## Luca Venturini

List of Publications by Year in descending order

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LUCA VENTURINI

#	Article	IF	CITATIONS
1	Multiple wheat genomes reveal global variation in modern breeding. Nature, 2020, 588, 277-283.	13.7	513
2	Multi-parent populations in crops: a toolbox integrating genomics and genetic mapping with breeding. Heredity, 2020, 125, 396-416.	1.2	124
3	Cenetic analysis of wheat sensitivity to the ToxB fungal effector from Pyrenophora tritici-repentis, the causal agent of tan spot. Theoretical and Applied Genetics, 2020, 133, 935-950.	1.8	31
4	A chromosome-anchored eggplant genome sequence reveals key events in Solanaceae evolution. Scientific Reports, 2019, 9, 11769.	1.6	179
5	Efficient and accurate detection of splice junctions from RNA-seq with Portcullis. GigaScience, 2018, 7, .	3.3	98
6	Leveraging multiple transcriptome assembly methods for improved gene structure annotation. GigaScience, 2018, 7, .	3.3	122
7	The transcriptional landscape of polyploid wheat. Science, 2018, 361, .	6.0	768
8	Shifting the limits in wheat research and breeding using a fully annotated reference genome. Science, 2018, 361, .	6.0	2,424
9	An improved assembly and annotation of the allohexaploid wheat genome identifies complete families of agronomic genes and provides genomic evidence for chromosomal translocations. Genome Research, 2017, 27, 885-896.	2.4	464
10	Decreased Nucleotide and Expression Diversity and Modified Coexpression Patterns Characterize Domestication in the Common Bean. Plant Cell, 2014, 26, 1901-1912.	3.1	103
11	De novo transcriptome characterization of Vitis vinifera cv. Corvina unveils varietal diversity. BMC Genomics, 2013, 14, 41.	1.2	110
12	TP53 Alterations Including Missense Mutations, Aberrant Exon-Junctions and Internal Intron Retentions Are Frequent and May Contribute to MDM2 Antagonist-Resistance in B-Acute Lymphoblastic	0.6	0

leukemia. Blood, 2011, 118, 1484-1484.