

Identification and Expression Analysis of the *SWEET* *pratensis* Under Abiotic Stresses

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Citation Report

#	ARTICLE	IF	CITATIONS
1	SWEET Transporters and the Potential Functions of These Sequences in Tea (<i>Camellia sinensis</i>). <i>Frontiers in Genetics</i> , 2021, 12, 655843.	2.3	9
2	Plant SWEETs: from sugar transport to plantâ€“pathogen interaction and more unexpected physiological roles. <i>Plant Physiology</i> , 2021, 186, 836-852.	4.8	90
3	Systematic Genome-Wide Study and Expression Analysis of SWEET Gene Family: Sugar Transporter Family Contributes to Biotic and Abiotic Stimuli in Watermelon. <i>International Journal of Molecular Sciences</i> , 2021, 22, 8407.	4.1	21
4	De novo transcriptome sequencing, assembly, and gene expression profiling of a saltâ€“stressed halophyte (<i>Salsola drummondii</i>) from a saline habitat. <i>Physiologia Plantarum</i> , 2021, 173, 1695-1714.	5.2	2
5	Understanding the role of SWEET genes in fruit development and abiotic stress in pomegranate (<i>Punica granatum</i> L.). <i>Molecular Biology Reports</i> , 2022, 49, 1329-1339.	2.3	6
6	Genome-wide identification of the SWEET gene family in <i>Phaseolus vulgaris</i> L. and their patterns of expression under abiotic stress. <i>Journal of Plant Interactions</i> , 2022, 17, 390-403.	2.1	8
7	Emerging Roles of SWEET Sugar Transporters in Plant Development and Abiotic Stress Responses. <i>Cells</i> , 2022, 11, 1303.	4.1	27
8	Genome-wide identification and expression analysis of the SWEET gene family in daylily (<i>Hemerocallis</i>) Tj ETQq1 1 0.784314 rgBT /Over 211.	3.6	11
9	Genome-wide identification of the <i>SWEET</i> gene family mediating the cold stress response in <i>Prunus mume</i> . <i>PeerJ</i> , 2022, 10, e13273.	2.0	4
10	Characterization and Functional Analysis of <i>ZmSWEET15a</i> in Maize. <i>DNA and Cell Biology</i> , 2022, 41, 564-574.	1.9	3
11	An overview of sucrose transporter (SUT) genes family in rice. <i>Molecular Biology Reports</i> , 2022, 49, 5685-5695.	2.3	12
12	Validation of a QTL on Chromosome 1DS Showing a Major Effect on Salt Tolerance in Winter Wheat. <i>International Journal of Molecular Sciences</i> , 2022, 23, 13745.	4.1	0
13	Genome-Wide Identification of the RsSWEET Gene Family and Functional Analysis of RsSWEET17 in Root Growth and Development in Radish. <i>Horticulturae</i> , 2023, 9, 698.	2.8	1
14	Genome-Wide Identification and Expression Analysis of the SWEET Gene Family in Annual Alfalfa (<i>Medicago polymorpha</i>). <i>Plants</i> , 2023, 12, 1948.	3.5	1
15	Identification and expression analysis of the SWEET genes in radish reveal their potential functions in reproductive organ development. <i>Molecular Biology Reports</i> , 0, , .	2.3	0
16	The SWEET gene family in watermelon: genome-wide identification, phylogeny, duplication and expression analyses in male sterile buds and under sucrose, fructose and glucose treatments. , 0, , .		0