

# Gaihua Qin

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8725303/publications.pdf>

Version: 2024-02-01

13  
papers

1,200  
citations

1039880

9  
h-index

1199470

12  
g-index

13  
all docs

13  
docs citations

13  
times ranked

1348  
citing authors

#	ARTICLE	IF	CITATIONS
1	The genome of the pear ( <i>Pyrus bretschneideri</i> Rehd.). <i>Genome Research</i> , 2013, 23, 396-408.	2.4	832
2	The pomegranate ( <i>Punica granatum</i> L.) genome and the genomics of punicalagin biosynthesis. <i>Plant Journal</i> , 2017, 91, 1108-1128.	2.8	109
3	Evaluation of the volatile profile of 33 <i>Pyrus ussuriensis</i> cultivars by HS-SPME with GC-MS. <i>Food Chemistry</i> , 2012, 134, 2367-2382.	4.2	83
4	Evolution of the Aroma Volatiles of Pear Fruits Supplemented with Fatty Acid Metabolic Precursors. <i>Molecules</i> , 2014, 19, 20183-20196.	1.7	41
5	Diversity of metabolite accumulation patterns in inner and outer seed coats of pomegranate: exploring their relationship with genetic mechanisms of seed coat development. <i>Horticulture Research</i> , 2020, 7, 10.	2.9	38
6	Molecular cloning and expression analysis of a gene for sucrose transporter from pear ( <i>Pyrus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542	2.8	32
7	Calcium treatments promote the aroma volatiles emission of pear ( <i>Pyrus ussuriensis</i> "Nanguoli"™) fruit during post-harvest ripening process. <i>Scientia Horticulturae</i> , 2017, 215, 102-111.	1.7	27
8	Transcriptome profiling reveals the candidate genes associated with aroma metabolites and emission of pear ( <i>Pyrus ussuriensis</i> cv.). <i>Scientia Horticulturae</i> , 2016, 206, 33-42.	1.7	15
9	Genome-wide distribution of simple sequence repeats in pomegranate and their application to the analysis of genetic diversity. <i>Tree Genetics and Genomes</i> , 2020, 16, 1.	0.6	10
10	Genome-wide identification of candidate aquaporins involved in water accumulation of pomegranate outer seed coat. <i>PeerJ</i> , 2021, 9, e11810.	0.9	6
11	Transcriptome analysis of colouration-related genes in two white-fleshed nectarine varieties and their yellow-fleshed mutants. <i>Biotechnology and Biotechnological Equipment</i> , 2018, 32, 899-907.	0.5	4
12	Identification of Candidate Auxin Response Factors Involved in Pomegranate Seed Coat Development. <i>Frontiers in Plant Science</i> , 2020, 11, 536530.	1.7	3
13	Role of membrane lipid hydrolysis genes in the aroma formation of Chinese white pear "Xiang Mian Li"™. <i>Indian Journal of Horticulture</i> , 2020, 77, 56.	0.1	0