

# Andrew F Bent

## List of Publications by Year in descending order

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47006

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times ranked

22794  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Exploring Soybean Resistance to Soybean Cyst Nematode. Annual Review of Phytopathology, 2022, 60, 379-409.   | 7.8  | 10        |
| 2  | Soybean Cyst Nematode Resistance Quantitative Trait Locus <i>ScqSCN-006</i> Alters the Expression of a $\beta$ -SNAP Protein. Molecular Plant-Microbe Interactions, 2021, 34, 1433-1445.   | 2.6  | 10        |
| 3  | Coordinated regulation of plant immunity by poly(ADP-ribosyl)ation and K63-linked ubiquitination. Molecular Plant, 2021, 14, 2088-2103.  | 8.3  | 14        |
| 4  | Detection of rare nematode resistance <i>Rhg1</i> haplotypes in Glycine soja and a novel <i>Rhg1</i> $\beta$ -SNAP. Plant Genome, 2021, , e20152.  | 2.8  | 1         |
| 5  | Soybean Resistance Locus <i>Rhg1</i> Confers Resistance to Multiple Cyst Nematodes in Diverse Plant Species. Phytopathology, 2019, 109, 2107-2115.   | 2.2  | 16        |
| 6  | The <i>rhg1</i> ( <i>Rhg1</i> low-copy) nematode resistance source harbors a copia family retrotransposon within the <i>Rhg1</i> -encoded $\beta$ -SNAP gene. Plant Direct, 2019, 3, e00164.                                       | 1.9  | 27        |
| 7  | Agrobacterium-mediated vacuum infiltration and floral dip transformation of rapid-cycling Brassica rapa. BMC Plant Biology, 2019, 19, 246.   | 3.6  | 18        |
| 8  | An atypical N-ethylmaleimide sensitive factor enables the viability of nematode-resistant <i>Rhg1</i> soybeans. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E4512-E4521.           | 7.1  | 58        |
| 9  | 3-Aminobenzamide Blocks MAMP-Induced Callose Deposition Independently of Its Poly(ADP-ribose) Inhibiting Activity. Frontiers in Plant Science, 2018, 9, 1907.  | 3.6  | 10        |
| 10 | A transcriptomics approach uncovers novel roles for poly(ADP-ribose)ation in the basal defense response in Arabidopsis thaliana. PLoS ONE, 2017, 12, e0190268.   | 2.5  | 16        |
| 11 | Directed Evolution of FLS2 towards Novel Flagellin Peptide Recognition. PLoS ONE, 2016, 11, e0157155.  | 2.5  | 11        |
| 12 | Disease resistance through impairment of $\beta$ -SNAP-NSF interaction and vesicular trafficking by soybean <i>Rhg1</i> . Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E7375-E7382. | 7.1  | 71        |
| 13 | Resistance from relatives. Nature Biotechnology, 2016, 34, 620-621.  | 17.5 | 10        |
| 14 | Rice OsFLS2-Mediated Perception of Bacterial Flagellins Is Evaded by Xanthomonas oryzae pvs. oryzae and orycolicola. Molecular Plant, 2015, 8, 1024-1037.  | 8.3  | 60        |
| 15 | PARP2 Is the Predominant Poly(ADP-Ribose) Polymerase in Arabidopsis DNA Damage and Immune Responses. PLoS Genetics, 2015, 11, e1005200.  | 3.5  | 90        |
| 16 | Microbial Pathogens Trigger Host DNA Double-Strand Breaks Whose Abundance Is Reduced by Plant Defense Responses. PLoS Pathogens, 2014, 10, e1004030.   | 4.7  | 99        |
| 17 | Distinct Copy Number, Coding Sequence, and Locus Methylation Patterns Underlie <i>Rhg1</i> -Mediated Soybean Resistance to Soybean Cyst Nematode. Plant Physiology, 2014, 165, 630-647.  | 4.8  | 136       |
| 18 | FLS2-BAK1 Extracellular Domain Interaction Sites Required for Defense Signaling Activation. PLoS ONE, 2014, 9, e111185.  | 2.5  | 23        |

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|----|--|------|-----------|
| 19 | Mutations in FLS2 Ser-938 Dissect Signaling Activation in FLS2-Mediated Arabidopsis Immunity. <i>PLoS Pathogens</i> , 2013, 9, e1003313.   | 4.7  | 57        |
| 20 | FLS2-Mediated Responses to Ax21-Derived Peptides: Response to the Mueller et al. Commentary. <i>Plant Cell</i> , 2012, 24, 3174-3176.  | 6.6  | 5         |
| 21 | Copy Number Variation of Multiple Genes at <i>Rhg1</i> Mediates Nematode Resistance in Soybean. <i>Science</i> , 2012, 338, 1206-1209.   | 12.6 | 535       |
| 22 | Probing the Arabidopsis Flagellin Receptor: FLS2-FLS2 Association and the Contributions of Specific Domains to Signaling Function. <i>Plant Cell</i> , 2012, 24, 1096-1113.  | 6.6  | 104       |
| 23 | Pathogens Drop the Hint: Don't Forget Phytoalexin Pathways. <i>Cell Host and Microbe</i> , 2011, 9, 169-170.   | 11.0 | 2         |
| 24 | Poly(ADP-ribosyl)ation in plants. <i>Trends in Plant Science</i> , 2011, 16, 372-380.  | 8.8  | 94        |
| 25 | LRR Conservation Mapping to Predict Functional Sites within Protein Leucine-Rich Repeat Domains. <i>PLoS ONE</i> , 2011, 6, e21614.  | 2.5  | 46        |
| 26 | Type III secretion-dependent host defence elicitation and type III secretion-independent growth within leaves by <i>Xanthomonas campestris</i> pv. <i>campestris</i> . <i>Molecular Plant Pathology</i> , 2011, 12, 731-745.       | 4.2  | 20        |
| 27 | Arabidopsis TTR1 Causes LRR-Dependent Lethal Systemic Necrosis, rather than Systemic Acquired Resistance, to Tobacco Ringspot Virus. <i>Molecules and Cells</i> , 2011, 32, 421-430.   | 2.6  | 17        |
| 28 | The Arabidopsis flagellin receptor FLS2 mediates the perception of Xanthomonas Ax21 secreted peptides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 9286-9291.              | 7.1  | 62        |
| 29 | A nematode demographics assay in transgenic roots reveals no significant impacts of the Rhg1 locus LRR-Kinase on soybean cyst nematode resistance. <i>BMC Plant Biology</i> , 2010, 10, 104.                                       | 3.6  | 77        |
| 30 | Fine Mapping of the SCN Resistance Locus <i>rhg1</i> from PI 88788. <i>Plant Genome</i> , 2010, 3, .   | 2.8  | 56        |
| 31 | Underexplored Niches in Research on Plant Pathogenic Bacteria. <i>Plant Physiology</i> , 2009, 150, 1631-1637.   | 4.8  | 17        |
| 32 | Disruption of Poly(ADP-ribosyl)ation Mechanisms Alters Responses of Arabidopsis to Biotic Stress. <i>Plant Physiology</i> , 2009, 152, 267-280.  | 4.8  | 118       |
| 33 | Signaling Pathways That Regulate the Enhanced Disease Resistance of Arabidopsis Defense, No Death Mutants. <i>Molecular Plant-Microbe Interactions</i> , 2008, 21, 1285-1296.  | 2.6  | 92        |
| 34 | Discovery of ADP-Ribosylation and Other Plant Defense Pathway Elements Through Expression Profiling of Four Different Arabidopsis Pseudomonas R-avr Interactions. <i>Molecular Plant-Microbe Interactions</i> , 2008, 21, 646-657. | 2.6  | 57        |
| 35 | MEKK1 Is Required for flg22-Induced MPK4 Activation in Arabidopsis Plants. <i>Plant Physiology</i> , 2007, 143, 661-669.   | 4.8  | 306       |
| 36 | Identification and Mutational Analysis of Arabidopsis FLS2 Leucine-Rich Repeat Domain Residues That Contribute to Flagellin Perception. <i>Plant Cell</i> , 2007, 19, 3297-3313.   | 6.6  | 97        |

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|----|---|------|-----------|
| 37 | Elicitors, Effectors, and R Genes: The New Paradigm and a Lifetime Supply of Questions. Annual Review of Phytopathology, 2007, 45, 399-436.   | 7.8  | 668       |
| 38 | Global expression analysis of nucleotide binding site-leucine rich repeat-encoding and related genes in Arabidopsis. BMC Plant Biology, 2007, 7, 56.  | 3.6  | 166       |
| 39 | <i>Arabidopsis thaliana</i> Floral Dip Transformation Method. , 2006, 343, 87-104.  |      | 155       |
| 40 | Within-Species Flagellin Polymorphism in <i>Xanthomonas campestris</i> pv <i>campestris</i> and Its Impact on Elicitation of Arabidopsis FLAGELLIN SENSING2-Dependent Defenses. Plant Cell, 2006, 18, 764-779.    | 6.6  | 181       |
| 41 | Disease- and Performance-Related Traits of Ethylene-Sensitive Soybean. Crop Science, 2006, 46, 893-901.   | 1.8  | 12        |
| 42 | Flagellin Is Not a Major Defense Elicitor in <i>Ralstonia solanacearum</i> Cells or Extracts Applied to <i>Arabidopsis thaliana</i> . Molecular Plant-Microbe Interactions, 2004, 17, 696-706.                    | 2.6  | 111       |
| 43 | Identification and functional analysis of Arabidopsis proteins that interact with resistance gene product RPS2 in yeast. Physiological and Molecular Plant Pathology, 2004, 65, 257-267.                          | 2.5  | 14        |
| 44 | Arabidopsis DND2, a Second Cyclic Nucleotide-Gated Ion Channel Gene for Which Mutation Causes the "Defense, No Death" Phenotype. Molecular Plant-Microbe Interactions, 2004, 17, 511-520.                         | 2.6  | 190       |
| 45 | Deciphering host resistance and pathogen virulence: the Arabidopsis / <i>Pseudomonas</i> interaction as a model. Molecular Plant Pathology, 2003, 4, 517-530.   | 4.2  | 57        |
| 46 | A Cyclic Nucleotide-Gated Ion Channel, CNGC2, Is Crucial for Plant Development and Adaptation to Calcium Stress. Plant Physiology, 2003, 132, 728-731.  | 4.8  | 106       |
| 47 | AGRICULTURE: Reconnecting Farms and Ecosystems- If It Pays. Science, 2002, 298, 1340-1341.  | 12.6 | 2         |
| 48 | Probing plant-pathogen interactions and downstream defense signaling using DNA microarrays. Functional and Integrative Genomics, 2002, 2, 259-273.  | 3.5  | 102       |
| 49 | Molecular Markers Linked to Brown Stem Rot Resistance Genes, <i>Rbs1</i> and <i>Rbs2</i> , in Soybean. Crop Science, 2001, 41, 527-535.   | 1.8  | 53        |
| 50 | Plant mitogen-activated protein kinase cascades: Negative regulatory roles turn out positive. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 784-786.                 | 7.1  | 65        |
| 51 | The Leucine-Rich Repeat Domain Can Determine Effective Interaction Between <i>RPS2</i> and Other Host Factors in Arabidopsis <i>RPS2</i> -Mediated Disease Resistance. Genetics, 2001, 158, 439-450.              | 2.9  | 66        |
| 52 | A Second T-Region of the Soybean-Supervirulent Chrysope-Type Ti Plasmid pTiChry5, and Construction of a Fully Disarmed vir Helper Plasmid. Molecular Plant-Microbe Interactions, 2000, 13, 1081-1091.             | 2.6  | 47        |
| 53 | Identification of Arabidopsis Mutants Exhibiting an Altered Hypersensitive Response in Gene-for-Gene Disease Resistance. Molecular Plant-Microbe Interactions, 2000, 13, 277-286.                                 | 2.6  | 51        |
| 54 | The Arabidopsis <i>dnd1</i> "defense, no death" gene encodes a mutated cyclic nucleotide-gated ion channel. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 9323-9328. | 7.1  | 523       |

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|----|---|------|-----------|
| 55 | Arabidopsis in Planta Transformation. Uses, Mechanisms, and Prospects for Transformation of Other Species. <i>Plant Physiology</i> , 2000, 124, 1540-1547.  | 4.8  | 190       |
| 56 | Female Reproductive Tissues Are the Primary Target of <i>Agrobacterium</i> -Mediated Transformation by the Arabidopsis Floral-Dip Method <sup>1</sup> . <i>Plant Physiology</i> , 2000, 123, 895-904.                         | 4.8  | 237       |
| 57 | Applications of Molecular Biology to Plant Disease and Insect Resistance. <i>Advances in Agronomy</i> , 1999, , 251-298.  | 5.2  | 18        |
| 58 | Isolation of Ethylene-Insensitive Soybean Mutants That Are Altered in Pathogen Susceptibility and Gene-for-Gene Disease Resistance <sup>1</sup> . <i>Plant Physiology</i> , 1999, 119, 935-950.                               | 4.8  | 187       |
| 59 | Regulation of Soybean Nodulation Independent of Ethylene Signaling <sup>1</sup> . <i>Plant Physiology</i> , 1999, 119, 951-960.   | 4.8  | 105       |
| 60 | Floral dip: a simplified method for <i>Agrobacterium</i> -mediated transformation of <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 1998, 16, 735-743.  | 5.7  | 19,148    |
| 61 | Plant disease reality. <i>Trends in Plant Science</i> , 1998, 3, 405-406.   | 8.8  | 0         |
| 62 | Gene-for-gene disease resistance without the hypersensitive response in Arabidopsis dnd1 mutant. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 7819-7824.                | 7.1  | 432       |
| 63 | <i>Agrobacterium</i> Germ-Line Transformation: Transformation of Arabidopsis without Tissue Culture. , 1998, , 17-30.   |      | 30        |
| 64 | Plant Disease Resistance Genes: Function Meets Structure. <i>Plant Cell</i> , 1996, 8, 1757.  | 6.6  | 93        |
| 65 | Identification and Map Location of TTR1, a Single Locus in <i>Arabidopsis thaliana</i> that Confers Tolerance to Tobacco Ringspot Nepovirus. <i>Molecular Plant-Microbe Interactions</i> , 1996, 9, 729.                      | 2.6  | 40        |
| 66 | RPS2 of Arabidopsis thaliana: a leucine-rich repeat class of plant disease resistance genes. <i>Science</i> , 1994, 265, 1856-1860.   | 12.6 | 929       |
| 67 | Genetic Analysis of Bacterial Disease Resistance in Arabidopsis and Cloning of the RPS2 Resistance Gene. <i>Current Plant Science and Biotechnology in Agriculture</i> , 1994, , 283-288.                                     | 0.0  | 0         |
| 68 | Identification of a disease resistance locus in Arabidopsis that is functionally homologous to the RPG1 locus of soybean. <i>Plant Journal</i> , 1993, 4, 813-820.  | 5.7  | 92        |
| 69 | RPS2, an Arabidopsis Disease Resistance Locus Specifying Recognition of Pseudomonas syringae Strains Expressing the Avirulence Gene avrRpt2. <i>Plant Cell</i> , 1993, 5, 865.  | 6.6  | 5         |
| 70 | RPS2, an Arabidopsis disease resistance locus specifying recognition of Pseudomonas syringae strains expressing the avirulence gene avrRpt2.. <i>Plant Cell</i> , 1993, 5, 865-875.   | 6.6  | 303       |
| 71 | Molecular analysis of avirulence gene avrRpt2 and identification of a putative regulatory sequence common to all known Pseudomonas syringae avirulence genes. <i>Journal of Bacteriology</i> , 1993, 175, 4859-4869.          | 2.2  | 196       |
| 72 | Disease Development in Ethylene-Insensitive <i>Arabidopsis thaliana</i> Infected with Virulent and Avirulent <i>Pseudomonas</i> and <i>Xanthomonas</i> Pathogens. <i>Molecular Plant-Microbe Interactions</i> , 1992, 5, 372. | 2.6  | 252       |

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|----|---|-----|-----------|
| 73 | Identification of <i>Pseudomonas syringae</i> pathogens of <i>Arabidopsis</i> and a bacterial locus determining avirulence on both <i>Arabidopsis</i> and soybean.. <i>Plant Cell</i> , 1991, 3, 49-59. | 6.6 | 632       |
| 74 | <i>Arabidopsis</i> as a Model System for Studying Plant Disease Resistance Mechanisms. <i>Annals of the New York Academy of Sciences</i> , 1991, 646, 228-230.  | 3.8 | 0         |
| 75 | Identification of <i>Pseudomonas syringae</i> Pathogens of <i>Arabidopsis</i> and a Bacterial Locus Determining Avirulence on Both <i>Arabidopsis</i> and Soybean. <i>Plant Cell</i> , 1991, 3, 49.     | 6.6 | 137       |
| 76 | <i>Rhizobium meliloti</i> <i>suhR</i> suppresses the phenotype of an <i>Escherichia coli</i> RNA polymerase sigma 32 mutant. <i>Journal of Bacteriology</i> , 1990, 172, 3559-3568.                     | 2.2 | 1         |
| 77 | Induction of Lactate Dehydrogenase Isozymes by Oxygen Deficit in Barley Root Tissue. <i>Plant Physiology</i> , 1986, 82, 658-663.   | 4.8 | 123       |