

# Rongmei Wu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/494684/publications.pdf>

Version: 2024-02-01

12  
papers

1,813  
citations

759233

12  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

2699  
citing authors

#	ARTICLE	IF	CITATIONS
1	Protocol: a highly sensitive RT-PCR method for detection and quantification of microRNAs. <i>Plant Methods</i> , 2007, 3, 12.	4.3	1,048
2	Analysis of expressed sequence tags from <i>Actinidia</i> : applications of a cross species EST database for gene discovery in the areas of flavor, health, color and ripening. <i>BMC Genomics</i> , 2008, 9, 351.	2.8	178
3	A manually annotated <i>Actinidia chinensis</i> var. <i>chinensis</i> (kiwifruit) genome highlights the challenges associated with draft genomes and gene prediction in plants. <i>BMC Genomics</i> , 2018, 19, 257.	2.8	167
4	SVP-like MADS Box Genes Control Dormancy and Budbreak in Apple. <i>Frontiers in Plant Science</i> , 2017, 08, 477.	3.6	121
5	Kiwifruit SVP2 gene prevents premature budbreak during dormancy. <i>Journal of Experimental Botany</i> , 2017, 68, 1071-1082.	4.8	62
6	Overexpression of the kiwifruit SVP3 gene affects reproductive development and suppresses anthocyanin biosynthesis in petals, but has no effect on vegetative growth, dormancy, or flowering time. <i>Journal of Experimental Botany</i> , 2014, 65, 4985-4995.	4.8	59
7	Kiwifruit floral gene APETALA2 is alternatively spliced and accumulates in aberrant indeterminate flowers in the absence of miR172. <i>Plant Molecular Biology</i> , 2012, 78, 417-429.	3.9	51
8	Identification and characterization of flowering genes in kiwifruit: sequence conservation and role in kiwifruit flower development. <i>BMC Plant Biology</i> , 2011, 11, 72.	3.6	43
9	Histone modification and activation by SOC1-like and drought stress-related transcription factors may regulate AcSVP2 expression during kiwifruit winter dormancy. <i>Plant Science</i> , 2019, 281, 242-250.	3.6	28
10	A MADS-box gene with similarity to <i>FLC</i> is induced by cold and correlated with epigenetic changes to control budbreak in kiwifruit. <i>New Phytologist</i> , 2022, 233, 2111-2126.	7.3	25
11	Kiwifruit SVP2 controls developmental and drought-stress pathways. <i>Plant Molecular Biology</i> , 2018, 96, 233-244.	3.9	17
12	Overexpression of both AcSVP1 and AcSVP4 delays budbreak in kiwifruit <i>A. chinensis</i> var. <i>deliciosa</i> , but only AcSVP1 delays flowering in model plants. <i>Environmental and Experimental Botany</i> , 2018, 153, 262-270.	4.2	14