## **Stefano Gattolin**

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A mutation in amino acid permease AAP6 reduces the amino acid content of the Arabidopsis sieve elements but leaves aphid herbivores unaffected. Journal of Experimental Botany, 2010, 61, 55-64.	2.4	126
2	Mapping of Tonoplast Intrinsic Proteins in Maturing and Germinating Arabidopsis Seeds Reveals Dual Localization of Embryonic TIPs to the Tonoplast and Plasma Membrane. Molecular Plant, 2011, 4, 180-189.	3.9	102
3	In vivo imaging of the tonoplast intrinsic protein family in Arabidopsis roots. BMC Plant Biology, 2009, 9, 133.	1.6	81
4	Spatial and temporal expression of the response regulators ARR22 and ARR24 in Arabidopsis thaliana. Journal of Experimental Botany, 2006, 57, 4225-4233.	2.4	54
5	An <scp>A</scp> rabidopsis reticulon and the atlastin homologue <scp><i>RHD3â€like2</i></scp> act together in shaping the tubular endoplasmic reticulum. New Phytologist, 2013, 197, 481-489.	3.5	50
6	Deletion of the miR172 target site in a <scp>TOE</scp> â€ŧype gene is a strong candidate variant for dominant doubleâ€flower trait in Rosaceae. Plant Journal, 2018, 96, 358-371.	2.8	43
7	A Diurnal Component to the Variation in Sieve Tube Amino Acid Content in Wheat. Plant Physiology, 2008, 147, 912-921.	2.3	42
8	Tonoplast intrinsic proteins and vacuolar identity. Biochemical Society Transactions, 2010, 38, 769-773.	1.6	29
9	A New Intra-Specific and High-Resolution Genetic Map of Eggplant Based on a RIL Population, and Location of QTLs Related to Plant Anthocyanin Pigmentation and Seed Vigour. Genes, 2020, 11, 745.	1.0	23
10	Secretion, purification and activity of two recombinant pepper endo-β-1,4-glucanases expressed in the yeast Pichia pastoris. FEBS Letters, 1998, 422, 23-26.	1.3	22
11	Integrative genomics approaches validate PpYUC11-like as candidate gene for the stony hard trait in peach (P. persica L. Batsch). BMC Plant Biology, 2018, 18, 88.	1.6	21
12	Mutations in orthologous PETALOSA TOE-type genes cause a dominant double-flower phenotype in phylogenetically distant eudicots. Journal of Experimental Botany, 2020, 71, 2585-2595.	2.4	20
13	Genetic dissection of Sharka disease tolerance in peach (P. persica L. Batsch). BMC Plant Biology, 2017, 17, 192.	1.6	19
14	Analysis of monoâ€; di―and oligosaccharides by CE using a twoâ€stage derivatization method and LIF detection. Electrophoresis, 2009, 30, 1399-1405.	1.3	16
15	A SmelAAT Acyltransferase Variant Causes a Major Difference in Eggplant (Solanum melongena L.) Peel Anthocyanin Composition. International Journal of Molecular Sciences, 2021, 22, 9174.	1.8	16
16	PeachVar-DB: A Curated Collection of Genetic Variations for the Interactive Analysis of Peach Genome Data. Plant and Cell Physiology, 2018, 59, e2-e2.	1.5	12
17	Milk protein polymorphism in Amiata donkey. Livestock Science, 2019, 230, 103845.	0.6	8
18	The effect of selection on casein genetic polymorphisms and haplotypes in Italian Holstein cattle. Italian Journal of Animal Science, 2020, 19, 833-839.	0.8	8

#	Article	IF	CITATIONS
19	The <i>Di2/pet</i> Variant in the <i>PETALOSA</i> Gene Underlies a Major Heat Requirement-Related QTL for Blooming Date in Peach [ <i>Prunus persica</i> (L.) Batsch]. Plant and Cell Physiology, 2021, 62, 356-365.	1.5	7
20	Scrapped but not neglected: Insights into the composition, molecular modulation and antioxidant capacity of phenols in peel of eggplant (Solanum melongena L.) fruits at different developmental stages. Plant Physiology and Biochemistry, 2021, 167, 678-690.	2.8	6
21	Detection of natural and induced mutations from next generation sequencing data in sweet orange bud sports. Acta Horticulturae, 2019, , 119-124.	0.1	2
22	Less is more: natural variation disrupting a miR172 gene at the di locus underlies the recessive double-flower trait in peach (P. persica L. Batsch). BMC Plant Biology, 2022, 22, .	1.6	2
23	The role of ARR22 and two-component systems during Arabidopsis pod development. Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology, 2007, 146, S270-S271.	0.8	0