

EGDB: A Comprehensive Multi-Omics Database for Energy Grasses and the Epigenomic Atlas of Pearl Millet

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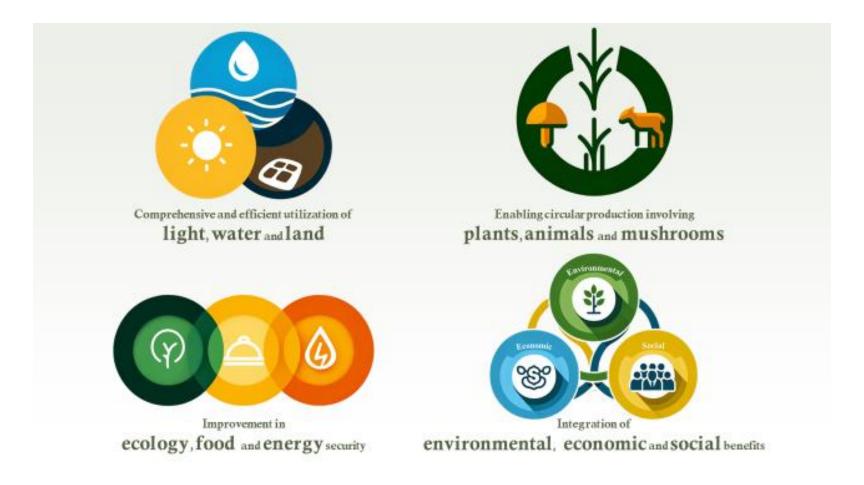
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Lin Luo, Dongmei Lin, Jinhui Li, Hao Chen, Qi Qu, Lin Zhang, Yuan Luo, et al. 2024. EGDB: A comprehensive multi-omics database for energy grasses and the epigenomic atlas of pearl millet. *iMeta* 3: e263. https://doi.org/10.1002/imt2.263



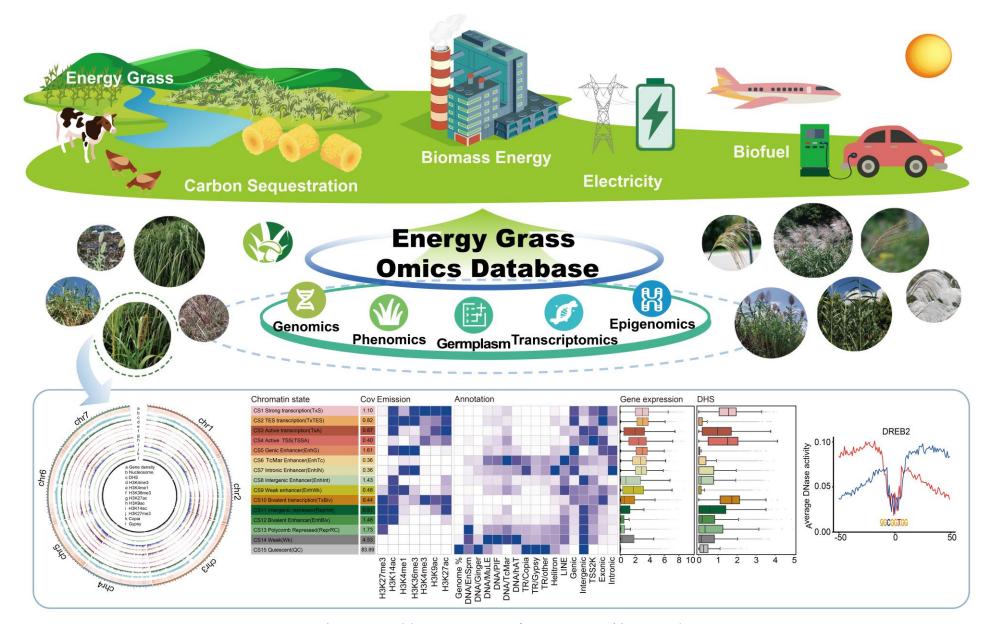
Introduction



- Energy grasses support carbon sequestration, soil health, and rural revitalization.
- EGDB (https://engrass.juncaodb.cn) provides multi-omics data for 11 energy grass species.
- It enables genome annotation and sustainable crop research.



Highlights



https://engrass.juncaodb.cn/



Overview of EGDB

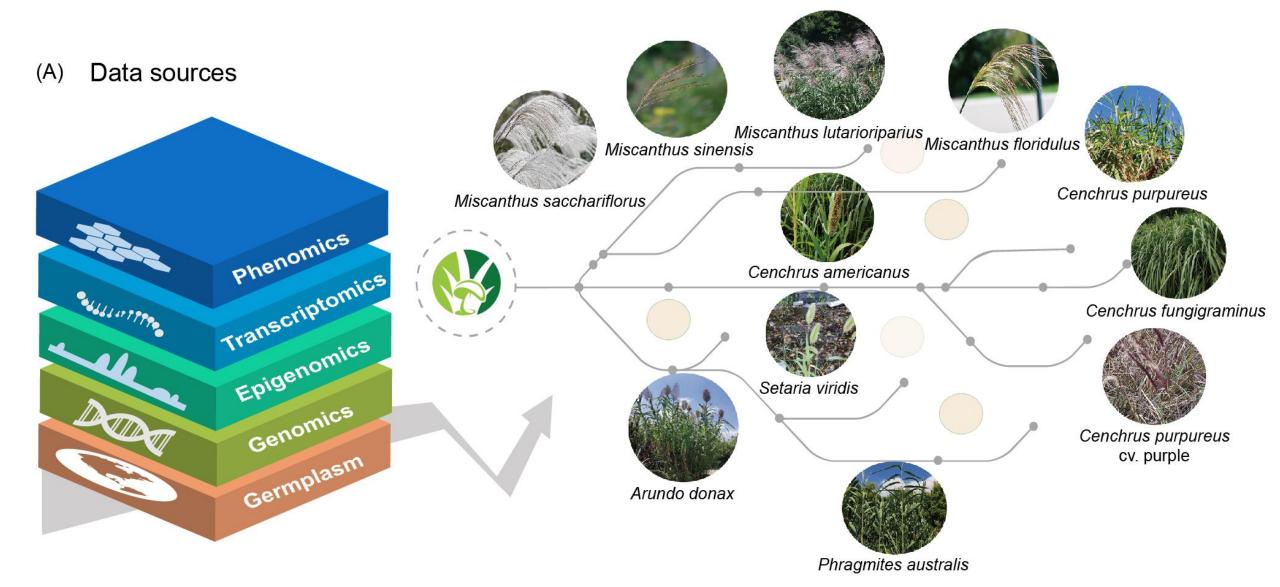


Figure 1. Energy grass database construction and overview and key functionalities. (A) Data source integration



Overview of EGDB



Web Interface

-AJAX -HTML

-CSS -JavaScript

Backend Service

-MySQL -Python -Perl -Nginx

-PHP -R -JavaScript

Front-end Interface

-HTML -Bootstrap -AJAX -CSS -iQuery -JavaScript

Dependency

-JBrowse2 -Sequence Server

-OrthoVenn3 -Primer3web

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Species Species

Browse Germplasm Phenotype

₩ Tools

JBrowse2 / BLAST+ Primer3 / OrthoVenn3 TE identification CRISPR-design

Q Search

Species search Gene search Gene classification Gene expression TF identification

Downloads

Genome / GFF / CDS Histone modification Chromatin accessibility TE annotation

About

About us Tutorial Guide

Middleware infrastructure

Modules and functions

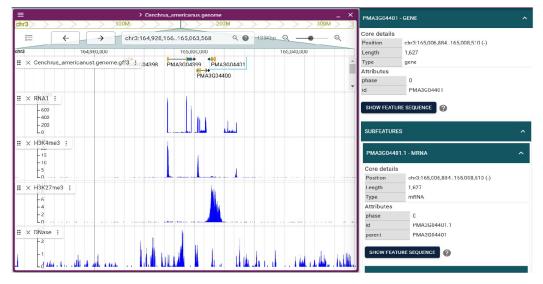


Overview of EGDB

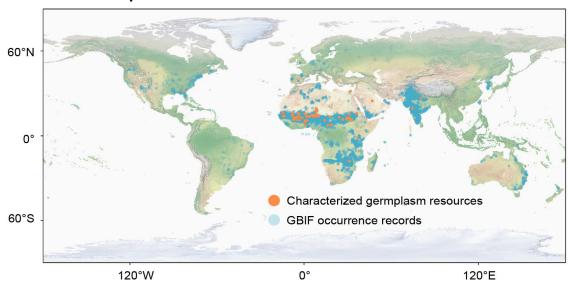
(D) Home page interface



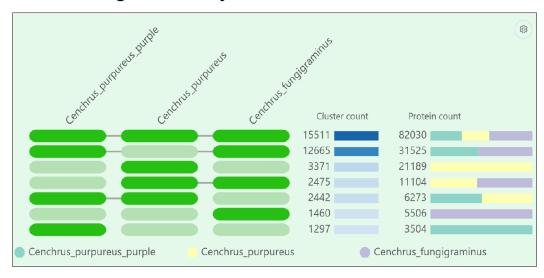
(F) JBrowse2



(E) Germplasm



(G) Orthologous analysis





Epigenetic features of pearl millet

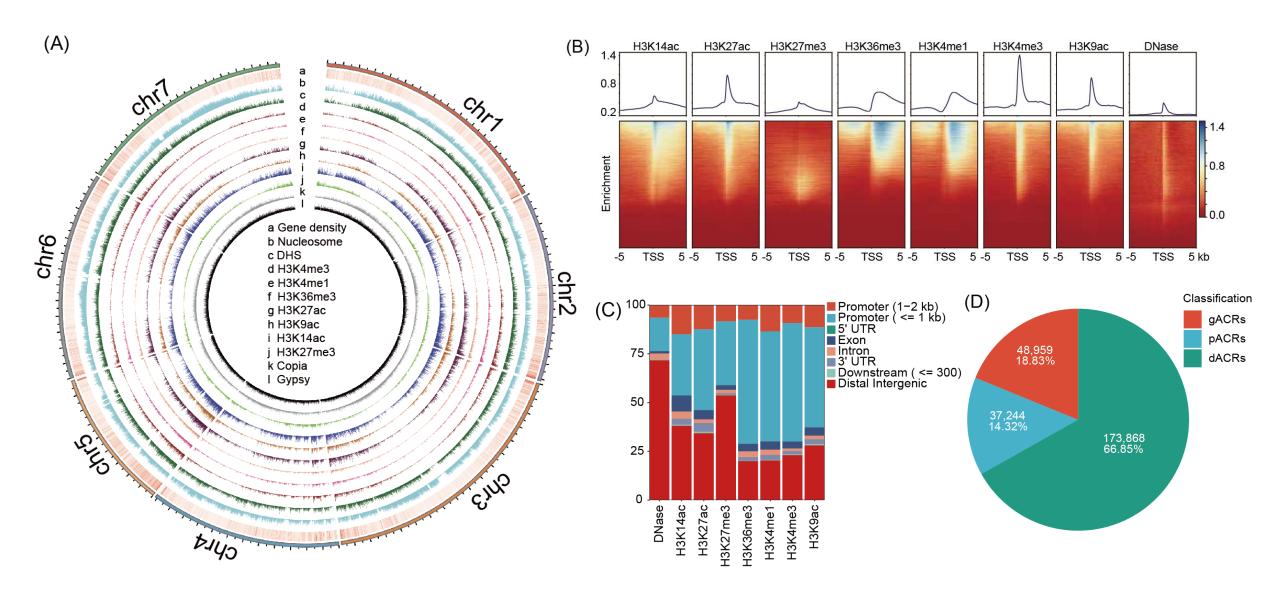


Figure 2. The epigenomic map of pearl millet helped to unveil the regulatory mechanism for gene expression.



Epigenetic features of pearl millet

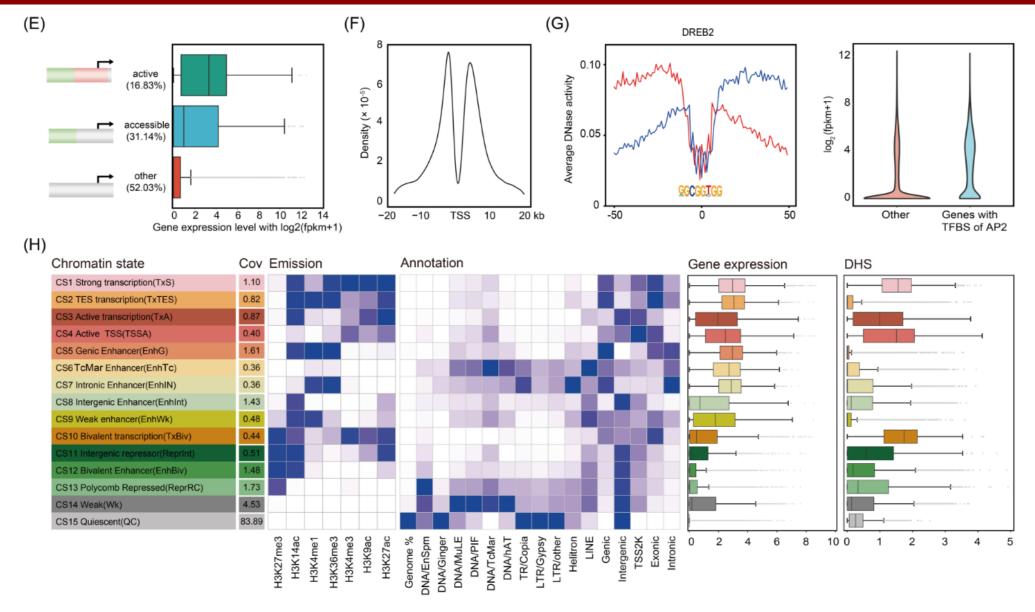


Figure 2. The epigenomic map of pearl millet helped to unveil the regulatory mechanism for gene expression.



Epigenetic features of pearl millet

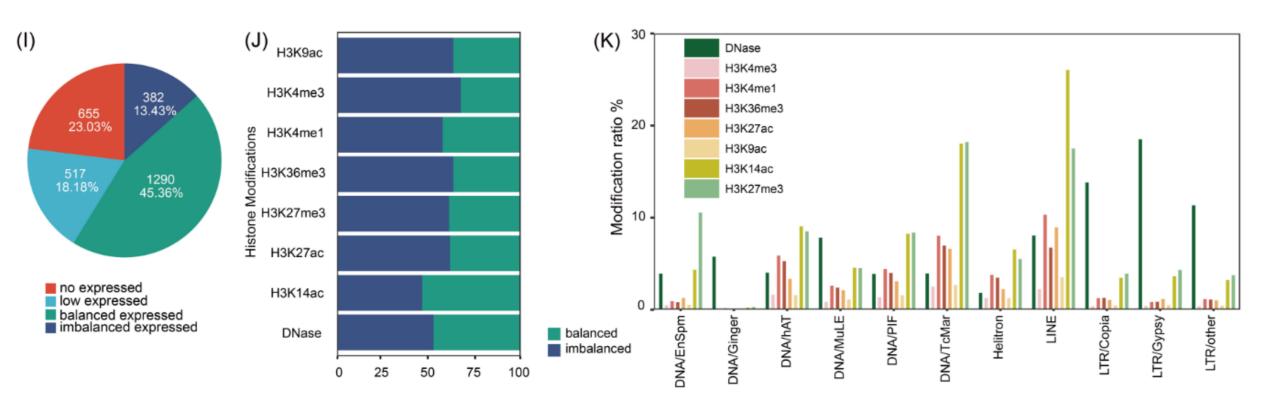


Figure 2. The epigenomic map of pearl millet helped to unveil the regulatory mechanism for gene expression.

- (I) Imbalanced Expression Patterns
- (J) Epigenomic Features and Expression Imbalance
- (K) Histone Modification in TE Families

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Summary

- ☐ In this study, We developed EGDB (https://engrass.juncaodb.cn/), a comprehensive multi-omics database integrating genomic, epigenomic, transcriptomic, and phenotypic data of energy grasses.
- We constructed the epigenomic atlas of pearl millet, covering key histone modifications and chromatin accessibility, providing valuable resources for epigenetic research.
- Our study supports future scientific and translational research, promoting sustainable agriculture and renewable energy development through energy grass studies.

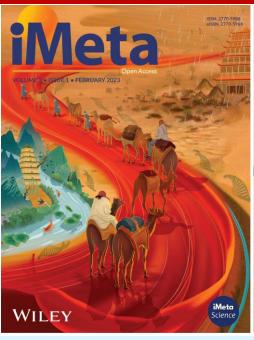
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