Resources

These pages include a curated set of bioinformatics and data resources that have been compiled by members of the International Arabidopsis Informatics Consortium (IAIC) and curators at TAIR. This is not a comprehensive list; if you know of a resource that you think should be included, please send us an email.

Search within these pages for Arabidopsis community bioinformatics tools and data resources. You can use labels for data types such as 'proteomics' or 'gene_expression' or actions such as data submission or data analysis.

Browse by subject portals

Epigenomics Resources

Education and Outreach Resources

General Databases

Gene Expression/Transcriptomics Resources

Metabolomics Resources

Mutant and Mapping Resources

Proteomics Resources

Stock Centers (DNA, Seeds, other Physical Resources)

Or you can **browse** the list of resources **based on tags**.

A-C	D-E	F-G	H-L	M-N
advocacy	data_analysis	fair	help_documents	mapping
analytics	data_management	functional_annotation	homologs	meeting-notes
arabidopsis	data_repository	gene	homology	metabolomics
bioinformatics	data_sets	gene_annotation	illustrations	metadata
brassicaceae	data_submission	gene_expression	interactome	methylation
chip-seq	data_visualization	gene_function		microarray
cis_elements	dataset	gene_ontology		mirna
clones	dna_methylation	general		mutants
comparative_genomics	dna_stocks	genomics		natural_variation
covid_19	education_outreach	governance		nomenclature
	epigenomics	gwas		

O	P-Q		R	S	T-Z
ontologies	phenotypes		reagents	seed_stocks	term_enrichment
organization	phylogenetics		rna_seq	seeds	tf_binding_sites
	phylogenomics			sequence	transcription_factor_binding_sites
	plant_transformation			single_cell	transcriptomics
	plasmids			small_rna	variants
	polymorphisms			software	
	post_translational_modification			standards	
	promoter			stock_center	
	proteomics			structures	
	protocols			synteny	
	publications				

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