# May 2025 SoyBase Update



#### New tool at SoyBase: Allele Search

The SoyBase team has recently created a new tool to assist the soybean research community. SoyBase has collected a representative sample of sequence variant collections ( VCF files ) from multiple studies including; SoySNP50K from Song et al., 2015 and GmHapMap from Torkamaneh et al., 2019 and many more. When using the Allele Search Tool it is important to ensure the gene ID (Step 2) matches the genome version.

#### Allele Search

Step 1. Select a variant collection. Note that there are three reference assemblies to choose from.

Synopsis: VCF and HapMap files containing genotype information of the whole U.S. soybean germplasm collection using the SoySNP50K BeadChip.

#### Step 2. Enter gene ID and size of left and right flanking regions...

Enter gene name	Enter flanking region	
e.g glyma.Wm82.gnm4.ann1.Glyma.16G040000	e.g 50000	

Downloadable output

#### Allele Search

SEARCH

glyma.Wm82.gnm4.ann1.Glyma.16G040000 **Region searched:** 

glyma.Wm82.gnm4.Gm16: 3637736 - 3839207

CHROMOSOME	VARIANT ID	POSITION	REF	ALT
glyma.Wm82.gnm4.Gm16	ss715624936	3665302	G	А
glyma.Wm82.gnm4.Gm16	ss715624979	3699838	А	G
glyma.Wm82.gnm4.Gm16	ss715625000	3723656	G	Т
glyma.Wm82.gnm4.Gm16	ss715625005	3727020	G	Α
CLOSE		E	XPORT TXT	EXPORT CSV

# **Soybean Community Deadlines**

#### Soybean Carrer Awards

Nomination packets are to be emailed to the SoyGEC chair by June 1st.

### Soy2025 Abstract Submission Deadline

The submission deadline for abstracts to be included in the program book is <u>June</u> **16, 2025**.



# Soy2025 July 23-26, 2025 Madison, WI Registration Open!

# Soybean Breeders Tour September 3-5, 2025 Tifton, GA Registration Open!

## SBW 2026 February 16 - 18, 2026 St. Louis, MO Save the Date!









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