

# Epigenomics Resources

- [1001 EpiGenomes Browser](#) —

<b>Resource Link</b>	<a href="#">1001 Genomes Epigenome Browser</a>
<b>Description</b>	AnnoJ Browser for 1001 genomes dataset from Schmitz Lab. Data also available to download.
<b>Maintained by</b>	Bob Schmitz


- [epigenomics](#)
- [dna\\_methylation](#)
- [natural\\_variation](#)

- [AraENCODE](#) —

<b>Resource Link</b>	<a href="#">AraENCODE</a>
<b>Description</b>	<a href="#">Arabidopsis thaliana Encyclopedia of DNA Elements</a> integrates public datasets for histone modification, chromatin accessibility, DNA methylation, transcriptome, and chromatin interactions from different tissues in wild type or mutants. Data is available as browsable, searchable and downloadable formats.
<b>Maintained by</b>	<a href="#">Guoling Li Lab</a>

- [epigenomics](#)

- [ChromDB](#) —

<b>Resource Link</b>	<a href="#">ChromDB</a>
	
<b>Description</b>	The broad mission of ChromDB is display, annotate, and curate sequences of two broad functional classes of biologically important proteins: chromatin-associated proteins (CAPs) and RNA interference-associated proteins.
<b>Maintained by</b>	Carolyn Napoli, University of Arizona

- [epigenomics](#)

- [Circadian Epigenome Browser](#) —

<b>Resource Link</b>	<a href="#">Circadian Epigenome Browser</a>
<b>Description</b>	AnnoJ Browser for Circadian epigenome dataset from Schmitz Lab. Data also available to download.
<b>Maintained by</b>	Bob Schmitz

- [epigenomics](#)
- [dna\\_methylation](#)

- [Ecker Epigenomics Browser](#) —

<b>Resource Link</b>	<a href="#">Ecker Epigenomics Browser</a>
<b>Description</b>	AnnoJ browser of small RNA, expression and DNA methylation data.
<b>Maintained by</b>	Joe Ecker

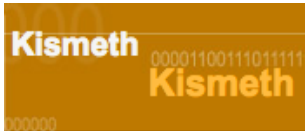
- [small\\_rna](#)
- [epigenomics](#)
- [dna\\_methylation](#)

- [Jacobsen Epigenomics Browser](#) —

<b>Resource Link</b>	<a href="#">Jacobsen Epigenomics Browser</a>
<b>Description</b>	DNA methylation, small RNA, ChIP data from Jacobsen lab publications, in searchable browser/
<b>Maintained by</b>	Steve Jacobsen

- [epigenomics](#)
- [dna\\_methylation](#)
- [small\\_rna](#)
- [gene\\_expression](#)

• [Kismeth](#) —

<b>Resource Link</b>	<a href="#">Kismeth</a>
	
<b>Description</b>	An online tool for analyzing plant DNA methylation patterns.
<b>Maintained by</b>	Ravi Sachidanandam


- [epigenomics](#)
- [data\\_analysis](#)
- [dna\\_methylation](#)

• [Plant Methyloome](#) —

<b>Resource Link</b>	<a href="#">Plant Methyloome DB</a>
<b>Description</b>	AnnoJ Browser for Methyloome data from Arabidopsis and many other plant species
<b>Maintained by</b>	Bob Schmitz

- [epigenomics](#)
- [dna\\_methylation](#)

• [RepBase](#) —

<b>Resource Link</b>	<a href="#">RepBase</a>
	
<b>Description</b>	This is a database of representative repetitive sequences from eukaryotic species. Contains a dataset of <i>Arabidopsis</i> repeats.
<b>Maintained by</b>	Genetic Information Research Institute (GIRI)

- [epigenomics](#)
- [data\\_repository](#)

• [Spontaneous Epiallele \(30 Generation\) Browser](#) —

<b>Resource Link</b>	<a href="#">Spontaneous Epiallele Browser</a>
<b>Description</b>	AnnoJ Browser for 30 generation epiallele dataset from Schmitz Lab. Data also available to download.
<b>Maintained by</b>	Bob Schmitz

- [epigenomics](#)
- [dna\\_methylation](#)