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Identification Of Triticum Aestivum L

Wheat (*Triticum aestivum* L., $2n = 6x = 42$, BBAADD) is one of the most important crops in the world. Extensive research has been conducted on the AP2/ERF superfamily transcription factors in wheat, but attention has

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mainly focused on the ERF family because of their significant roles in abiotic and biotic stresses (Xu et al., 2011 ; Kulkarni et al., 2017).

Genome-Wide Identification and Analysis of the AP2 ...

Identification of candidate genes, regions and markers for pre-harvest

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sprouting resistance in wheat (*Triticum aestivum* L.) Adrian L Cabral , Mark C Jordan , Curt A McCartney , Frank M You , D Gavin Humphreys , Ron MacLachlan , and Curtis J Pozniak

Identification of candidate genes, regions and markers for ...

Although the WD40 family has been

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studied in many plant species, it has yet to be investigated in wheat (*Triticum aestivum* L.), which is an important cereal crop worldwide. Wheat is rich in proteins, carbohydrates, and minerals.

Genome-wide identification and analysis of WD40 proteins ...

The small YABBY plant-specific

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transcription factor has a prominent role in regulating plant growth and developmental activities. However, little information is available about YABBY gene family in *Triticum aestivum* L. Herein, we identified 21 TaYABBY genes in the Wheat genome database. Then, we performed the conserved motif and domain analysis of TaYABBY proteins.

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The phylogeny of the TaYABBY ...

Agronomy | Free Full-Text | Genome Wide Identification ...

Identification of Circular RNAs and Their
Targets in Leaves of Triticum aestivum
L. under Dehydration Stress
Introduction. Wheat (Triticum aestivum)
is the most widely grown and important

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food staple grain. During the life span...
Materials and Methods. Seeds of wheat
cultivars (Triticum aestivum ...

Frontiers | Identification of Circular RNAs and Their ...

Common wheat (Triticum aestivum) is
one of the most important crop species
and together with maize and rice

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provides > 60% of the calories and proteins for humans. Comprehensive identification of miRNAs in the common bread wheat is therefore of great importance.

Identification and characterization of a subset of ...

Identification of candidate genes,

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regions and markers for pre-harvest
sprouting resistance in wheat (*Triticum
aestivum* L.) Adrian L Cabral 1 , 3 Mark C
Jordan 1

Identification of candidate genes, regions and markers for ...

Identification and Validation of Markers
Linked to Broad-Spectrum Stem Rust

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Resistance Gene Sr2 in Wheat (Triticum aestivum L.)

Identification and Validation of Markers Linked to Broad ...

To better understand the molecular mechanisms underlying differential responses to dehydration, transcriptome changes of two contrasting wheat

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(Triticum aestivum L.) cultivars were evaluated in plants grown under unfavorable osmotic conditions. A total of 107 non-redundant transcripts were identified.

link.springer.com

Analysis of metabolic pathways related to fertility restoration and identification

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of fertility candidate genes associated with *Aegilops kotschyi* cytoplasm in wheat (*Triticum aestivum* L.)

Yulin Jia's research works | Northwest A & F University ...

Identification of genes involved in male sterility in wheat (*Triticum aestivum* L.) which could be used in a genic hybrid

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breeding system Matthew J. Milner. The John Bingham Laboratory, NIAB, Cambridge, UK. Search for more papers by this author. Melanie Craze.

Identification of genes involved in male sterility in ...

Wheat (*Triticum aestivum* L.), which is one of the most important staple food

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crops worldwide, is adversely affected by periodic drought in more than half of its area of production (Rajaram 2001). The productivity of wheat is often limited by a shortage of water (Aprile et al. 2009).

Identification of osmotic stress-responsive genes from ...

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Results: A total of 18 COI candidate sequences for 8 members of COI gene family were isolated in wheat (*Triticum aestivum* L.). Phylogenetic and structural analyses showed that these COI genes could be divided into seven distinct subfamilies. The COI genes showed high expression in stamens and glumes.

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Genome-wide identification and analysis of the COI gene ...

The generic name *Triticum* derives from the Latin for threshing or bruising, and the specific epithet *aestivum* is from the Latin for summer. *Triticum aestivum* is a cultigen (a plant that has been altered by humans through a process of selective breeding) and as such is only

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Triticum aestivum L. | Plants of the World Online

Amino acid transporters (AATs), which transport amino acids across cell membranes, play important roles in alleviating plant damage under stresses and...

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Genome-wide survey of the amino acid transporter gene ...

Triticum aestivum subsp. aestivum - MHNT Common wheat (Triticum aestivum), also known as bread wheat, is a cultivated wheat species. About 95% of wheat produced worldwide is common wheat; it is the most widely grown of all

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crops and the cereal with the highest
monetary yield.

Common wheat - Wikipedia

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do Caribe, acessíveis de forma universal na Internet de modo compatível com as bases internacionais.

Pesquisa | Portal Regional da BVS

Identification of wheat (*Triticum aestivum* L.) genotypes for food safety on two different cadmium contaminated soils Article in Environmental Science

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and Pollution Research 27 (1-2) ·
December 2019...

Identification of wheat (Triticum aestivum L.) genotypes ...

Identification of chromosomal regions
governing grain size and shape in wheat
(Triticum aestivum L) Journal of Genetics
& Breeding März 2004. Andere Autoren.

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Inheritance and identification of DNA markers associated with yellow berry tolerance in wheat (*Triticum aestivum* L) *Euphytica* Mai 2002.

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