

Ari Sadanandom

List of Publications by Year in descending order

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Version: 2024-02-01

60
papers

6,317
citations

117625

34
h-index

138484

58
g-index

63
all docs

63
docs citations

63
times ranked

6316
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Genome sequence and analysis of the Irish potato famine pathogen <i>Phytophthora infestans</i> . <i>Nature</i> , 2009, 461, 393-398. | 27.8 | 1,405 |
| 2 | The RAR1 Interactor SGT1, an Essential Component of R Gene-Triggered Disease Resistance. <i>Science</i> , 2002, 295, 2073-2076. | 12.6 | 574 |
| 3 | <i>Phytophthora infestans</i> effector AVR3a is essential for virulence and manipulates plant immunity by stabilizing host E3 ligase CMPG1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9909-9914. | 7.1 | 412 |
| 4 | Ubiquitin ligase-associated protein SGT1 is required for host and nonhost disease resistance in plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 10865-10869. | 7.1 | 385 |
| 5 | The ubiquitin-proteasome system: central modifier of plant signalling. <i>New Phytologist</i> , 2012, 196, 13-28. | 7.3 | 329 |
| 6 | Role of SGT1 in resistance protein accumulation in plant immunity. <i>EMBO Journal</i> , 2006, 25, 2007-2016. | 7.8 | 226 |
| 7 | RAR1 and NDR1 Contribute Quantitatively to Disease Resistance in Arabidopsis, and Their Relative Contributions Are Dependent on the R Gene Assayed. <i>Plant Cell</i> , 2002, 14, 1005-1015. | 6.6 | 218 |
| 8 | The E3 Ubiquitin Ligase Activity of Arabidopsis PLANT U-BOX17 and Its Functional Tobacco Homolog ACRE276 Are Required for Cell Death and Defense. <i>Plant Cell</i> , 2006, 18, 1084-1098. | 6.6 | 215 |
| 9 | Arabidopsis RAR1 Exerts Rate-Limiting Control of R Gene-Mediated Defenses against Multiple Pathogens. <i>Plant Cell</i> , 2002, 14, 979-992. | 6.6 | 197 |
| 10 | Root branching toward water involves posttranslational modification of transcription factor ARF7. <i>Science</i> , 2018, 362, 1407-1410. | 12.6 | 179 |
| 11 | Small Ubiquitin-Like Modifier Proteases OVERLY TOLERANT TO SALT1 and -2 Regulate Salt Stress Responses in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2008, 20, 2894-2908. | 6.6 | 173 |
| 12 | CMPG1-dependent cell death follows perception of diverse pathogen elicitors at the host plasma membrane and is suppressed by <i>Phytophthora infestans</i> RXLR effector AVR3a. <i>New Phytologist</i> , 2011, 190, 653-666. | 7.3 | 142 |
| 13 | E3 ubiquitin ligases and plant innate immunity. <i>Journal of Experimental Botany</i> , 2009, 60, 1123-1132. | 4.8 | 140 |
| 14 | Small Ubiquitin-like Modifier Protein SUMO Enables Plants to Control Growth Independently of the Phytohormone Gibberellin. <i>Developmental Cell</i> , 2014, 28, 102-110. | 7.0 | 139 |
| 15 | Deubiquitinating enzymes AtUBP12 and AtUBP13 and their tobacco homologue NtUBP12 are negative regulators of plant immunity. <i>New Phytologist</i> , 2011, 191, 92-106. | 7.3 | 94 |
| 16 | Timing is everything: regulatory overlap in plant cell death. <i>Trends in Plant Science</i> , 2008, 13, 589-595. | 8.8 | 93 |
| 17 | Towards understanding the virulence functions of RXLR effectors of the oomycete plant pathogen <i>Phytophthora infestans</i> . <i>Journal of Experimental Botany</i> , 2009, 60, 1133-1140. | 4.8 | 92 |
| 18 | Cauliflower mosaic virus protein P6 is a suppressor of RNA silencing. <i>Journal of General Virology</i> , 2007, 88, 3439-3444. | 2.9 | 81 |

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|----|--|------|-----------|
| 19 | Dealing With Stress: A Review of Plant SUMO Proteases. <i>Frontiers in Plant Science</i> , 2019, 10, 1122. | 3.6 | 71 |
| 20 | SUMOylation of phytochrome-B negatively regulates light-induced signaling in <i>Arabidopsis thaliana</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 11108-11113. | 7.1 | 69 |
| 21 | SUMO Is a Critical Regulator of Salt Stress Responses in Rice. <i>Plant Physiology</i> , 2016, 170, 2378-2391. | 4.8 | 63 |
| 22 | SUMO proteases: uncovering the roles of deSUMOylation in plants. <i>Journal of Experimental Botany</i> , 2016, 67, 2541-2548. | 4.8 | 61 |
| 23 | Rice <i>Os</i> SUMO protease <i>Os</i> Overly Tolerant to Salt 1 targets the transcription factor, <i>Os</i> ZIP23 to promote drought tolerance in rice. <i>Plant Journal</i> , 2017, 92, 1031-1043. | 5.7 | 59 |
| 24 | SUMOylation represses SnRK1 signaling in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2016, 85, 120-133. | 5.7 | 56 |
| 25 | Identification of Transgene-Free CRISPR-Edited Plants of Rice, Tomato, and <i>Arabidopsis</i> by Monitoring DsRED Fluorescence in Dry Seeds. <i>Frontiers in Plant Science</i> , 2019, 10, 1150. | 3.6 | 56 |
| 26 | SUMO conjugation to the pattern recognition receptor FLS2 triggers intracellular signalling in plant innate immunity. <i>Nature Communications</i> , 2018, 9, 5185. | 12.8 | 55 |
| 27 | <i>DAY NEUTRAL FLOWERING</i> Represses <i>CONSTANS</i> to Prevent <i>Arabidopsis</i> Flowering Early in Short Days. <i>Plant Cell</i> , 2010, 22, 1118-1128. | 6.6 | 50 |
| 28 | Stability of small ubiquitin-like modifier (SUMO) proteases <i>OVERLY TOLERANT TO SALT1</i> and -2 modulates salicylic acid signalling and SUMO1/2 conjugation in <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2016, 67, 353-363. | 4.8 | 48 |
| 29 | SUMO Conjugation to BZR1 Enables Brassinosteroid Signaling to Integrate Environmental Cues to Shape Plant Growth. <i>Current Biology</i> , 2020, 30, 1410-1423.e3. | 3.9 | 48 |
| 30 | U-box E3 ubiquitin ligase PUB17 acts in the nucleus to promote specific immune pathways triggered by <i>Phytophthora infestans</i> . <i>Journal of Experimental Botany</i> , 2015, 66, 3189-3199. | 4.8 | 47 |
| 31 | Fifty shades of SUMO: its role in immunity and at the fulcrum of the growth-defence balance. <i>Molecular Plant Pathology</i> , 2018, 19, 1537-1544. | 4.2 | 45 |
| 32 | Biosensors in plants. <i>Current Opinion in Plant Biology</i> , 2010, 13, 736-743. | 7.1 | 43 |
| 33 | SUMO Suppresses the Activity of the Jasmonic Acid Receptor <i>CORONATINE INSENSITIVE1</i> . <i>Plant Cell</i> , 2018, 30, 2099-2115. | 6.6 | 43 |
| 34 | BTB-BACK Domain Protein POB1 Suppresses Immune Cell Death by Targeting Ubiquitin E3 ligase PUB17 for Degradation. <i>PLoS Genetics</i> , 2017, 13, e1006540. | 3.5 | 41 |
| 35 | Ubiquitin chain topology in plant cell signaling: a new facet to an evergreen story. <i>Frontiers in Plant Science</i> , 2014, 5, 122. | 3.6 | 35 |
| 36 | Post-translational modifications in priming the plant immune system: ripe for exploitation?. <i>FEBS Letters</i> , 2018, 592, 1929-1936. | 2.8 | 31 |

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|----|---|-----|-----------|
| 37 | OTS1 and OTS2 SUMO proteases link plant development and survival under salt stress. <i>Plant Signaling and Behavior</i> , 2009, 4, 225-227. | 2.4 | 23 |
| 38 | SUMO mediated regulation of transcription factors as a mechanism for transducing environmental cues into cellular signaling in plants. <i>Cellular and Molecular Life Sciences</i> , 2021, 78, 2641-2664. | 5.4 | 21 |
| 39 | SUMO enables substrate selectivity by mitogen-activated protein kinases to regulate immunity in plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, . | 7.1 | 21 |
| 40 | Revised nomenclature and functional overview of the ULP gene family of plant deSUMOylating proteases. <i>Journal of Experimental Botany</i> , 2018, 69, 4505-4509. | 4.8 | 20 |
| 41 | Rice OVERLY TOLERANT TO SALT 1 (OTS1) SUMO protease is a positive regulator of seed germination and root development. <i>Plant Signaling and Behavior</i> , 2016, 11, e1173301. | 2.4 | 19 |
| 42 | SUMO proteases OTS1 and 2 control filament elongation through a DELLA-dependent mechanism. <i>Plant Reproduction</i> , 2016, 29, 287-290. | 2.2 | 17 |
| 43 | A functional Small Ubiquitin-like Modifier (SUMO) interacting motif (SIM) in the gibberellin hormone receptor GID1 is conserved in cereal crops and disrupting this motif does not abolish hormone dependency of the DELLA-GID1 interaction. <i>Plant Signaling and Behavior</i> , 2015, 10, e987528. | 2.4 | 16 |
| 44 | Functional analysis of a <i>Wheat</i> heat shock domain protein, <i>TaR1</i> , reveals that host chromatin remodelling influences the dynamics of the switch to necrotrophic growth in the phytopathogenic fungus <i>Zymoseptoria tritici</i> . <i>New Phytologist</i> , 2015, 206, 598-605. | 7.3 | 16 |
| 45 | Ubiquitination in plant nutrient utilization. <i>Frontiers in Plant Science</i> , 2013, 4, 452. | 3.6 | 15 |
| 46 | Exploiting protein modification systems to boost crop productivity: SUMO proteases in focus. <i>Journal of Experimental Botany</i> , 2018, 69, 4625-4632. | 4.8 | 14 |
| 47 | Understanding and Exploiting Post-Translational Modifications for Plant Disease Resistance. <i>Biomolecules</i> , 2021, 11, 1122. | 4.0 | 14 |
| 48 | Destabilization of interaction between cytokinin signaling intermediates <i>AHP1</i> and <i>ARR4</i> modulates <i>Arabidopsis</i> development. <i>New Phytologist</i> , 2015, 206, 726-737. | 7.3 | 13 |
| 49 | Towards understanding the multifaceted role of <i>SUMOylation</i> in plant growth and development. <i>Physiologia Plantarum</i> , 2021, 171, 77-85. | 5.2 | 13 |
| 50 | CHPA, a Cysteine- and Histidine-Rich-Domain-Containing Protein, Contributes to Maintenance of the Diploid State in <i>Aspergillus nidulans</i> . <i>Eukaryotic Cell</i> , 2004, 3, 984-991. | 3.4 | 11 |
| 51 | An Insight into the Factors Influencing Specificity of the SUMO System in Plants. <i>Plants</i> , 2020, 9, 1788. | 3.5 | 11 |
| 52 | The conjugation of SUMO to the transcription factor MYC2 functions in blue light-mediated seedling development in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2022, 34, 2892-2906. | 6.6 | 8 |
| 53 | Plant proteostasis “shaping the proteome: a research community aiming to understand molecular mechanisms that control protein abundance. <i>New Phytologist</i> , 2020, 227, 1028-1033. | 7.3 | 7 |
| 54 | Response to Cacas and Diamond: Is the autophagy machinery an executioner of programmed cell death in plants?. <i>Trends in Plant Science</i> , 2009, 14, 300-301. | 8.8 | 5 |

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|----|---|-----|-----------|
| 55 | Structure and Mechanism of Dimer→Monomer Transition of a Plant Poly(A)-Binding Protein upon RNA Interaction: Insights into Its Poly(A) Tail Assembly. <i>Journal of Molecular Biology</i> , 2015, 427, 2491-2506. | 4.2 | 5 |
| 56 | HEARTBREAK Controls Post-translational Modification of INDEHISCENT to Regulate Fruit Morphology in <i>Capsella</i> . <i>Current Biology</i> , 2020, 30, 3880-3888.e5. | 3.9 | 5 |
| 57 | TaWRKY10 transcription factor is a novel jasmonic acid signalling regulator involved in immunity against <i>Septoria tritici</i> blotch disease in wheat. <i>Plant Pathology</i> , 2021, 70, 1397-1408. | 2.4 | 3 |
| 58 | Preface. <i>Journal of Experimental Botany</i> , 2009, 60, 1083-1083. | 4.8 | 0 |
| 59 | A Tropical Plant with Friends in Cold Places: The Formation of the UK Rice Research Community. <i>Trends in Plant Science</i> , 2020, 25, 421-422. | 8.8 | 0 |
| 60 | MARylation meets ubiquitination in the ART of plant immunity. <i>Molecular Cell</i> , 2021, 81, 4572-4574. | 9.7 | 0 |