

## **Grasslands are an Integral Part of American Identity**

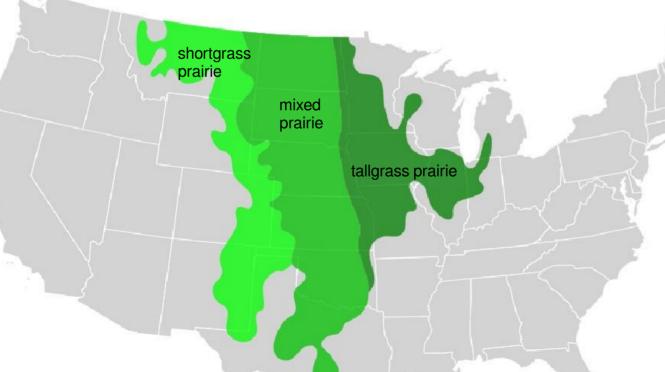
Banished from the thoroughfare and the field, it bides its time to return, and when vigilance is relaxed, or the dynasty has perished, it silently resumes the throne from which it has been expelled, but which it never abdicates.

-John James Ingalls

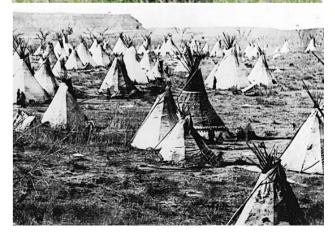




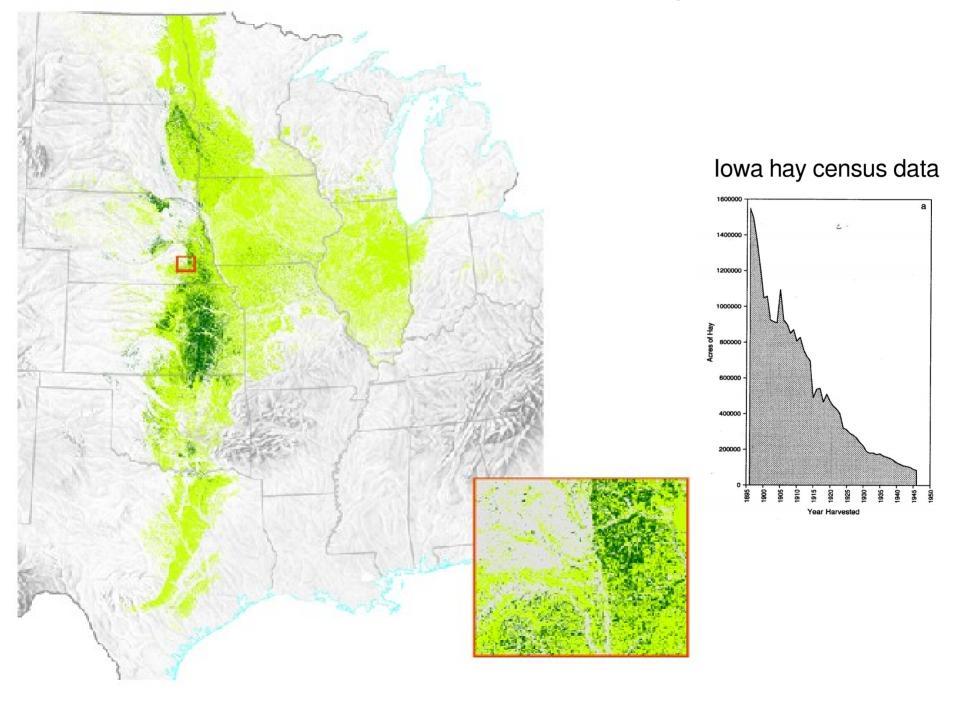








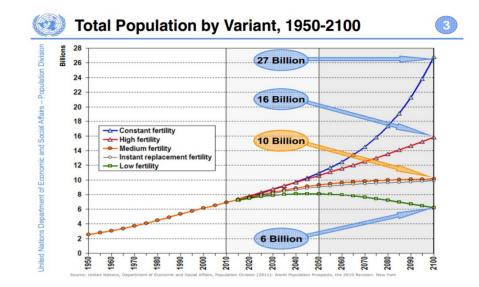
## 95% of the Great Plains Prairie Ecosystem is Gone

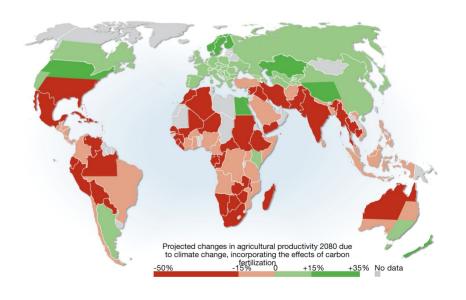


Smith, D.D. 1998 J. Iowa Acad. Sci.

# **Food Production Requirements**

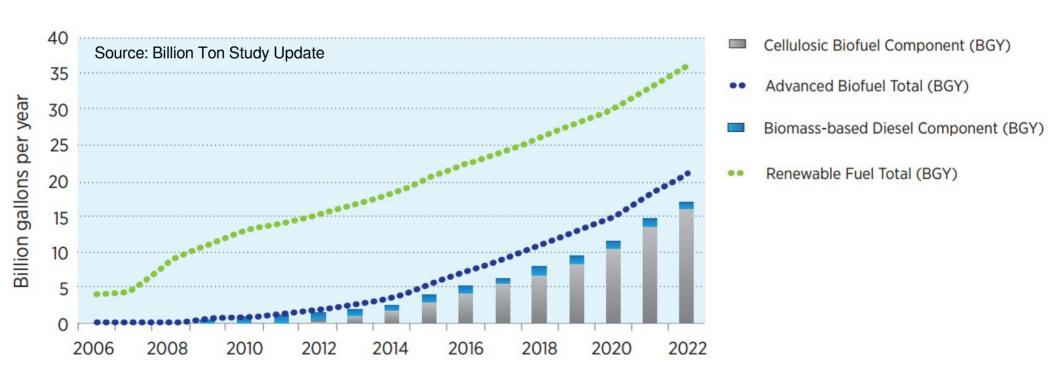
- •Assuming no change in population growth, food consumption patterns and food waste management, the following production increases must take place by 2050:
  - •cereals production must increase by 940 million tonnes to reach 3 billion tonnes;
  - •meat production must increase by 196 million tonnes to reach 455 million tonnes;
  - •and oilcrops by must increase by 133 million tonnes to reach 282 million tonnes.





**Renewable Energy Legislation/Mandates** 

Year	Milestone
2005	Billion Ton Study
	Energy Policy Act
2006	Advanced Energy Initiative
2007	Energy Independence and Security Act (EISA)
2008	Farm Bill Bioenergy Crop Assistance Program
2009	Establishment of BIWG
2010	USDA roadmap
	USDA Farm to Fly Partnership
2011	Billion Ton Study Update
2012	USDA Establishes Five Regional Biomass Research Centers
2013	USDA Farm to Fly 2.0 Partnership
2016	Billion Ton Update



# Second Generation Cellulosic Fuel Industry is Viable but Growing Slowly

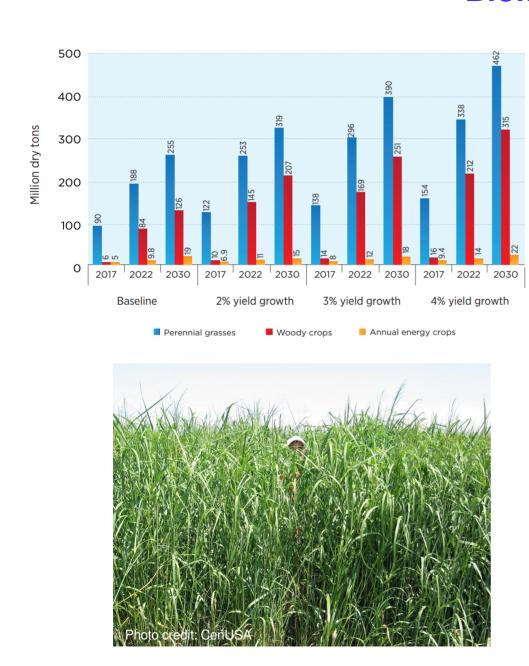
<ul> <li>Hugoton KS</li> </ul>	25-30 Mga/yr	Ethanol	wheat straw
<ul> <li>Alpena MI</li> </ul>	1 Mga/yr	Ethanol	wood residues
<ul> <li>Galva IA</li> </ul>	3.75 Mga/yr	Ethanol	corn kernel fiber
<ul> <li>Emmetsberg IA</li> </ul>	20-25Mga/yr	Ethanol	corn stover
Crescentino IT	13 Mga/yr	Ethanol	wheat straw, Arundo
<ul> <li>Nevada IA</li> </ul>	25 Mga/yr	Ethanol	corn stover
<ul> <li>Fulton MS</li> </ul>	(19 Mga/yr)*	Ethanol	multiple sources
<ul> <li>Reno, NV</li> </ul>	(11 Mga/yr)	JetFuel/Diesel	MSW
<ul> <li>Alagoas BZ</li> </ul>	(22 Mga/yr)	Ethanol	sugarcane straw
<ul> <li>Las Cruces NM</li> </ul>	(~7 Mga/yr)	<b>Green Crude</b>	Algae

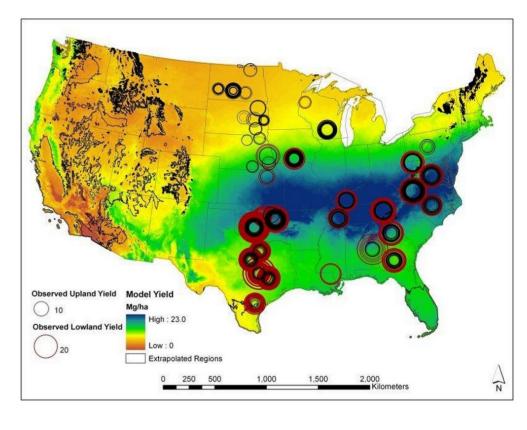
### US Cellulosic Biofuel Production (Mgal): EPA RFS data

Year	All Cellulosic	Ethanol	EISA target
2014	33.3	0.7	2500
2015	140	2.2	3000
2016 (to 7.1)	78.8	2.0	4000

<sup>\*</sup>Plants under construction

## Herbaceous Perennials are a Significant Source of Available Biomass





## The Big Three: Dominant Grasses of the Tallgrass Prairie









Switchgrass
Panicum virgatum

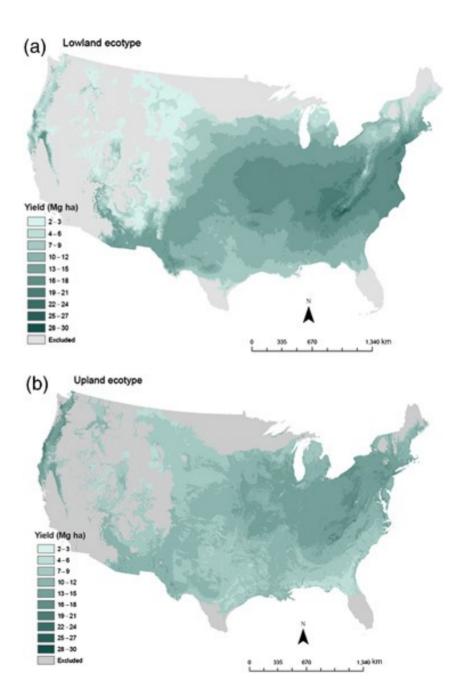


Indiangrass
Sorghastrum nutans



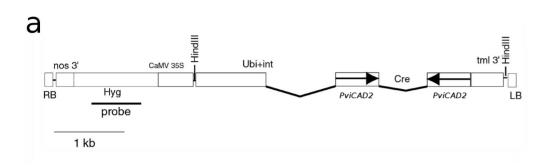
Big Bluestem Andropogon gerardi

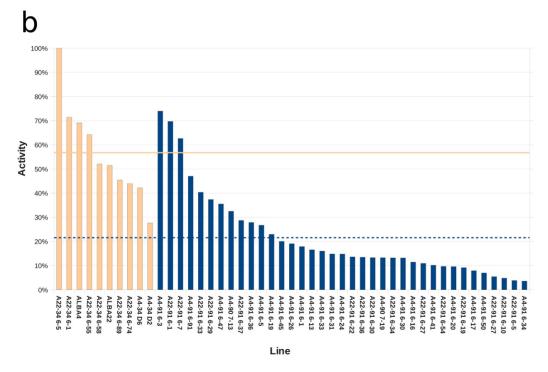
- Switchgrass as a feedstock is ready to hit the road in terms of it's utilization for renewable biofuels.
- Outcrossing and Polyploidy complicate genetics
- Upland/Lowland 8x/4x cytotypes
- It is amenable to transformation and molecular approaches
- It is highly variable and there is high potential for yield improvements however biotechnological approaches remain unproven but with high potential.

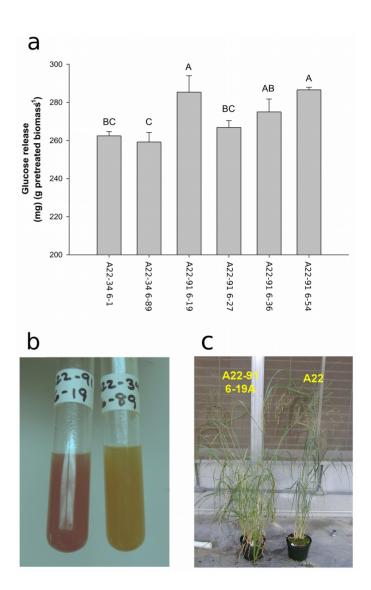


Jager et al. 2010

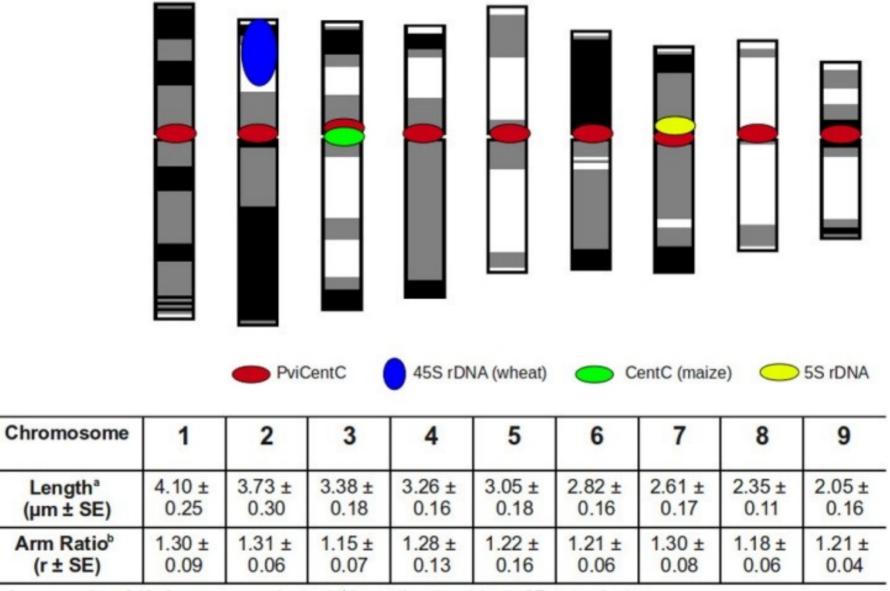
# Silencing of a Switchgrass Gene Involved in Lignin Formation Results in Greater Digestbility



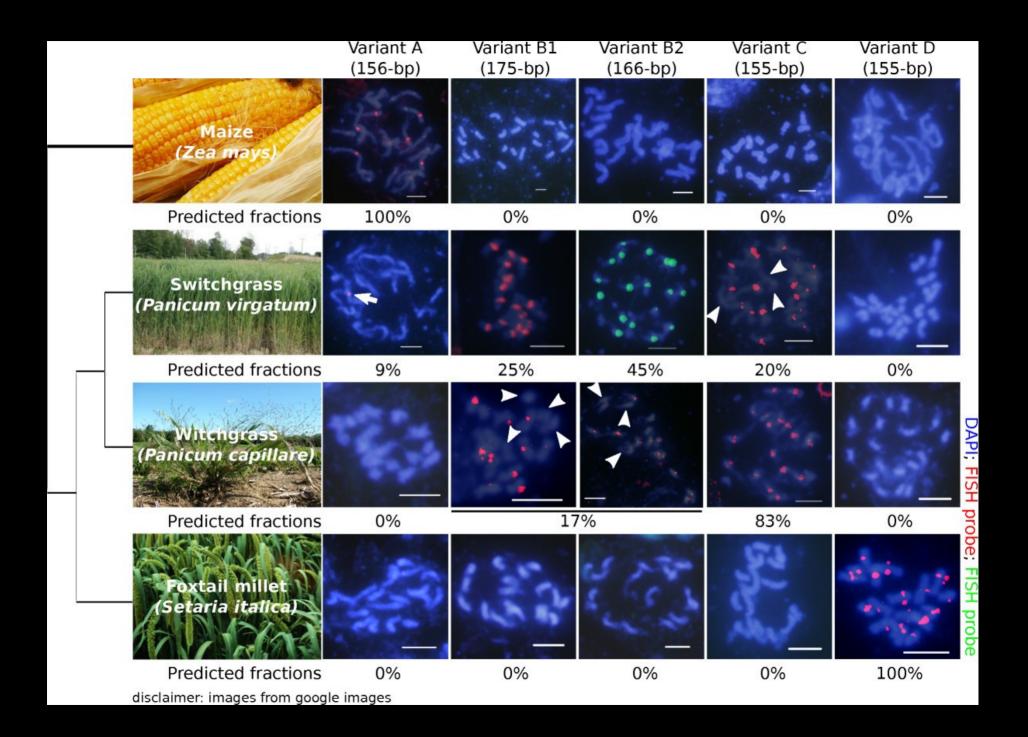




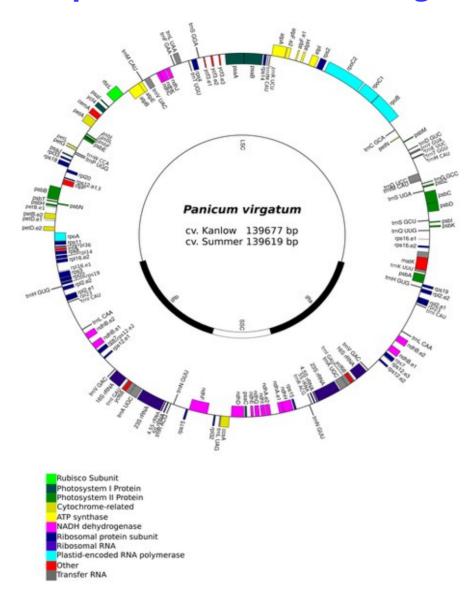
## Cytogenetic Features Reveal Structural Differences Between Subgenomes and Switchgrass Ecotypes



<sup>\*</sup>Average value of 10 chromosome pairs (μm); \*Arm ratio r (long/short); SE = standard error



### Map of the Chloroplast Genome of P. virgatum cv. Kanlow

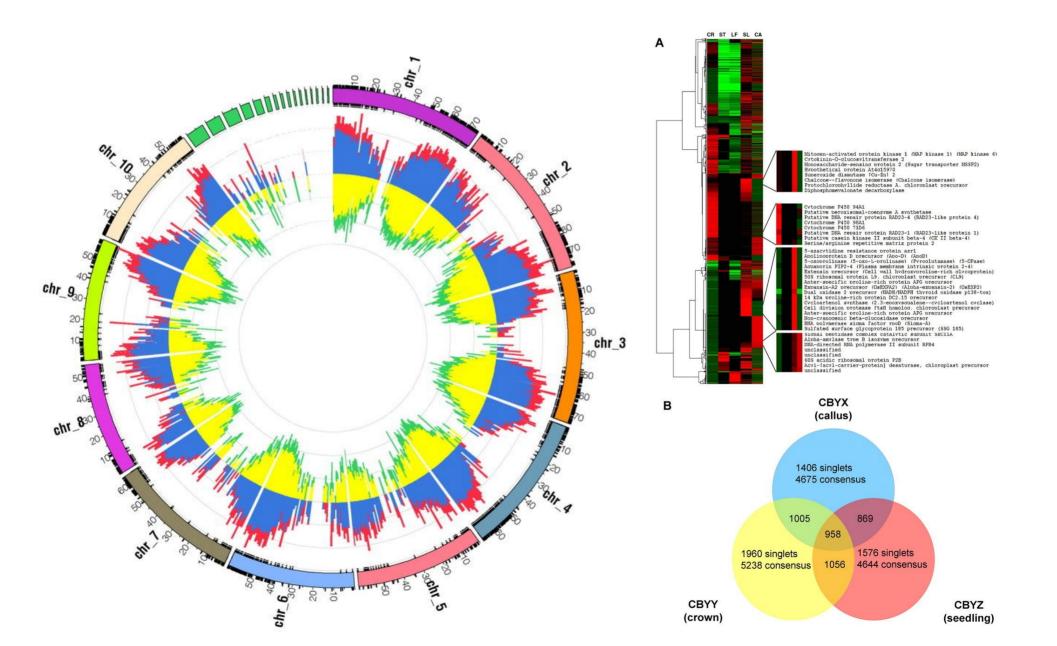


Young HA, Lanzatella CL, Sarath G, Tobias CM (2011) Chloroplast Genome Variation in Upland and Lowland Switchgrass. PLoS ONE 6(8): e23980. doi:10.1371/journal.pone.0023980

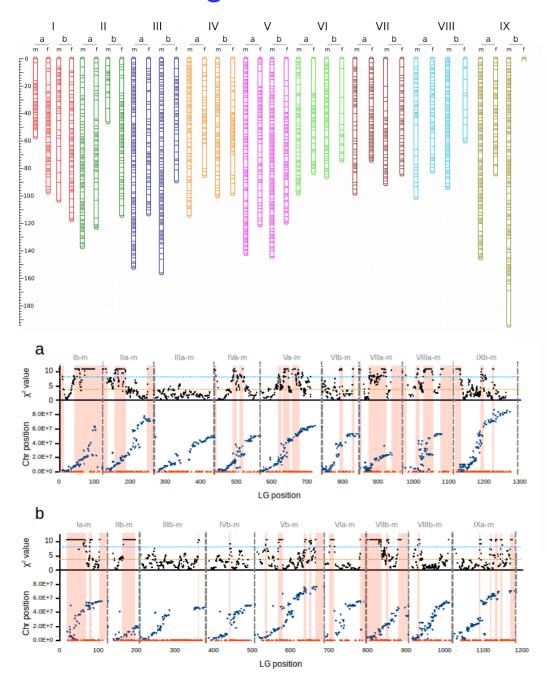
http://www.plosone.org/article/info:doi/10.1371/journal.pone.0023980

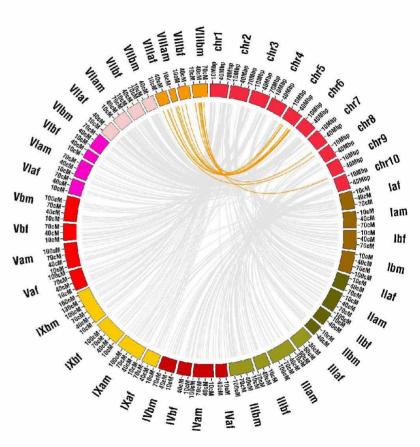


# **Sequencing of the Switchgrass Transcriptome Polymorphism, Duplication, and Conservation**



## Linkage Mapping has Allowed Comparisons Among Grasses, Across Subgenomes and Has Facilitated Genome Assembly





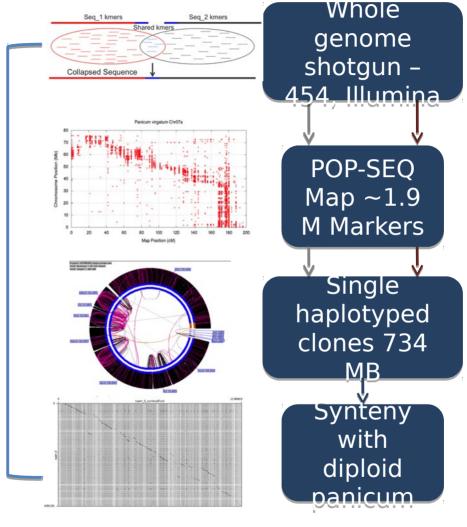
## Switchgrass Genome



Switchgrass Panicum virgatum

Outbred tetraploid 1.4 Gb genome Advanced Custom Assembly & Integratio

	V1.0	V2.0
Chromosome Bases	636.1 Mb	1.031 Gb
Chromosome Contig N50	9.2 Kb	33.1 Kb
Unassigned	48.0%	11.7%



#### **Phenotypic selection**

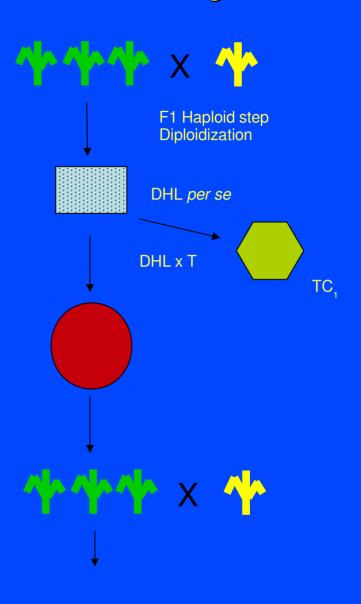
## Space-planted source nursery Mass. Sel. Replicated polycross C1 of selected genotypes **Sward** Trial Replicated space planted Half-sib progeny evaluation nursury **Sward** Trial Cycle 2

2 generations 6 years

Cycle1

B&W FS

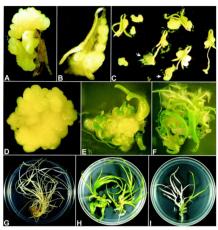
#### **DH Breeding**



2 generations 6 years 2x genetic variance

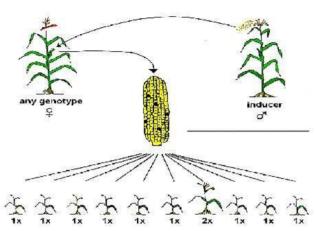
## **Uniparental Genome Elimination**

#### Gametopyte culture



canola/barley

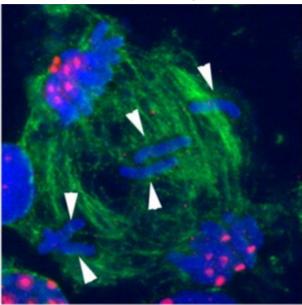
#### Inducer lines



hybrid seed to reject

approx. 5 % maternal haploids or doubled haploid:

#### CMCE (Arabidopsis only)



Sanei 2011 PNAS

#### Interspecific crossing



Hordeum bulbosum



Masu salmon



Nasonia

### **Acknowledgements**

#### **UC Davis**

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Samuel Roberts Noble Foundation

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#### **WRRC**

**Roger Thilmony** 

**John Vogel** 

Sanwoong Yoon (genome editing)

**Christina Lanzatella Craig (Mapping, Plant Transformation)** 

**Hugh Young (FISH)** 

**Jason Fiedler (Mapping)** 

Miki Okada (Mapping)



