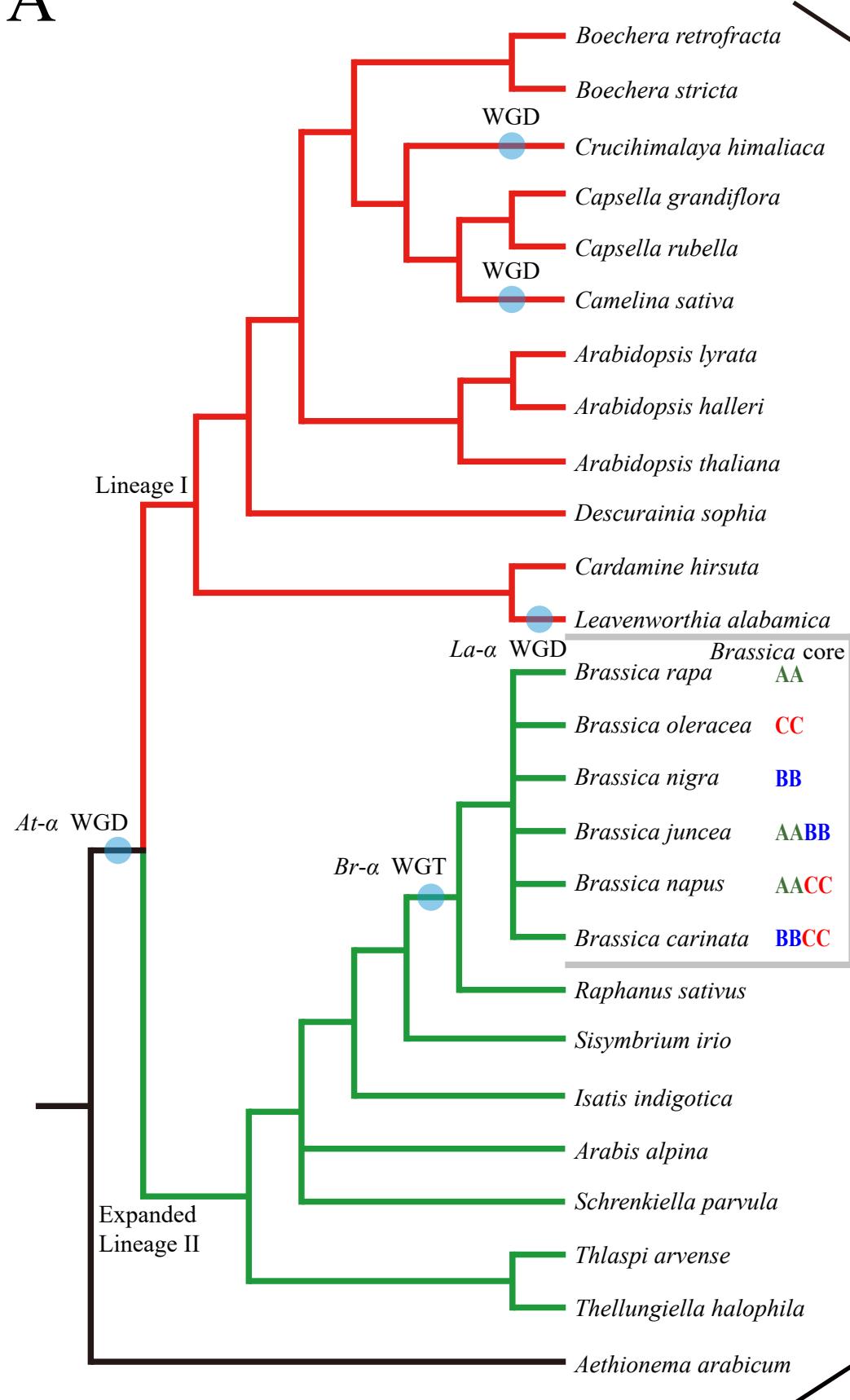


A**Data sources**

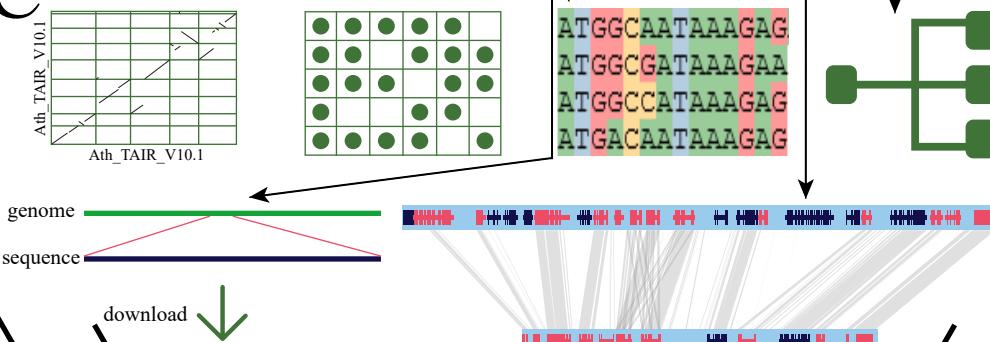
- Genome assembly
- Gene model
- Gene family
- Sequencing data
- Variations and primer
- Gene expression profile

B**Annotations**

- InterPro domain
- Pfam domain
- Gene ontology
- KEGG ontology
- Best hit to *A.thaliana*

Analysis

- Genome Synteny
- Syntenic gene and subgenome
- Multiple sequence alignment
- Genomic sequence retrieval
- Phylogenetic tree
- MicroSynteny

C**D**

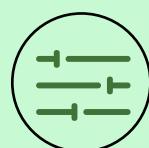
MySQL
Django
Vue
D3.js
Element UI

E**Browse**

- Species info
- Gene family

**Search**

- by Gene id
- by Annotations
- by Coordinate

**Genome browser**

- Variation tracks
- Marker tracks
- RNA-seq tracks

**Tools**

- BLAST
- Primer design
- MicroSynteny
- Genomic sequence retrieval

**Download**

- Genome assembly
- MicroSynteny figure
- Result table