

# Caixi Zhang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8907539/publications.pdf>

Version: 2024-02-01

7  
papers

115  
citations

1478505

6  
h-index

1720034

7  
g-index

7  
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7  
docs citations

7  
times ranked

118  
citing authors

#	ARTICLE	IF	CITATIONS
1	FRUITFULL is involved in double fruit formation at high temperature in sweet cherry. <i>Environmental and Experimental Botany</i> , 2022, 201, 104986.	4.2	3
2	SVP-like gene PavSVP potentially suppressing flowering with PavSEP, PavAP1, and PavJONITLESS in sweet cherries ( <i>Prunus avium</i> L.). <i>Plant Physiology and Biochemistry</i> , 2021, 159, 277-284.	5.8	20
3	Dormancy-Associated MADS-Box (DAM) Genes Influence Chilling Requirement of Sweet Cherries and Co-Regulate Flower Development with SOC1 Gene. <i>International Journal of Molecular Sciences</i> , 2020, 21, 921.	4.1	34
4	MADS-Box Genes are Involved in Cultivar- and Temperature-Dependent Formation of Multi-pistil and Polycarpy in <i>Prunus avium</i> L.. <i>Journal of Plant Growth Regulation</i> , 2019, 38, 1017-1027.	5.1	6
5	Hydrogen cyanamide improves endodormancy release and blooming associated with endogenous hormones in "Summit"™ sweet cherry trees. <i>New Zealand Journal of Crop and Horticultural Science</i> , 2017, 45, 14-28.	1.3	14
6	Proteomic analysis of pear ( <i>Pyrus pyrifolia</i> ) ripening process provides new evidence for the sugar/acid metabolism difference between core and mesocarp. <i>Proteomics</i> , 2016, 16, 3025-3041.	2.2	16
7	Root restriction affected anthocyanin composition and up-regulated the transcription of their biosynthetic genes during berry development in "Summer Black"™ grape. <i>Acta Physiologiae Plantarum</i> , 2013, 35, 2205-2217.	2.1	22