



MONTANA
STATE UNIVERSITY

College of
AGRICULTURE &
MONTANA AGRICULTURAL
EXPERIMENT STATION

2025 DURUM FIELD DAY NOTES

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- Dr. Linda Dykes, USDA-ARS, Fargo, ND
- Montana Wheat and Barley Committee Support – Sam Anderson / Lori Wickett/ Kent
- Montana State Foundation Seed Program – Doug Holen / BranDee Johnston

MSU durum breeding is:

- Testing 19 advanced selections against currently grown varieties.
- Working with MSU Foundation Seed to promote our releases.
 - **MT Raska** is high yielding, semi-dwarf, early flowering, very high test weight, low sawfly cutting, low lipoxxygenase for semolina and pasta color retention, resistant to fungal leaf spot
 - **MT Blackbeard** is highest yielding, standard height and flowering time, very high pasta quality, low cadmium, tolerant to fungal leaf spot
 - **WB8148** is a high yielding semi-dwarf, early flowering, high yielding, low sawfly cutting, high pasta quality.
 - **Lustre** is high yielding, standard height, good pasta quality and protein content, low cadmium.
 - **MTD19011**- High yield potential across locations, large seeds, good test weight, good quality traits, low cadmium uptake.
 - **MTD19241** – Solid Stem, good yield potential in saw fly prone environments, good test weight, low cadmium uptake
- **The MSU durum breeding team thanks MT producers and MSU Research Center faculty and personnel for their support.**

MSU durum line Lustre (MTD16005) was released January 2020. MT Raska (MTD18313) and MT Blackbeard (MTD18348) were released January 2022. WB8148 was released in January 2023 and licensed by WestBred.

MT Raska pedigree: Alkabo/Brigade//Alzada/Strongfield

MT Blackbeard pedigree: Alzada/Strongfield//Brigade/Carpio

WB8148 pedigree Joppa/Alzada

MTD19011 pedigree Alkabo/Brigade(F1)//Durafield

MTD19241 pedigree MTD16002/Fortitude

Yield Data

	2021		2022		2023		2024	
Cultivar	Triangle ^{a*}	Eastern MT ^b	Triangle ^a	Eastern MT ^b	Triangle ^a	Eastern MT ^{b*}	Triangle ^a	Eastern MT ^{b*}
MT Blackbeard	<u>28.9</u>	<u>35</u>	31.7	<u>41.2</u>	42.4	48.4	38.5	47.9
MT Raska	26.6	30.6	31.2	39	44.2	<u>52.3</u>	37.8	44.5
WB8148	25.6	<u>35</u>	<u>32.2</u>	40	42.7	49.3	<u>40.9</u>	47.9
ND Riveland	28	33.1	30.2	36.1	39.4	46.3	34.2	45.8
Alzada	/	/	31.4	38.8	<u>47.7</u>	48.2	39.4	47.4
Joppa	24.4	28.5	27.5	38.2	41.7	48.3	35.9	45.8
Mountrail	25.6	29.6	30.8	41	40.4	50.7	38.6	49.0
Divide	25	29.9	29.4	39.5	38.3	49.8	37.0	47.8
Carpio	27.6	27.9	28.3	36.8	38.8	48.1	34.3	47.7
Lustre	23.8	27.1	29.8	36.9	35.7	50.4	38.2	<u>49.2</u>
ANOVA p	0.113	<0.001	0.002	0.04	<0.001	0.26	0.005	0.318
LSD (0.05)		3.71	2.15	3.51	3.87		3.5	
No. Tests	6	2	6	4	6	4	6	5

^a Test locations included EARC dryland, Sheridan Co, Roosevelt Co, Valley Co, McCone Co.

^b Test locations included WTARC, NARC, CARC, Conrad, Ft. Benton. 2023 NARC and 2024 CARC were not included.

IMPROVING WHEAT WITH TARGETED GENE EDITING

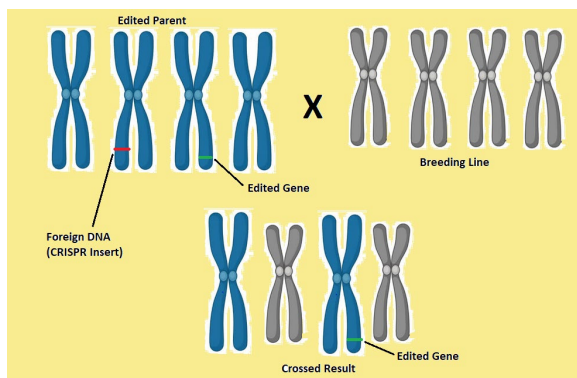
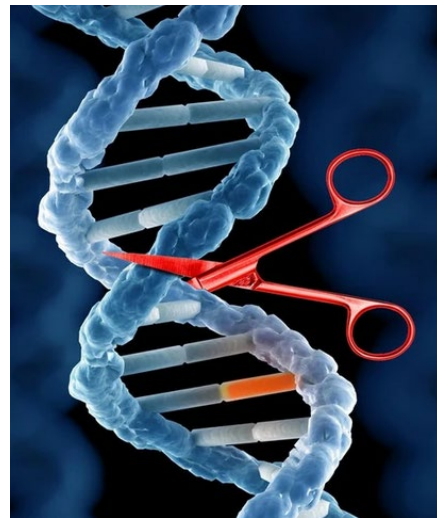
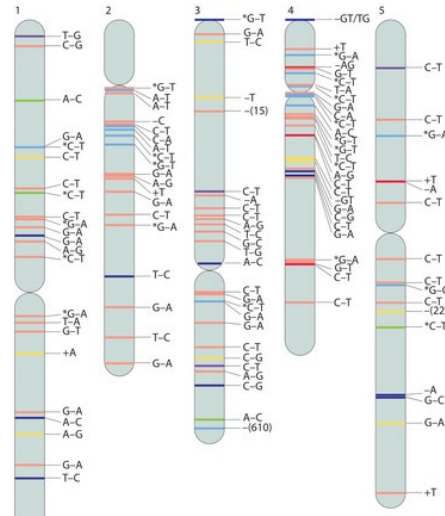
Brandon Tillett

EMS – The Old Way

- Many random mutations
- Time consuming to find the right mutation
- Unavoidable linkage drag

CRISPR – The New Way

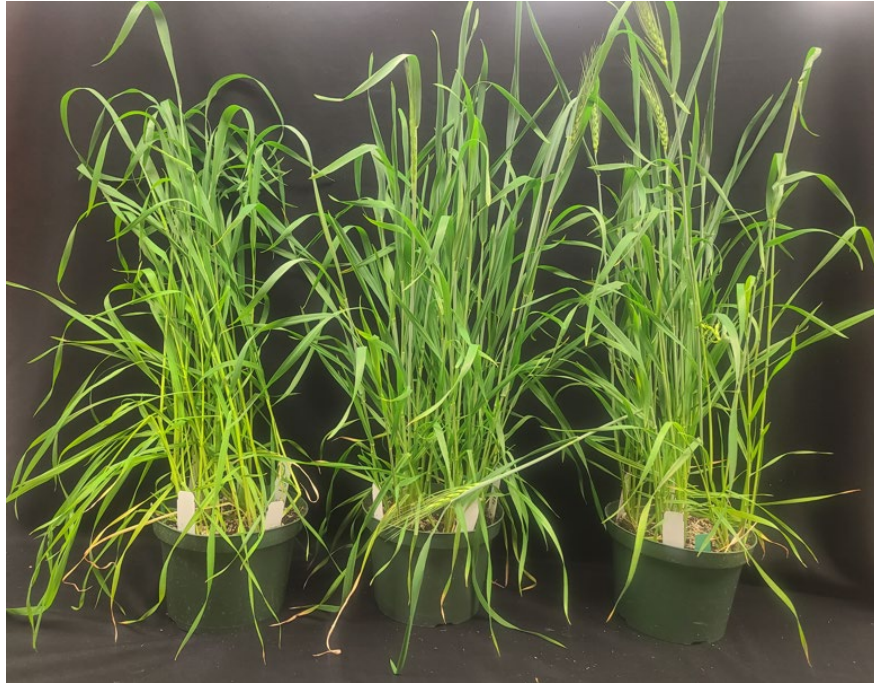
- Precise guided edits
- Can be much faster than EMS
- Significantly less linkage drag than EMS



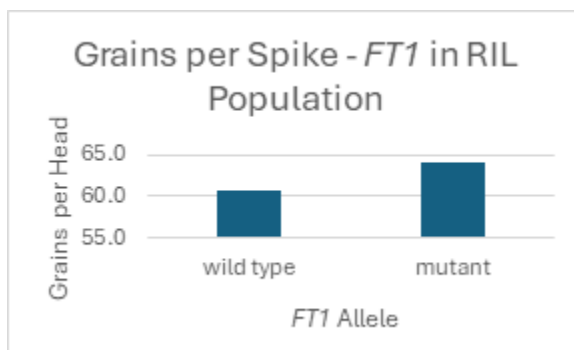
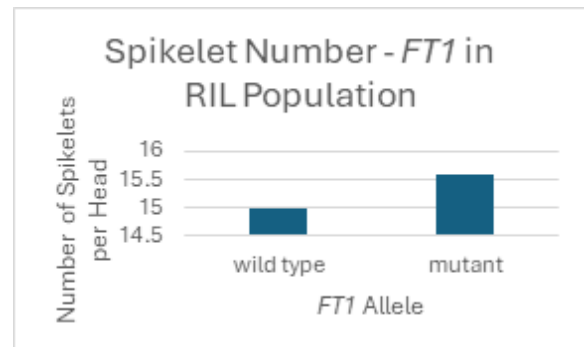
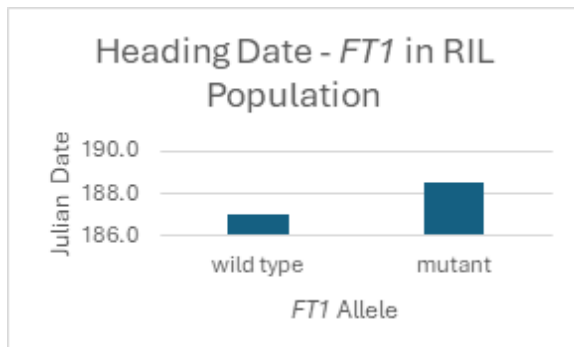
CRISPR insert is separated from edited gene through crossing.

Result = No foreign DNA & a useful new allele

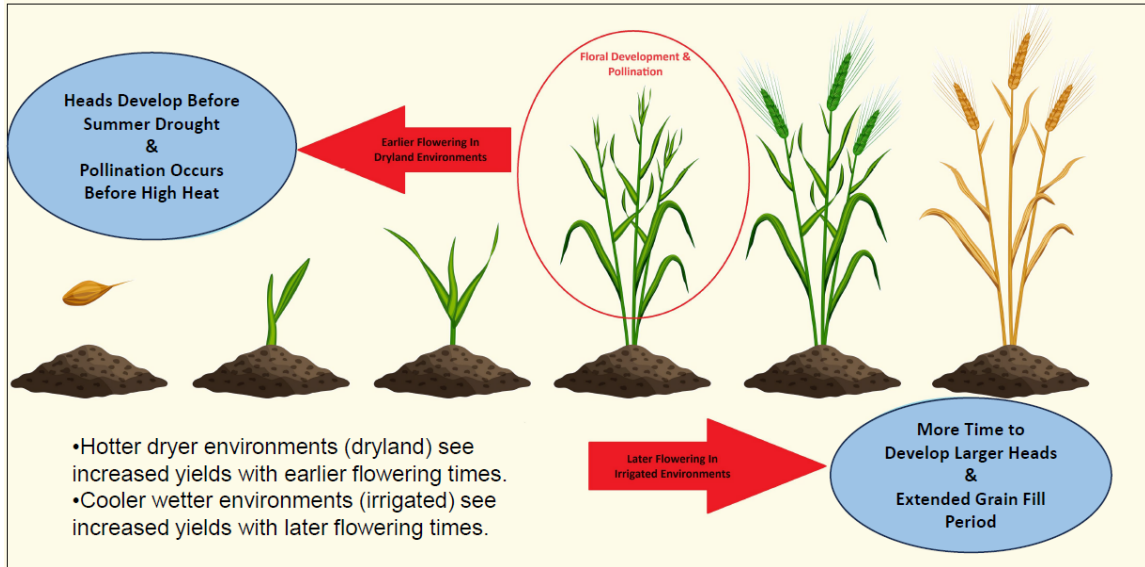
GENE EDITING TARGETS



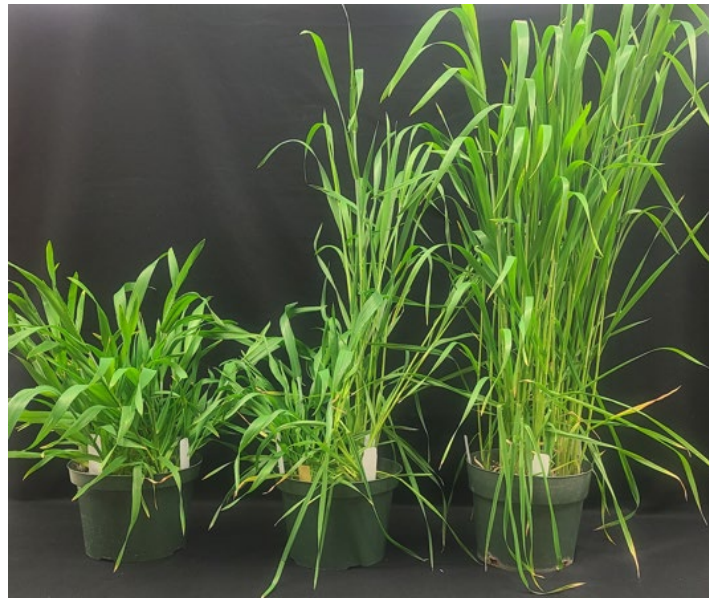
Spring Wheat – MT Carlson varying for flowering time caused by direct edits to flowering genes. The timing of floral development has significant impacts on yield based on environment. Different flowering times will be evaluated for yield stability in the different environmental regions around the state.



The mutant allele of *FT1* delays flowering by 1.5 days, allowing for longer head development which results in on an average of 0.6 more spikelets per spike and 4 grains per spike in a recombinant inbred spring wheat population. In Bozeman 2024 the later flowering allele yielded 7% more than the normal allele.



Other Editing Successes



Tall durum wheat was edited to make semi-dwarf durum wheat. While semi-dwarf wheat already exists, the hope is the gene edited allele that is causing the dwarfing condition will be more agronomically valuable than the current natural allele being utilized in all semi-dwarf durum lines.

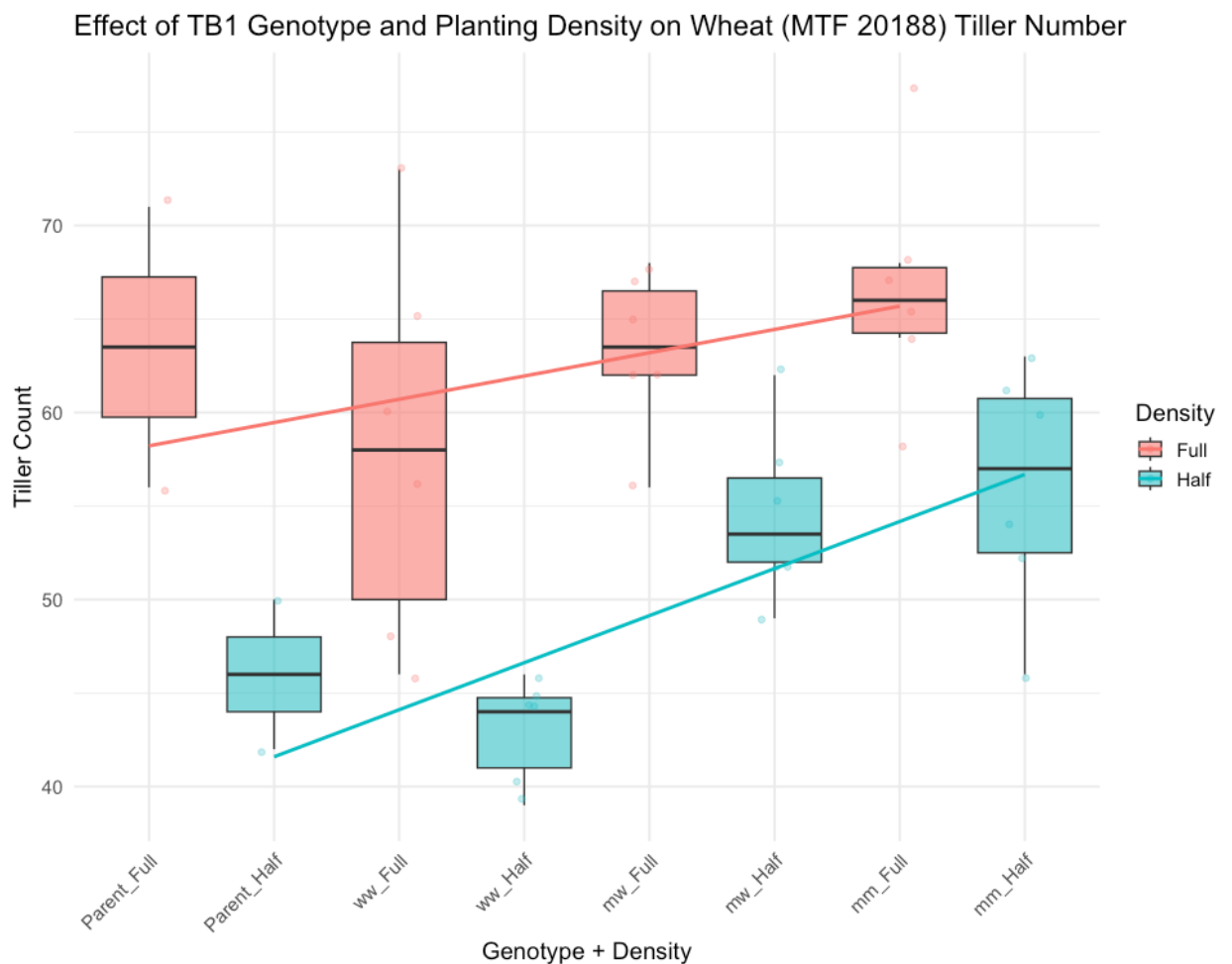
Additional editing projects are underway to pulse crops to increase branching and rooting.

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Teosinte Branched1 mutations on Tillering for Winter Wheat Dual Purpose

Sergei O'Sullivan

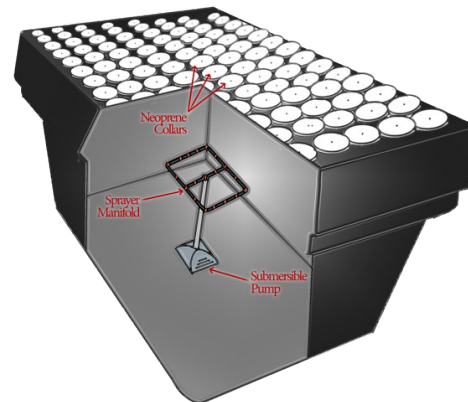
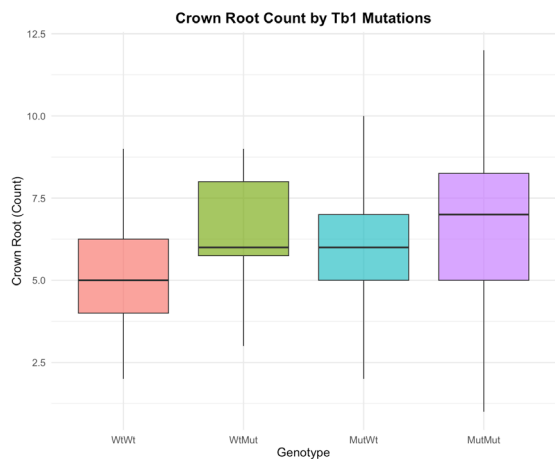
This figure shows how changes to a single gene, Teosinte Branched1 (TB1), affect tiller production in winter wheat grown under two different seeding rates. Each group represents a set of genetically similar lines (composite near-isogenic lines) with specific TB1 mutations, tested in both full and half seeding rate conditions. The boxplots show the number of tillers per plant, while the trend lines highlight a clear pattern: plants with more TB1 mutations tended to produce more tillers, especially at the lower seeding rate. For farmers, this trait could be valuable—extra tillers may help compensate for thin stands caused by winterkill or reduced seeding rates, offering potential for more resilient forage or dual-purpose wheat in challenging conditions.



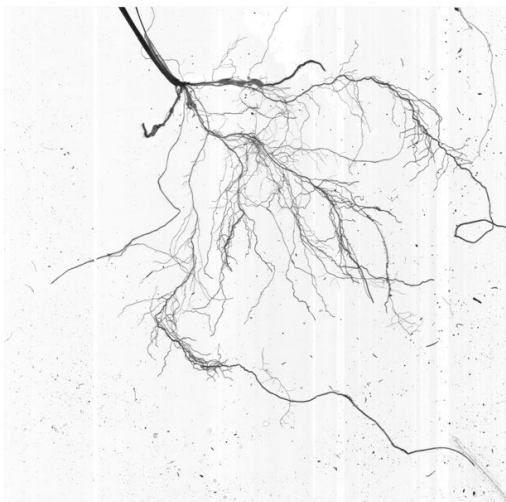
***Teosinte Branched1* for Durum Wheat Rooting**

This figure shows how changes to a single gene, *Teosinte Branched1* (*TB1*), can affect the way wheat roots develop. Plants with more *TB1* mutations tend to produce more crown roots, as shown in both the boxplot (top left) and the root images (bottom). These plants were grown in a soil-free system (top right) that allows us breeders to clearly observe root development over time. The roots of plants with two *TB1* mutations are noticeably denser and more branched compared to those with no mutations. This denser, more branched root system may offer improved access to water and nutrients below ground.

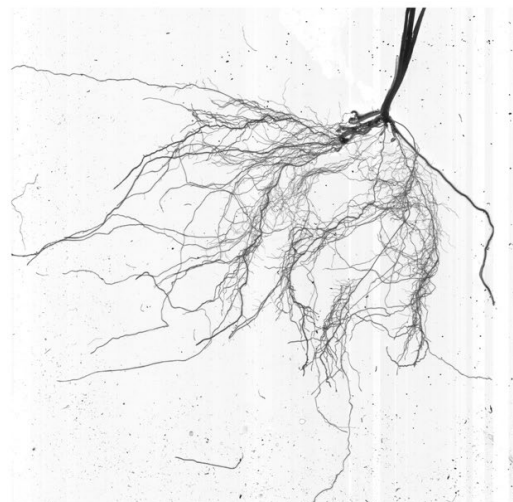
Differences in root architecture like this may influence how plants interact with their environment—particularly in terms of water and nutrient uptake. By understanding the genetic factors that shape root systems, breeders can make more informed decisions when developing wheat varieties suited for diverse and challenging conditions.



No Mutations



2x *tb1* Mutations



Slow-Release Hydrogen Sulfide Seed Treatments Enhances Growth in Pulse Crops

Ethan Harbo

Project Goal



Develop a DBDTP seed treatment to boost plant growth and agronomic yield



Pulses treated with a DBDTP concentration curve at EARC



Chickpeas exhibiting physiological differences in greenhouse experiments.

- DBDTP Releases H_2S at a slow rate as a salt and in solution. Hydrolysis of DBDTP occurs within the plant.
- Exogenous applications of H_2S up to a point increase drought, heat, pathogen tolerance as well as plant growth.
- We are evaluating efficacy of seed treatments under normal field conditions and elevated stress conditions in greenhouse settings.

Revolutionizing Wheat with HaHB-4 Transcription Factor for Drought Tolerance in Montana

Luis Esquivel Cervantes

