sequencing identifies somatic ATRX mutations in pheochromocytomas and paragangliomas. <i>Nature Communications</i> , <b>2015</b> , 6, 6140                     | 17.4 | 115       |
| 42 | Nuclear Acetyl-CoA Production by ACLY Promotes Homologous Recombination. <i>Molecular Cell</i> , <b>2017</b> , 67, 252-265.e6  | 17.6 | 110       |

|    |  |      |    |
|----|--|------|----|
| 41 | ATM Dependent Silencing Links Nucleolar Chromatin Reorganization to DNA Damage Recognition. <i>Cell Reports</i> , <b>2015</b> , 13, 251-9  | 10.6 | 94 |
| 40 | Noncanonical views of homology-directed DNA repair. <i>Genes and Development</i> , <b>2016</b> , 30, 1138-54   | 12.6 | 92 |
| 39 | HGG-33. PATIENT DERIVED CELL LINES TO STUDY ATRX AND ALT IN PEDIATRIC BRAIN TUMORS. <i>Neuro-Oncology</i> , <b>2018</b> , 20, i96-i96  | 1    | 78 |
| 38 | A BRISC-SHMT complex deubiquitinates IFNAR1 and regulates interferon responses. <i>Cell Reports</i> , <b>2013</b> , 5, 180-93  | 10.6 | 62 |
| 37 | Combining PARP with ATR inhibition overcomes PARP inhibitor and platinum resistance in ovarian cancer models. <i>Nature Communications</i> , <b>2020</b> , 11, 3726                                | 17.4 | 61 |
| 36 | Deciphering the BRCA1 Tumor Suppressor Network. <i>Journal of Biological Chemistry</i> , <b>2015</b> , 290, 17724-17732  | 17.4 | 59 |
| 35 | Differential regulation of JAMM domain deubiquitinating enzyme activity within the RAP80 complex. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 30971-81                             | 5.4  | 57 |
| 34 | A Radiotracer Strategy to Quantify PARP-1 Expression In Vivo Provides a Biomarker That Can Enable Patient Selection for PARP Inhibitor Therapy. <i>Cancer Research</i> , <b>2016</b> , 76, 4516-24 | 10.1 | 55 |
| 33 | RAD52 and SLX4 act nonredundantly to ensure telomere stability during alternative telomere lengthening. <i>Genes and Development</i> , <b>2019</b> , 33, 221-235                                   | 12.6 | 50 |
| 32 | Recognition of DNA double strand breaks by the BRCA1 tumor suppressor network. <i>Chromosoma</i> , <b>2008</b> , 117, 305-17   | 2.8  | 49 |
| 31 | Nuclear body phase separation drives telomere clustering in ALT cancer cells. <i>Molecular Biology of the Cell</i> , <b>2020</b> , 31, 2048-2056   | 3.5  | 44 |
| 30 | Type I interferon controls propagation of long interspersed element-1. <i>Journal of Biological Chemistry</i> , <b>2015</b> , 290, 10191-9   | 5.4  | 44 |
| 29 | ALC1 links chromatin accessibility to PARP inhibitor response in homologous recombination-deficient cells. <i>Nature Cell Biology</i> , <b>2021</b> , 23, 160-171                                  | 23.4 | 37 |
| 28 | Metabolic control of BRISC-SHMT2 assembly regulates immune signalling. <i>Nature</i> , <b>2019</b> , 570, 194-199  | 50.4 | 33 |
| 27 | Higher-Order Assembly of BRCC36-KIAA0157 Is Required for DUB Activity and Biological Function. <i>Molecular Cell</i> , <b>2015</b> , 59, 970-83  | 17.6 | 31 |
| 26 | As a Nucleus Enters a Small Pore, Chromatin Stretches and Maintains Integrity, Even with DNA Breaks. <i>Biophysical Journal</i> , <b>2017</b> , 112, 446-449                                       | 2.9  | 29 |
| 25 | Cell Cycle Checkpoints Cooperate to Suppress DNA- and RNA-Associated Molecular Pattern Recognition and Anti-Tumor Immune Responses. <i>Cell Reports</i> , <b>2020</b> , 32, 108080                 | 10.6 | 29 |
| 24 | [(18)F]FluorThanatrace uptake as a marker of PARP1 expression and activity in breast cancer. <i>American Journal of Nuclear Medicine and Molecular Imaging</i> , <b>2016</b> , 6, 94-101           | 2.2  | 27 |

|    |   |      |    |
|----|---|------|----|
| 23 | Meiosis-specific proteins MEIOB and SPATA22 cooperatively associate with the single-stranded DNA-binding replication protein A complex and DNA double-strand breaks. <i>Biology of Reproduction</i> , <b>2017</b> , 96, 1096-1104 | 3.9  | 26 |
| 22 | Histone tails: Directing the chromatin response to DNA damage. <i>FEBS Letters</i> , <b>2011</b> , 585, 2883-90   | 3.8  | 26 |
| 21 | Targeting PARP-1 with Alpha-Particles Is Potently Cytotoxic to Human Neuroblastoma in Preclinical Models. <i>Molecular Cancer Therapeutics</i> , <b>2019</b> , 18, 1195-1204  | 6.1  | 21 |
| 20 | MERIT40 cooperates with BRCA2 to resolve DNA interstrand cross-links. <i>Genes and Development</i> , <b>2015</b> , 29, 1955-68  | 12.6 | 17 |
| 19 | Choreographing the Double Strand Break Response: Ubiquitin and SUMO Control of Nuclear Architecture. <i>Frontiers in Genetics</i> , <b>2016</b> , 7, 103  | 4.5  | 11 |
| 18 | Case-control analysis of truncating mutations in DNA damage response genes connects TEX15 and FANCD2 with hereditary breast cancer susceptibility. <i>Scientific Reports</i> , <b>2017</b> , 7, 681                               | 4.9  | 10 |
| 17 | The BRISC deubiquitinating enzyme complex limits hematopoietic stem cell expansion by regulating JAK2 K63-ubiquitination. <i>Blood</i> , <b>2019</b> , 133, 1560-1571   | 2.2  | 9  |
| 16 | Assembling a protective shield. <i>Nature Cell Biology</i> , <b>2018</b> , 20, 862-863  | 23.4 | 8  |
| 15 | Chromatin yo-yo: expansion and condensation during DNA repair. <i>Trends in Cell Biology</i> , <b>2014</b> , 24, 616-618  | 18.3 | 6  |
| 14 | MERIT40 deficiency expands hematopoietic stem cell pools by regulating thrombopoietin receptor signaling. <i>Blood</i> , <b>2015</b> , 125, 1730-8  | 2.2  | 6  |
| 13 | DNA repair: Familiar ends with alternative endings. <i>Nature</i> , <b>2015</b> , 518, 174-6  | 50.4 | 6  |
| 12 | Cancer. BRCA1, everything but the RING?. <i>Science</i> , <b>2011</b> , 334, 459-60   | 33.3 | 6  |
| 11 | Phosphorylation of TIP60 Suppresses 53BP1 Localization at DNA Damage Sites. <i>Molecular and Cellular Biology</i> , <b>2019</b> , 39,   | 4.8  | 6  |
| 10 | Direct Quantitative Monitoring of Homology-Directed DNA Repair of Damaged Telomeres. <i>Methods in Enzymology</i> , <b>2018</b> , 600, 107-134  | 1.7  | 5  |
| 9  | The abscopal effect: a sense of DNA damage is in the air. <i>Journal of Clinical Investigation</i> , <b>2021</b> , 131,   | 15.9 | 5  |
| 8  | In vivo imaging of DNA double-strand break induced telomere mobility during alternative lengthening of telomeres. <i>Methods</i> , <b>2017</b> , 114, 54-59   | 4.6  | 4  |
| 7  | Morning for Irofulven, What Could be fiNER?. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 1833-1835  | 12.9 | 3  |
| 6  | Telomere erosion in human pluripotent stem cells leads to ATR-mediated mitotic catastrophe. <i>Journal of Cell Biology</i> , <b>2021</b> , 220,   | 7.3  | 2  |

|   |   |      |   |
|---|---|------|---|
| 5 | Communication between chromatin and homologous recombination. <i>Current Opinion in Genetics and Development</i> , <b>2021</b> , 71, 1-9  | 4.9  | 1 |
| 4 | Putting PHDs to work: PHF11 clears the way for EXO1 in double-strand break repair. <i>Genes and Development</i> , <b>2017</b> , 31, 3-5   | 12.6 |   |
| 3 | David Livingston (1941-2021).. <i>Molecular Cell</i> , <b>2022</b> , 82, 4-7  | 17.6 |   |
| 2 | RNF4-Dependent Hybrid SUMO-Ubiquitin Chains are Signals for RAP80 and thereby Mediate the Recruitment of BRCA1 to Sites of DNA Damage. <i>FASEB Journal</i> , <b>2013</b> , 27, 782.7 | 0.9  |   |
| 1 | The inner workings of replisome-dependent control of DNA damage tolerance.. <i>Genes and Development</i> , <b>2022</b> , 36, 103-105  | 12.6 |   |