## Youn-Sung Kim

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6712284/publications.pdf

Version: 2024-02-01

	1478505	1474206	
88	6	9	
citations	h-index	g-index	
9	9	91	
docs citations	times ranked	citing authors	
	citations 9	88 6 citations h-index	

#	Article	IF	CITATIONS
1	Identification of Flowering-Related Genes Responsible for Differences in Bolting Time between Two Radish Inbred Lines. Frontiers in Plant Science, 2016, 7, 1844.	3.6	26
2	Gibberellin Promotes Bolting and Flowering via the Floral Integrators RsFT and RsSOC1-1 under Marginal Vernalization in Radish. Plants, 2020, 9, 594.	3.5	13
3	Genome-wide identification of flowering time genes associated with vernalization and the regulatory flowering networks in Chinese cabbage. Plant Biotechnology Reports, 2018, 12, 347-363.	1.5	10
4	Nitrogen Signaling Genes and SOC1 Determine the Flowering Time in a Reciprocal Negative Feedback Loop in Chinese Cabbage (Brassica rapa L.) Based on CRISPR/Cas9-Mediated Mutagenesis of Multiple BrSOC1 Homologs. International Journal of Molecular Sciences, 2021, 22, 4631.	4.1	10
5	An Arabidopsis homologue of human seven-in-absentia-interacting protein is involved in pathogen resistance. Molecules and Cells, 2006, 21, 389-94.	2.6	10
6	The Arabidopsis cyclophilin CYP18-1 facilitates PRP18 dephosphorylation and the splicing of introns retained under heat stress. Plant Cell, 2022, 34, 2383-2403.	6.6	10
7	A single amino acid insertion in LCYB2 deflects carotenoid biosynthesis in red carrot. Plant Cell Reports, 2021, 40, 1793-1795.	5.6	5
8	An Arabidopsis GH3 Gene, Encoding an Auxin-Conjugating Enzyme, Mediates Phytochrome B-Regulated Light Signals in Hypocotyl Growth. Plant and Cell Physiology, 2007, 48, 1514-1514.	3.1	3
9	Physiological and molecular characterization of two inbred radish lines with different bolting times. Journal of Plant Biotechnology, 2015, 42, 215-222.	0.4	1