## 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.<br>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.<br>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.<br>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<br>Retentions Are Frequent and May Contribute to MDM2 Antagonist-Resistance in B-Acute Lymphoblastic                         | 0.6  | 0         |

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