Revealing the Past and Present of Bison Using Genome Analysis

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Presentation outline

1. What can genetics teach us about the history of the Yellowstone bison herd?

2. What genetic information can we gain from analyzing bison fecal samples?

3. How can we develop more powerful genetic testing technologies?

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The Yellowstone bison herd is unique



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- Well documented recent bottleneck



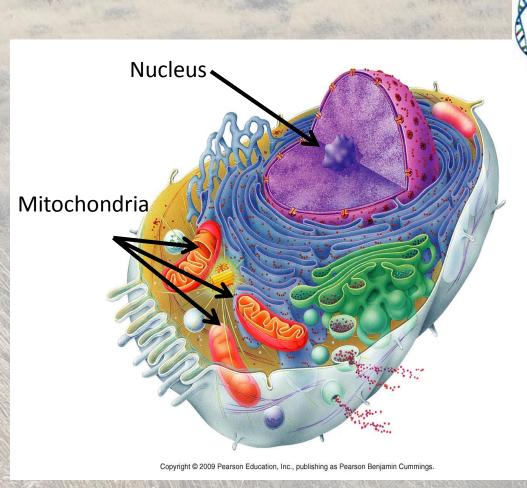
- The Yellowstone bison herd is unique
- Well documented recent bottleneck
- Intensive management and population control is necessary

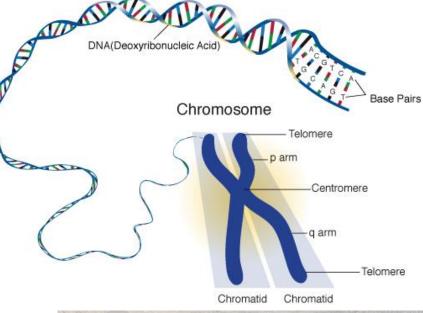




Objective: Analyze the genetic diversity and population structure of Yellowstone bison

The bison cell

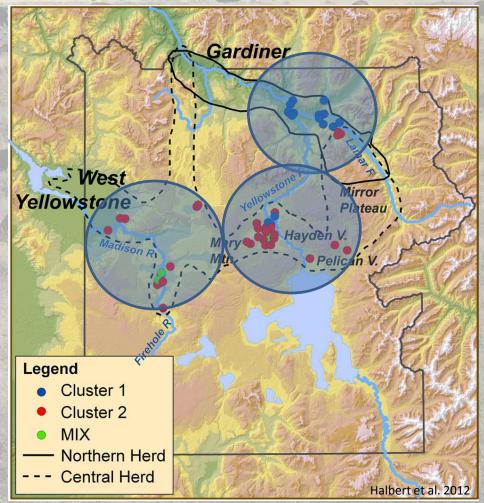




Population structure

1996

- Since the late 1990s:
 - Population size has increased significantly
 - Migration within the park has become commonplace
 - Leading topopulationdifferentiation loss

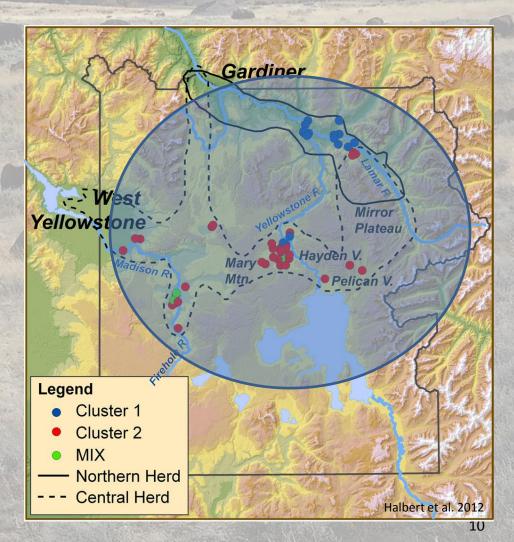


2003

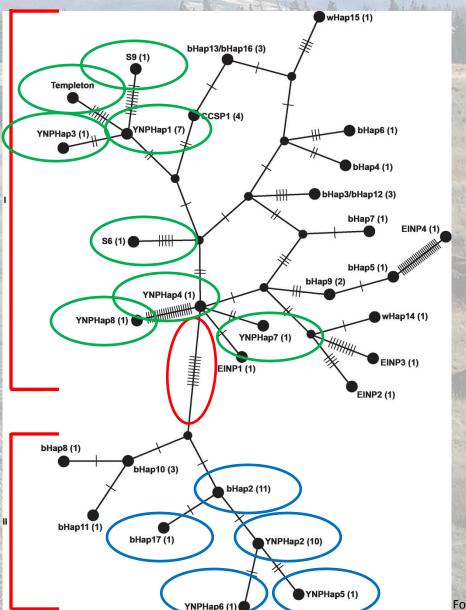
2012

Population structure

- Increase in migration results in:
 - Reduced diversity
 between populations
 - Increased diversity in the whole population
- High population genetic diversity, but only small differences between individuals
- Signature of healthy recovery after a major bottleneck



Bison diversity beyond Yellowstone



- Two groups with nearly equal number of bison separated by 10 mutations
- Yellowstone bison are also evenly split
- What causes this dichotomy if not current population structure?

The history of Yellowstone bison

- Lowest number of bison in Yellowstone recorded in 1902
- Remaining 22
 indigenous animals
 were supplemented by
 21 introduced bison
 initially corralled in
 Northern Yellowstone
- Released in 1915

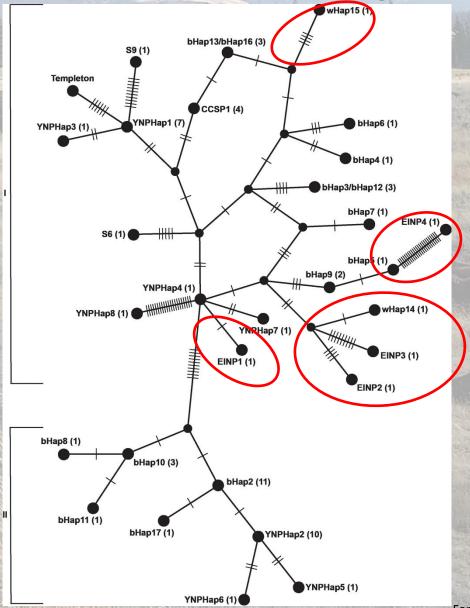


The history of wood bison



- Wood bison is a bison subspecies with clear conformational differences
- Large-scale
 displacement of plains
 bison into Canada in the
 1920s
- Some Canadian bison might look like a wood bison but have plains bison mitochondria

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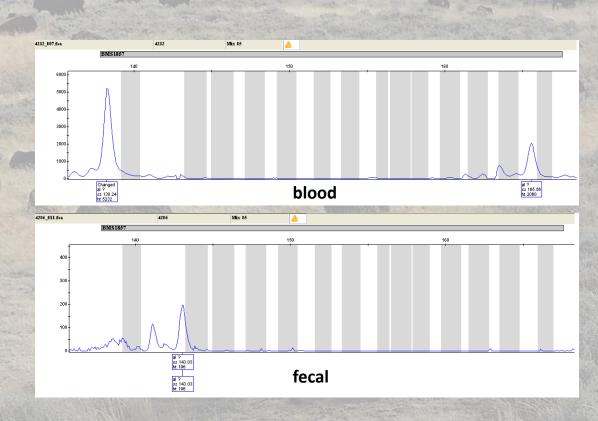
- Fecal DNA samples are widely used in wildlife research
- However, no comprehensive direct comparison between high quality and fecal DNA has ever been done



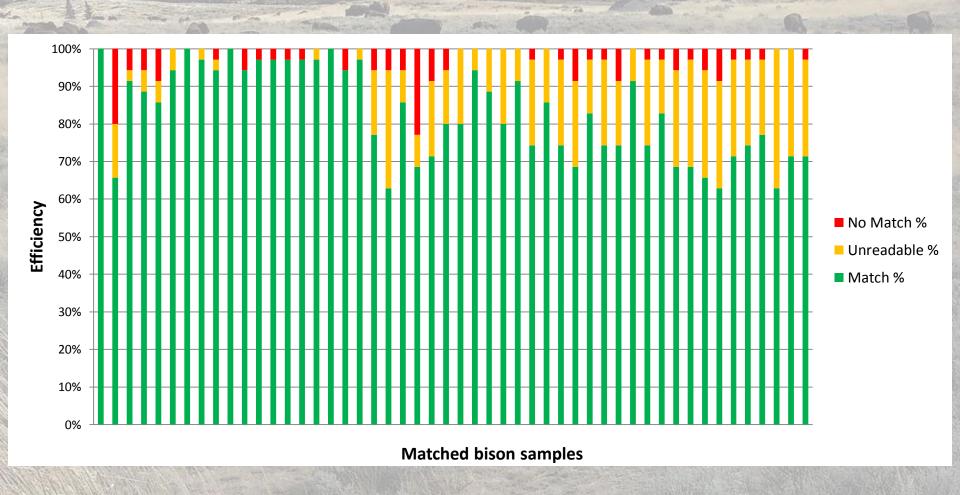
Objective: Analysis of fecal DNA and assembly of a panel of validated markers for genetic testing in bison

False alleles

- Occurs when neither allele matches
 - Misidentification of individual
 - Incorrect results
- 0.46% of cases, negligible



No evidence of contamination



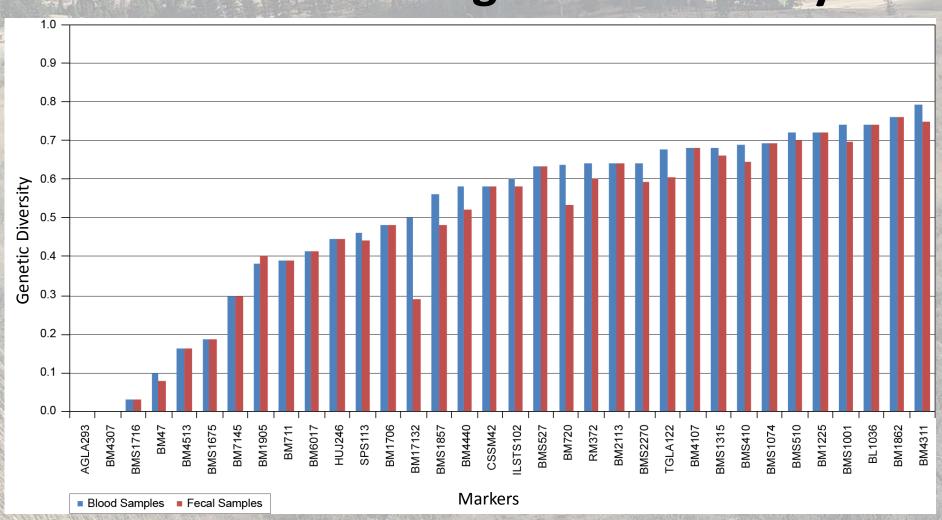
Allelic dropout

- Occurs in cases when only one allele shows up, leading to biased results
 - False identification
 - Underestimateperceived genetichealth
 - Increased inbreeding
 - Lower genetic diversity

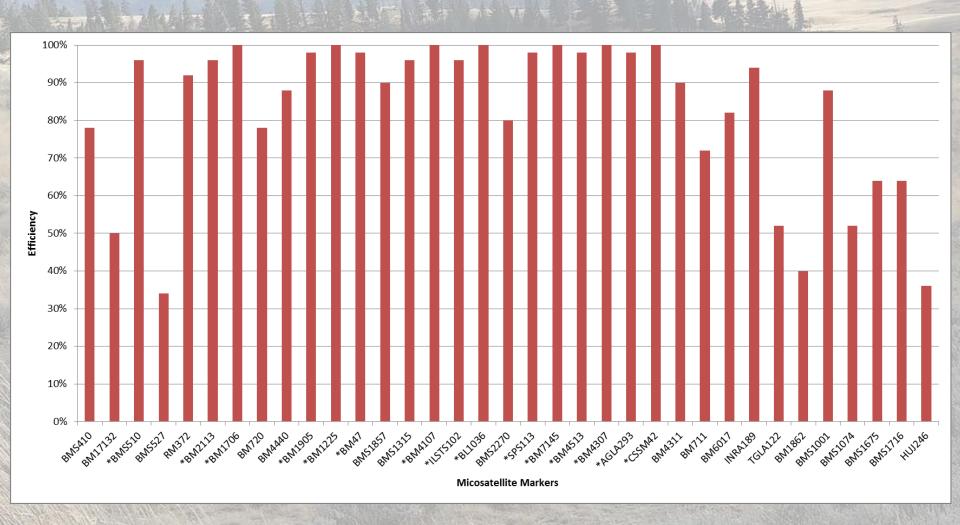


2.3% of cases

Allelic dropout can lead to underestimated genetic diversity



Genetic testing panel



Genetic testing panel

	The state of the s		The sales of the s
	Percent	Percent	
	matching	matching	
Marker	(excl. reruns)	(incl. reruns)	Chromosome
BM7145	100%	100%	1
BM4307	100%	100%	1
BM2113	96%	96%	2
CSSM42	100%	100%	2
AGLA293	100%	98%	5
SPS113	98%	98%	10
BL1036	100%	100%	14
BM4513	100%	98%	14
BM1706	100%	100%	16
BM1225	100%	100%	20
BM4107	100%	100%	20
BM1905	98%	98%	23
BM47	98%	98%	23
ILSTS102	98%	98%	25
BMS510	96%	96%	28



Chance of two bison having the same alleles at all these markers: 2.318 x 10⁻⁶

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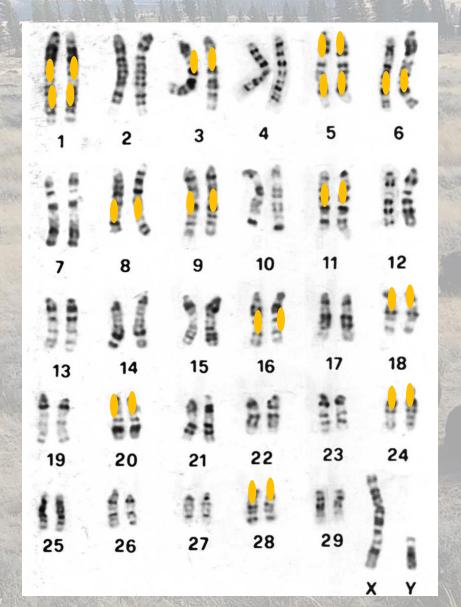
Current technologies

- Developed by the Derr lab over the last 20 years
- Based on nuclear microsatellites to test for:
 - parentage/kinship
 - genetic diversity and inbreeding
 - lineage ancestry
 - domestic cattle introgression
- Tens of thousands of bison successfully tested

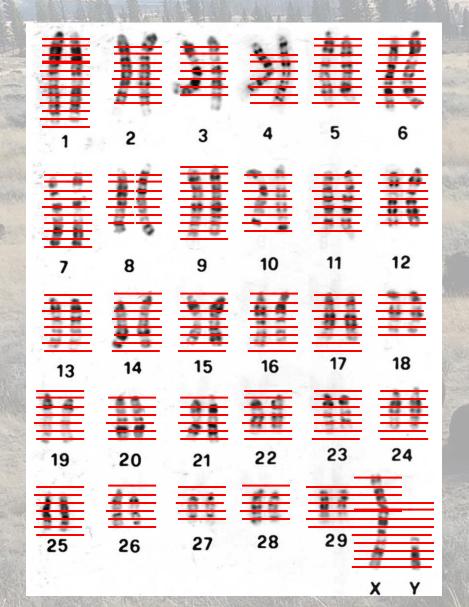


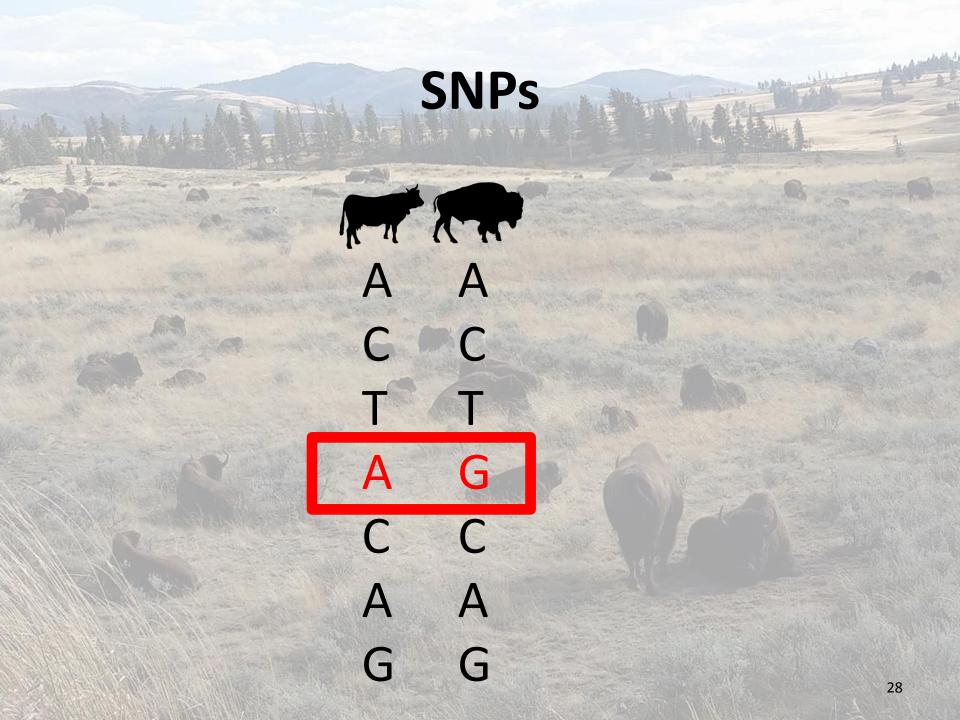
We are currently working on the development of a new Single Nucleotide Polymorphism (SNP) genetic tests for bison

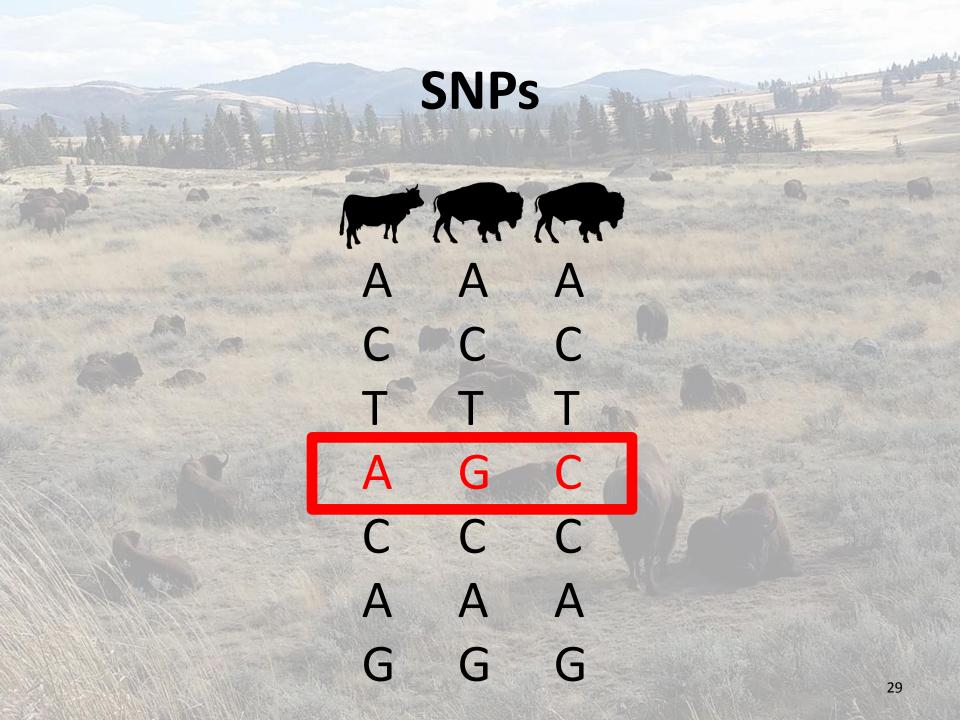
Microsatellites vs SNPs



Microsatellites vs SNPs







Advantages of SNP-based testing

- Increase in marker coverage across the bison genome resulting in a tremendous increase in genetic information per test
- SNP genotyping is cheaper than traditional microsatellite genotyping (about ½ the cost)
- SNP genotyping provides a much faster turnaround time in the laboratory (days <u>vs</u> weeks)
- Allows for future trait-specific marker detection

Final summary

- Yellowstone bison population genetic analysis reveals a healthy herd with high diversity, migration, and a constantly increasing population size
- We have developed a panel of markers for genotyping bison from fecal samples, but special care must be taken when dealing with fecal DNA
- Development of a SNP-based genotyping assay for a wide array of applications is under development



Acknowledgements



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