## Tamar Krugman

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genomic Architecture of Phenotypic Plasticity in Response to Water Stress in Tetraploid Wheat. International Journal of Molecular Sciences, 2021, 22, 1723.	4.1	13
2	Endopolyploidy Variation in Wild Barley Seeds across Environmental Gradients in Israel. Genes, 2021, 12, 711.	2.4	4
3	TdPm60 identified in wild emmer wheat is an ortholog of Pm60 and constitutes a strong candidate for PmG16 powdery mildew resistance. Theoretical and Applied Genetics, 2021, 134, 2777-2793.	3.6	12
4	Grain protein content and thousand kernel weight QTLs identified in a durum × wild emmer wheat mapping population tested in five environments. Theoretical and Applied Genetics, 2020, 133, 119-131.	3.6	47
5	Chromatin dynamics during interphase and cell division: similarities and differences between model and crop plants. Journal of Experimental Botany, 2020, 71, 5205-5222.	4.8	32
6	Variation in phosphorus and sulfur content shapes the genetic architecture and phenotypic associations within the wheat grain ionome. Plant Journal, 2020, 101, 555-572.	5.7	14
7	GenoTypeMapper: graphical genotyping on genetic and sequence-based maps. Plant Methods, 2020, 16, 123.	4.3	3
8	Glycerol-Induced Powdery Mildew Resistance in Wheat by Regulating Plant Fatty Acid Metabolism, Plant Hormones Cross-Talk, and Pathogenesis-Related Genes. International Journal of Molecular Sciences, 2020, 21, 673.	4.1	28
9	Exogenous sodium diethyldithiocarbamate, a Jasmonic acid biosynthesis inhibitor, induced resistance to powdery mildew in wheat. Plant Direct, 2020, 4, e00212.	1.9	11
10	Epigenetics: possible applications in climate-smart crop breeding. Journal of Experimental Botany, 2020, 71, 5223-5236.	4.8	84
11	Durum Wheat as a Bridge Between Wild Emmer Wheat Genetic Resources and Bread Wheat. , 2019, , 201-230.		20
12	The Institute of Evolution Wild Cereal Gene Bank at the University of Haifa. Israel Journal of Plant Sciences, 2018, 65, 129-146.	0.5	14
13	Compassionate approaches for the conservation and protection of fire salamanders. Israel Journal of Ecology and Evolution, 2017, 63, 43-51.	0.6	8
14	Ancestral QTL Alleles from Wild Emmer Wheat Enhance Root Development under Drought in Modern Wheat. Frontiers in Plant Science, 2017, 8, 703.	3.6	42
15	Ancestral QTL Alleles from Wild Emmer Wheat Improve Drought Resistance and Productivity in Modern Wheat Cultivars. Frontiers in Plant Science, 2016, 7, 452.	3.6	82
16	Ancestral QTL alleles from wild emmer wheat improve grain yield, biomass and photosynthesis across enviroinments in modern wheat. Plant Science, 2016, 251, 23-34.	3.6	37
17	Evolution and Adaptation of Wild Emmer Wheat Populations to Biotic and Abiotic Stresses. Annual Review of Phytopathology, 2016, 54, 279-301.	7.8	67
18	Distribution and haplotype diversity of WKS resistance genes in wild emmer wheat natural populations. Theoretical and Applied Genetics, 2016, 129, 921-934.	3.6	24

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19	Ecological transcriptomics – a nonâ€lethal sampling approach for endangered fire salamanders. Methods in Ecology and Evolution, 2015, 6, 1417-1425.	5.2	16
20	Transcriptome profiling of wheat glumes in wild emmer, hulled landraces and modern cultivars. BMC Genomics, 2015, 16, 777.	2.8	16
21	Alteration in expression of hormone-related genes in wild emmer wheat roots associated with drought adaptation mechanisms. Functional and Integrative Genomics, 2011, 11, 565-583.	3.5	74
22	Multilevel regulation and signalling processes associated with adaptation to terminal drought in wild emmer wheat. Functional and Integrative Genomics, 2010, 10, 167-186.	3.5	67
23	Chromosomal regions controlling seedling drought resistance in Israeli wild barley, Hordeum spontaneum C. Koch. Genetic Resources and Crop Evolution, 2010, 57, 85-99.	1.6	54
24	Genomic dissection of drought resistance in durum wheatâ€f×â€fwild emmer wheat recombinant inbreed line population. Plant, Cell and Environment, 2009, 32, 758-779.	5.7	202
25	Allelic diversity associated with aridity gradient in wild emmer wheat populations. Plant, Cell and Environment, 2008, 31, 39-49.	5.7	80
26	Identification of a novel gene (Hsdr4) involved in water-stress tolerance in wild barley. Plant Molecular Biology, 2007, 64, 17-34.	3.9	80
27	Ecogeographic and genetic determinants of kernel weight and colour of wild barley (Hordeum) Tj ETQq1 1 0.784	314 <sub>.</sub> rgBT 1.7	/Oygrlock 10
28	Wild barley eibi1 mutation identifies a gene essential for leaf water conservation. Planta, 2004, 219, 684-93.	3.2	40
29	Title is missing!. Conservation Genetics, 2000, 1, 191-207.	1.5	24
30	Edaphic natural selection of allozyme polymorphisms in Aegilops peregrina at a Galilee microsite in Israel. Heredity, 1994, 72, 109-112.	2.6	61