

3rd International Biocuration Conference

April 16-19 Berlin, Germany

The Maize Genetics Cooperation Newsletter MNL

maize gene review

www.maizegenereview.org

A new on-line journal promotes
community curation of phenotypes

Mary Schaeffer



MaizeGDB
Maize Genetics and Genomics Database

MNL and MaizeGDB ties

1933 First formal listings of maize genes and genetic stocks M. Rhoades *MNL* 3:5-16

1991 Maize[G]DB online with data parsed from MNL

- **MGC Stock Center List E. Patterson. *MNL* 65:129-133.**
- **Gene List E. Coe, D. Hoisington, and S. Chao. *MNL* 65:134-148.**

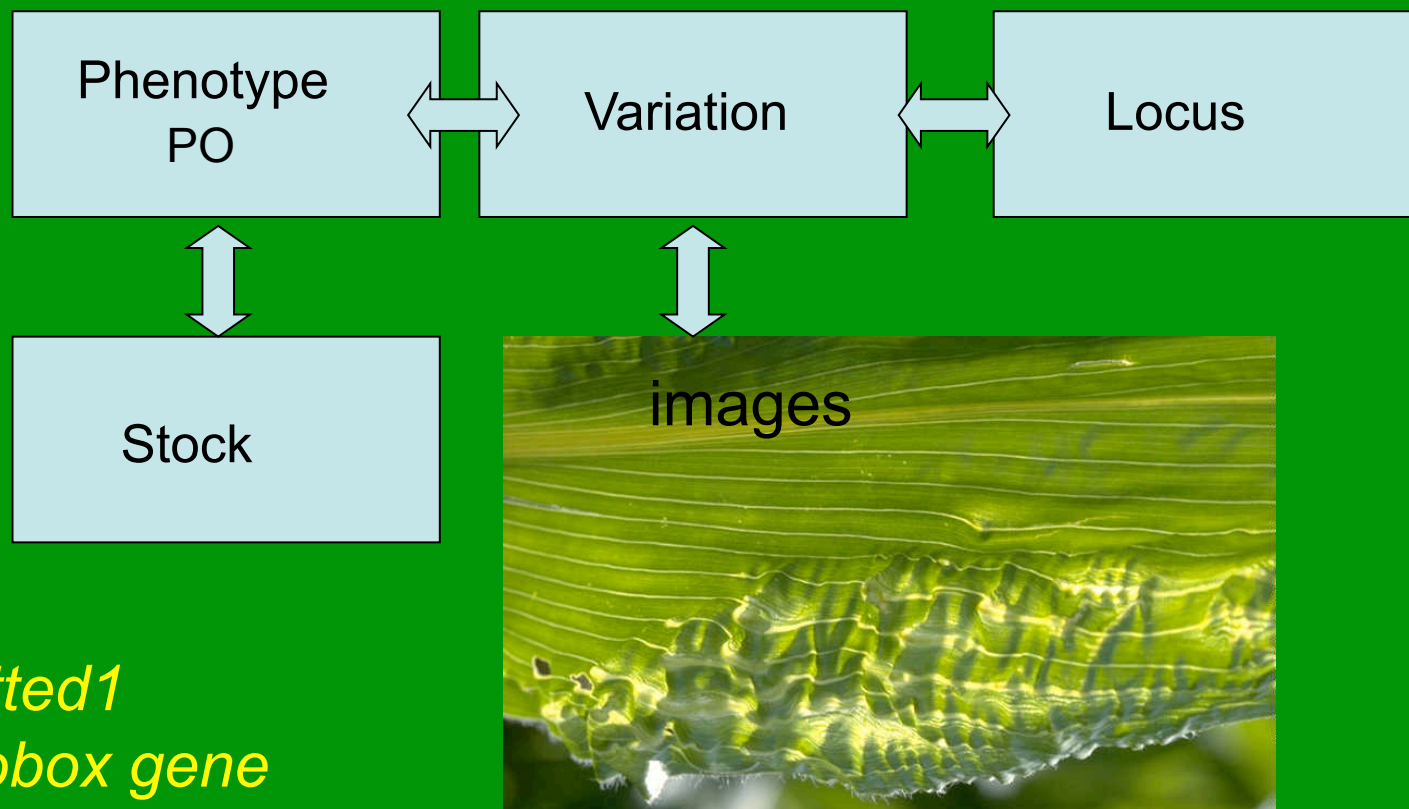
MaizeGDB hosts online MNL

Phenotype data at MaizeGDB

SOURCES:

- The MGCSC Maize Genetics Cooperation Stock Center, Illinois USA, Marty Sachs, Director
- High-throughput projects
- *Mutants of Maize*, eds MG Neuffer, EH Coe, SR Wessler (1997)
- Images - Community, MG Neuffer collection
- Literature

Phenotype data at MaizeGDB



kn1 knotted1
a homeobox gene



knotted leaf ([phenotype](#))

This is also known by the following names:

knotted leaf

Definition:

scattered proliferations of tissue at vascular bundles on leaf

[Add your own annotation to this record!](#)

Body Part(s) Affected: [leaf](#)

Genes:

[cr4](#) [crinkly4](#)

[kn1](#) [knotted1](#)

Variations: (see [all 13 images of these](#))

[cr4-6143](#) (variation of [cr4](#) [crinkly4](#))

[Kn*-N1234D](#)

[Kn*-N64B](#)

[Kn1-167::Mu](#) (variation of [kn1](#) [knotted1](#))

[Kn1-174::Mu](#) (variation of [kn1](#) [knotted1](#))

[Kn1-N2](#) (variation of [kn1](#) [knotted1](#))

[Kn1-Z3](#) (variation of [kn1](#) [knotted1](#))

[Kn*-HsuA1191](#)

[kn*-N1891](#)

[Kn1](#) (variation of [kn1](#) [knotted1](#))

[Kn1-169::Mu](#) (variation of [kn1](#) [knotted1](#))

[Kn1-2F11::Ds2](#) (variation of [kn1](#) [knotted1](#))

[Kn1-O](#) (variation of [kn1](#) [knotted1](#))

[Rs1-Z](#) (variation of [rs1](#) [rough sheath1](#))

Stocks: (stocks listed in **bold** are available from the Stock Center)

[117E](#)

[4011M](#)

▶ This phenotype has **4** references that describe it. Click the arrow to the left to view these references.

[See record in a printer-friendly format](#)

[Return to the homepage](#)

Research Tools

Here are some internet resources to aid your investigation into knotted leaf.

Phenotype Browser

Search Tools

Search [Google](#) for knotted leaf

Search [PubMed](#) for knotted leaf

Search [Usenet](#) for knotted leaf

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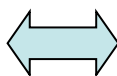
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Body Part(s) Affected: [leaf](#)



PO

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annotation by community at MaizeGDB

Register once with MaizeGDB

Add comments to any page.

Add new records

Soon, annotate genome.

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annotation window

MaizeGDB 
Maize Genetics and Genomics Database

[jobs](#) | [upcoming events](#)
Useful Pages

[home](#) |

Add Annotation For knotted leaf

Here, you can annotate MaizeGDB's information on knotted leaf.

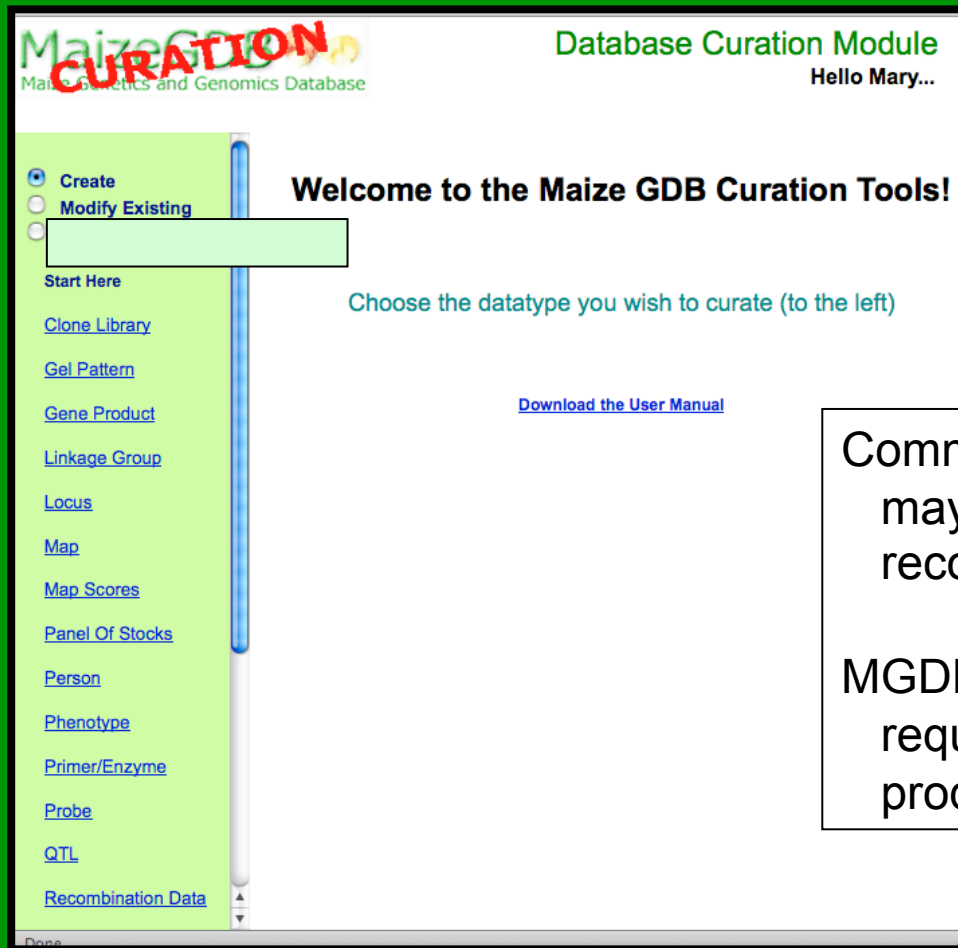
Enter annotation here (limit 1500 characters):

[Return to the homepage](#)

[home](#) |

If you have some comments about this page, or about the site in general, fill out our easy to use [feedback form](#) without leaving this page!

Community curation - detailed tool suite



Community curator
may create and update a new
record prior to production

MGDB Curator approval
required before advancing to
production

Phenotype data at MaizeGDB

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kn1 comment in MaizeGDB

Extracted from Mutants of Maize 1997 ed. Neuffer, Coe, & Wessler

Dominant Kn1, localized proliferation of tissue at vascular bundles on leaf to give a knotted appearance. Homozygotes more extreme. Primary effect is to lateral veins of leaf blades; veins are often more prominent, localized extra cell divisions (knots) occur sporadically along the veins, ligule fringes are found along the veins (Gelinas et al. 1969 Am J Bot 56:671). The phenotype is not determined by the epidermis (Hake and Freeling 1986 Nature 320:621), but by the inner middle mesophyll-bundle sheath layer (Sinha and Hake 1990 Dev Biol Symp 19:77). The dominant mutation is neomorphic (Freeling and Hake 1985 Genetics 111:617). A number of alleles have been cloned (Hake et al. 1989 Embo J 8:15, Veit et al 1990 Genetics 125:623; Gelinas et al. 1969 Am J Bot 56:671). Classification varies widely among alleles and according to genetic background (Hake)

Parenthetic acknowledgement of contributor

maize gene review- in brief

Unlike notes in the Newsletter, these contributions will be:

- tailored for entry into the MaizeGDB,
- peer-reviewed,
- freely available for use by anyone who cites the work, as is the practice for any MaizeGDB data.

authors provide

- * a short summary of the mutant with reference citations
- * an unpublished image with a caption
- * the reference(s) that first reports or defines this locus
- * key alleles, with phenotypes, information about viability or special conditions
- * regulation of/by other genes or genetic elements
- * expression--tissue/conditions/inhibitors/inducers
- * gene product, function, pathways (GO terms welcome)
- * map location, how mapped
- * any other information, e.g. paralogous loci.

MGDB curator/editor provides

- Links to relevant databases: MGDB, NCBI, UniProt
- consensus map coordinates
- known synonyms
- nomenclature checks
- other

Under development

automated generation of gene review
pages with XML markup

DOI registry

Maize Genetics Cooperation Newsletter

maize gene review

[HOME](#)[ABOUT](#)[BROWSE](#)[AUTHORS](#)[CONTACT](#)

Author: [David Braun](#), Department of Biology, 208 Mueller Lab Pennsylvania State University, University Park, PA 16802

DOI/

Pub date: Feb. 2009

Reviewed: Mar 2009

Name: [tdy1 tie-dyed1](#)

Chromosome [6L bin 6.07 BAC* c0256p22/AC204318](#)

*BAC determined by NCBI BLAST

Description: Leaf blades develop variegated chlorotic and green regions; chlorotic regions hyper-accumulate carbohydrates.



Images: Leaf of *tdy1*, courtesy Frank Baker and David Braun.

MGDB: [tdy1](#)
NCBI: [FJ376984](#)
UniProt: [B8XIB0](#)

Key Alleles:

tdy1-Reference (tdy1-R): recessive, EMS-induced, proline to arginine substitution at the 112th amino acid

tdy1-D190: recessive, *Mu1* insertion in the 5'UTR 101 base pairs upstream of the ATG

Summary: *tdy1* is a recessive mutation that results in leaf blades developing chlorotic, nonclonal regions that hyperaccumulate carbohydrates. Expression of the phenotype is dependent on growth in high light. The chlorotic regions appear soon after a leaf emerges from the whorl and stable once formed. *tdy1* is proposed to function in the same genetic pathway as *tdy2* based on their nearly identical mutant phenotypes and a dosage-sensitive genetic interaction between the loci (Baker and Braun, 2008). *tdy1* appears to be a grass-specific gene that encodes a novel, predicted membrane-localized protein (Ma et al., 2009). *tdy1* RNA is expressed in phloem cells in mature and immature leaves, stems and developing ears. *tdy1* RNA accumulates in protophloem sieve elements upon differentiation and is one of the earliest phloem-expressed genes reported in maize. *tdy1* is hypothesized to function as an osmotic stress or sugar sensor that prevents the over accumulation of carbohydrates in developing leaves.

First reported: Braun et al. (2006) *tie-dyed1 regulates carbohydrate accumulation in maize leaves*, Plant Physiology 142: 1511-1522
PUBMED:17071639

tdy1 tie-dyed1
carbohydrate transport mutant

Links to MaizeGDB

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Showing 161.7 kbp from Chr6, positions 159,627,300 to 159,789,000

MaizeGDB Tools - New!

[BLAST] [Bin Viewer] [FTP] [Locus Lookup] [Search]

Instructions

Searching: Search using a sequence name, BAC name or other landmark. The wildcard character * is allowed.

Navigation: Click one of the rulers to center on a location, or click and drag to select a region. Use the Scroll/Zoom buttons to change magnification and position.

Example: Chr6:159,627,300-159,789,000, BAC:AC177838, AC195844, c0151b16, FPCcontig:ctg1, ctg70, smt2, SOG0588, ESTs:149110846, cDNA:21217038, 1.01, MAGIV3.1_32562, MAGiv4.0_76504,

[Hide banner] [Share these tracks] [Link to Image] [High-res Image] [Help] [Reset]

Locus Lookup

Reports & Analysis:

Download GFF File

Configure...

Go

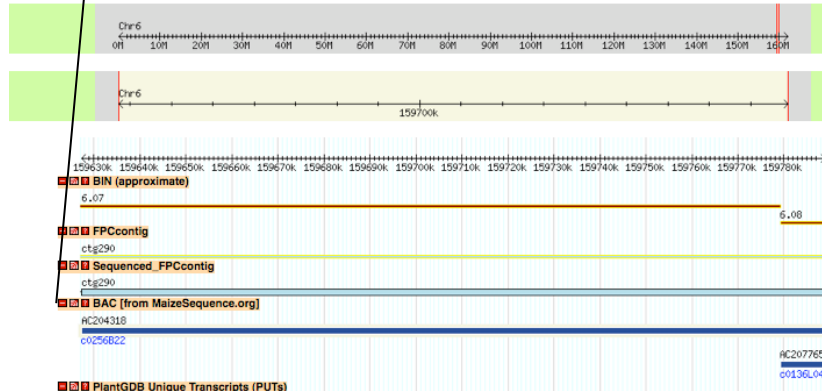
Scroll/Zoom:

Show 161.7 kbp

+

-

Flip



php20599	133.20	6.07
umc1653	133.65	6.07
mmp105	133.65	6.07
agp2	134.10	6.07
asg7a (CBM 6.08)	134.10	6.08
tdy1	134.13	6.07
umc2059	135.68	6.07
ago104	135.75	6.07
hir3	136.13	6.07
umc2324	136.13	6.07
cdo345c	136.45	6.07
cdo202a(mcf)	137.18	6.07
telomere6L	145.00 +/- 8	

Maize Genetics Cooperation Newsletter

maize gene review

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Data transfer to MaizeGDB

Summary description and image added,
citing the MGR

New records, including gene, reference
mutant allele, phenotype, and adding
sequence accessions, consensus map
placement, gene products.

scale

> 800 designated genes in MaizeGDB
with an assigned phenotype

< 250 with some description in
Neuffer et al 1997. Most of these were
not sequenced at the time.

~100 not yet listed at MaizeGDB

46 reviews from 8 authors in process
and evaluation for semi-automation.

BIN	2008 cM	NAME	FULL_NAME	Author
1.01		brk2	brick2	Laurie Smith
1.03	71.8	phyB1	phytochromeB1	Patrick Dubois, Tom Brutnell
1.05	121.5	bif2	barren inflorescence2	Paula McSteen
1.05	202.0	tha4	thylakoid assembly4	Alice Barkan
1.05		wtf1	what's this factor?1	Alice Barkan
1.07	164.8	crs1	chloroplast RNA splicing1	Alice Barkan
1.10	222.5	phyA1	phytochromeA1	Patrick Dubois, Tom Brutnell
1.11	262.8	ts6	tasselseed6	George Chuck
1.12		phyC1	phytochromeC1	Patrick Dubois, Tom Brutnell
2.06	57.3	tha8	thylakoid assembly8	Alice Barkan
3.02	24.4	cg1	corngrass1	George Chuck
3.03	32.8	ra2	ramosa2	Erik Vollbrecht, Sarah Hake
3.04	57.3	caf2	CRS2 associated factor2	Alice Barkan
3.04	57.3	tha1	thylakoid assembly1	Alice Barkan
3.05	78.0	ts4	tasselseed4	George Chuck
3.06		rnc1	ribonuclease1	Alice Barkan
3.08	161.8	spi1	sparse inflorescence1	Paula McSteen
4.07		ppr5	pentricopeptide5	Alice Barkan
5.00		crp2	chloroplast RNA processing2	Alice Barkan
5.01		phyA2	phytochromeA2	Patrick Dubois, Tom Brutnell
5.01		phyC2	phytochromeC2	Patrick Dubois, Tom Brutnell
5.03	50.0	csy1	chloroplast SecY-1	Alice Barkan
5.03	74.4	dvd1	developmental disaster1	Paula McSteen
5.08	167.3	brk1	brick1	Laurie Smith
6.01	30.3	si1	silky1	George Chuck
6.01		why1	whirly1	Alice Barkan
6.02		ppr10	pentricopeptide10	Alice Barkan
6.05	77.8	tan1	tangled1	Laurie Smith
6.05		afd1	absence of first division1	Rachel Wang
6.07	134.1	tdy1	tie-dyed1	David Braun

triage

New genes described by editorial board for MaizeGDB (5 papers a month)

Genes not in Neuffer 1997, that are sequenced, peer-reviewed literature

Updates to Neuffer 1997, but now sequenced

Acknowledgements

University of Missouri, Columbia, MO USA

Joseph Kokenge, Graduate Student, Community Journalism
Jim Birchler, MNL co-editor
Ed Coe ('retired') and Gerry Neuffer ('retired')

MaizeGDB team

Carolyn Lawrence, Director
Lisa Harper, Taner Sen, Carson Andorf, Darwin Campbell

MGC Endowment for the MNL; USDA-ARS

Maize Genetics Cooperation Newsletter - *maize gene review*