

Modeling spatio-temporal, genetic, and environmental patterns of anthrax outbreaks in Texas and Montana



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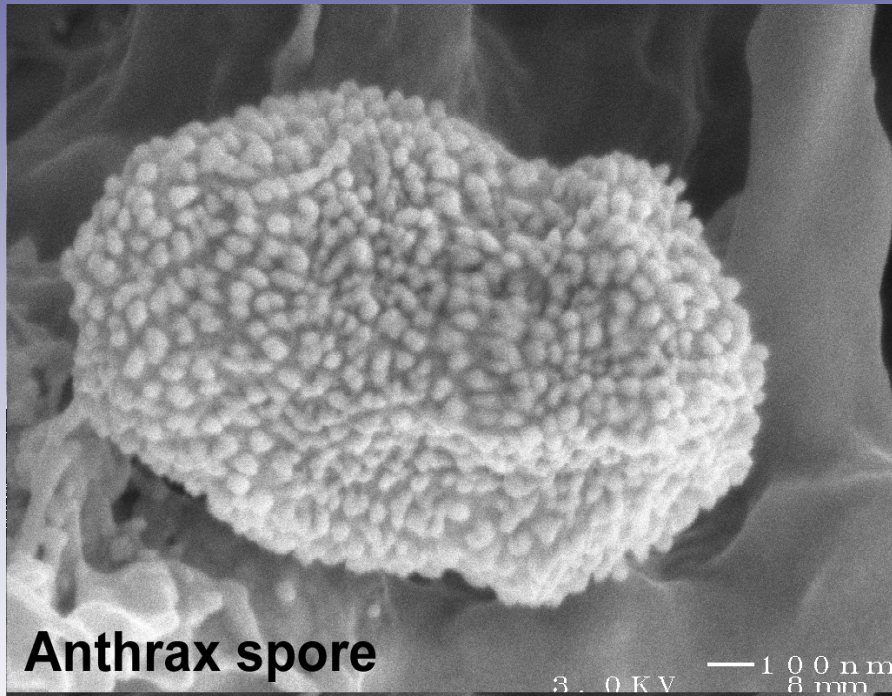
³MRIGlobal - Florida Division, Palm Bay, Florida USA

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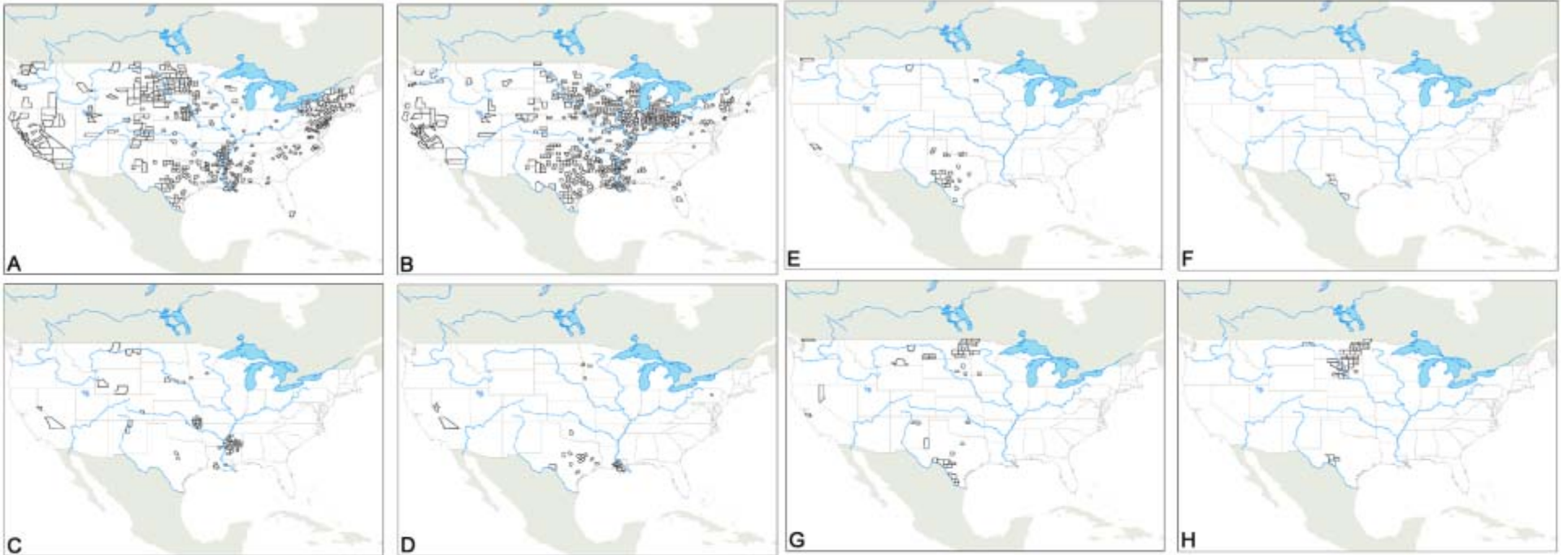
⁵ Turner Endangered Species Fund and Turner Enterprises, Inc, Bozeman, Montana, USA

⁶ Department of Environmental Sciences, School of the Coast and Environment, Louisiana State University, Baton Rouge, Louisiana USA

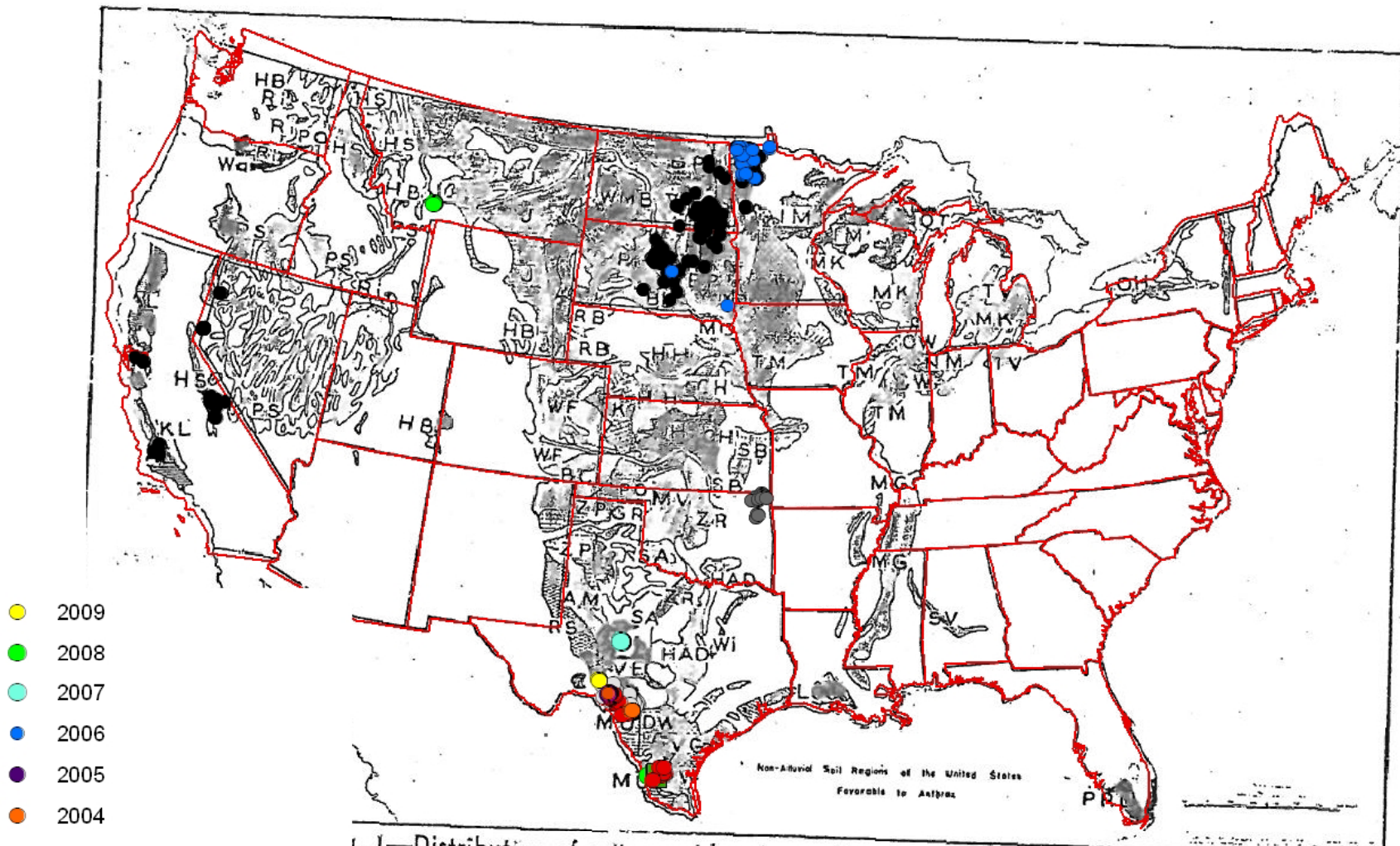
Two views of *Bacillus anthracis*



Spatial Epidemiology: Mapping Space-Time



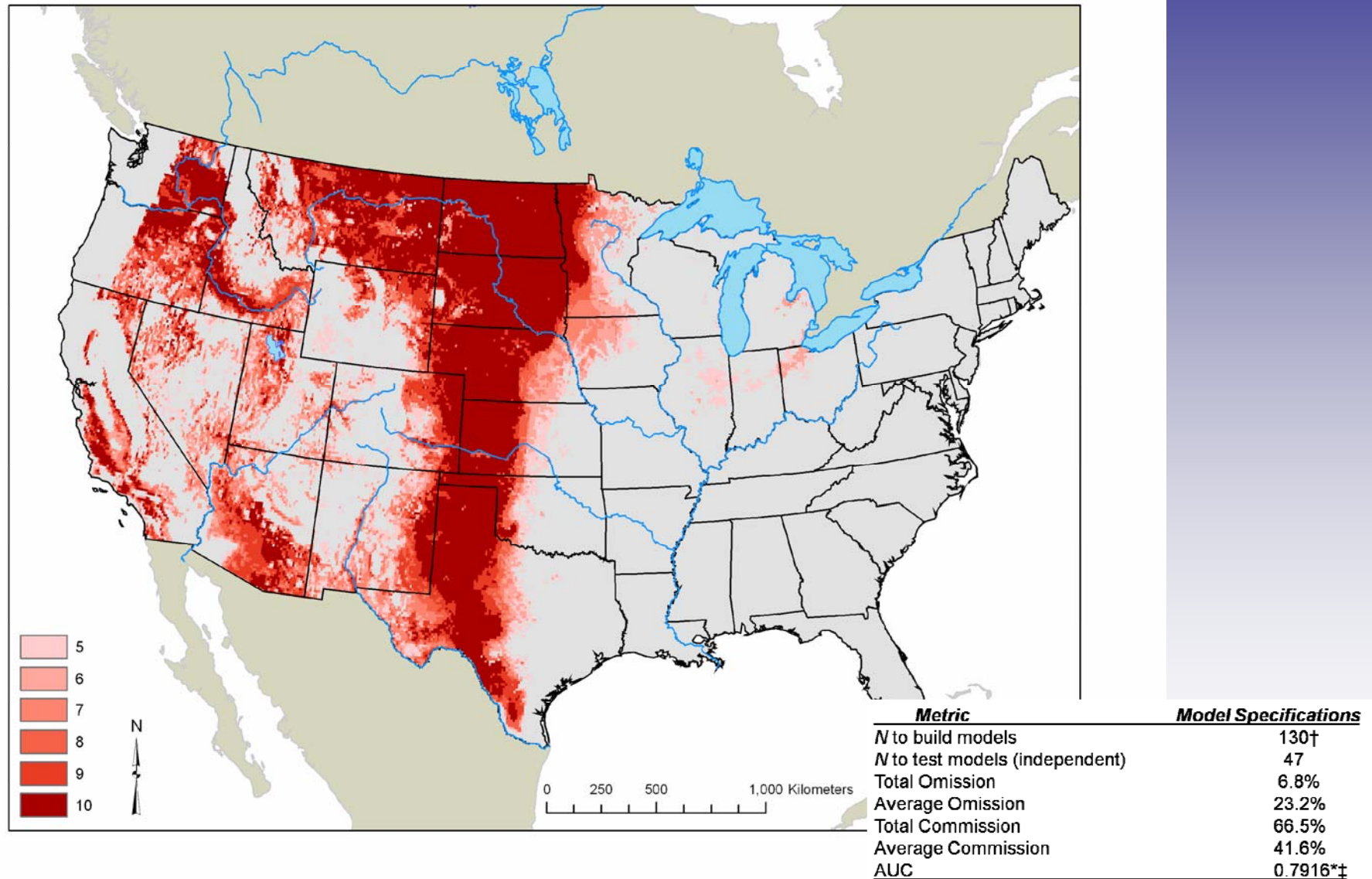
Blackburn (2006)



1. I—Distribution of soils considered capable of supporting anthrax.

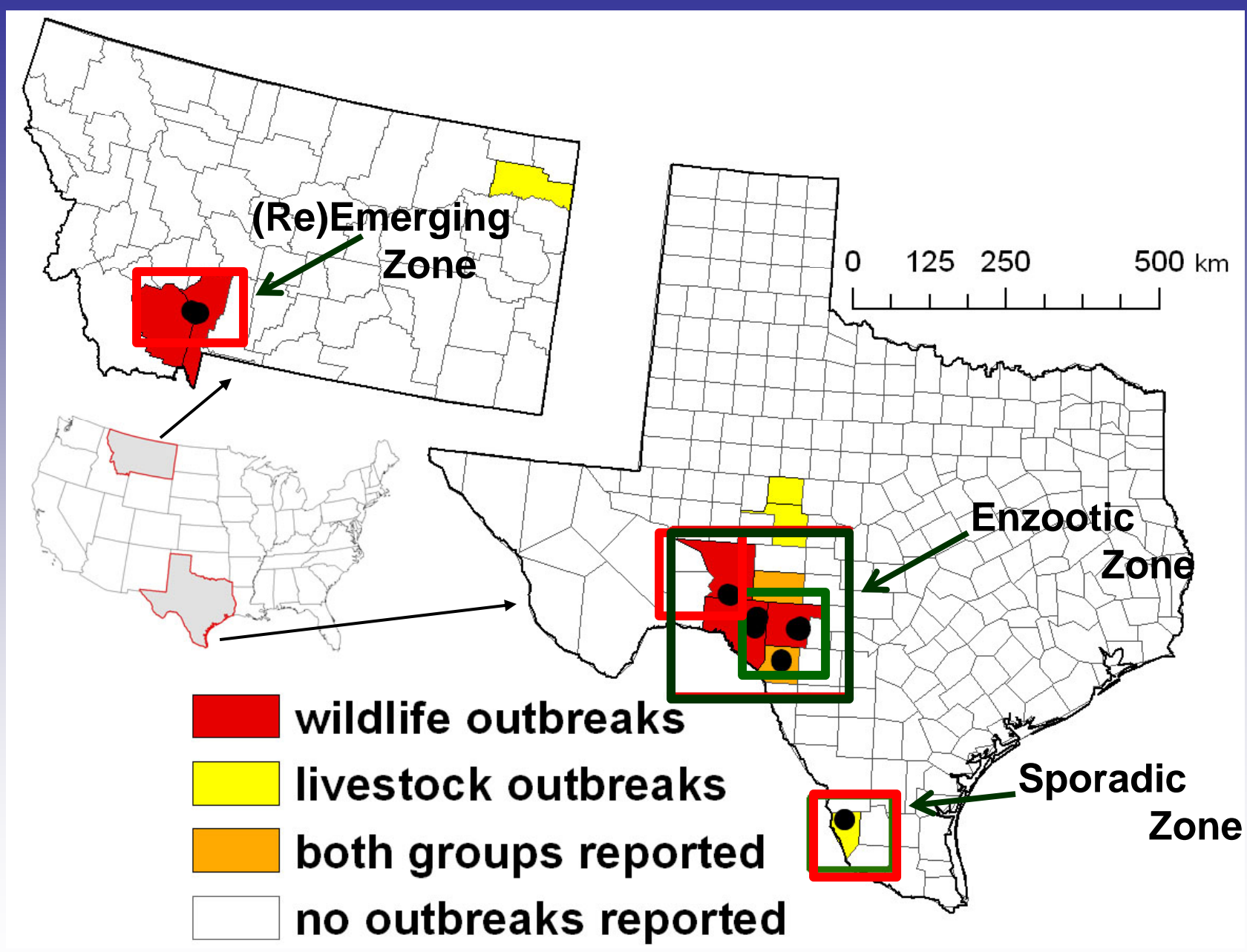
Base map from Van Ness and Stein 1956

Predicted distribution of *B. anthracis*



Blackburn et al. (2007)

†*N* was divided into 50% training / 50% testing at each model iteration
* $z=10.503$ ($p<0.01$), ‡SE = 0.0394





Geography of the *Sporadic Zone*

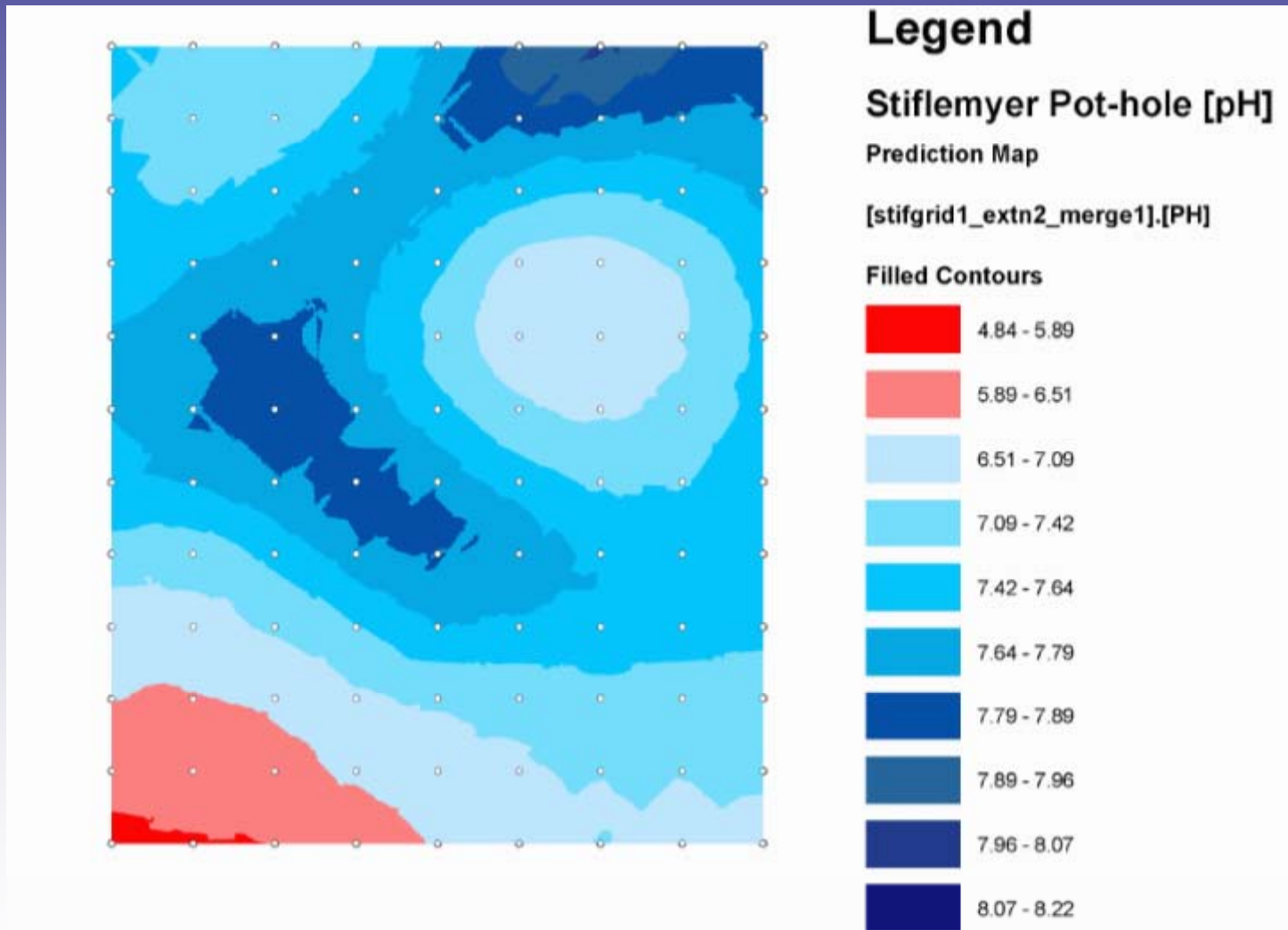
Q: Are there landscape characteristics that differentiate the epizootiology of south and west Texas? Can we detect them?



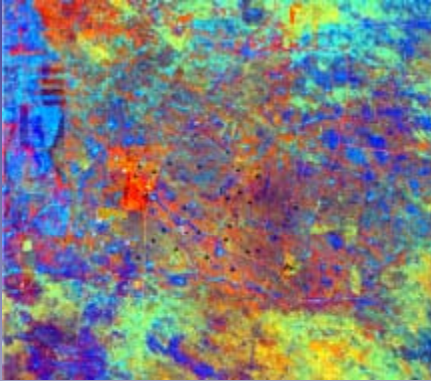


Photos: Deer © Jason Blackburn; Cows © M. Hugh-Jones

Mapping limiting factors to persistence

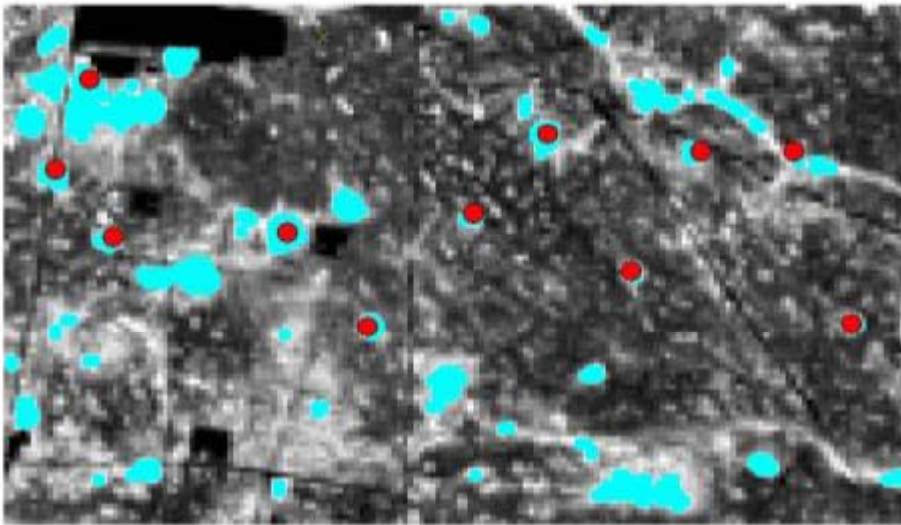


Defining local clusters: Getis' G statistic

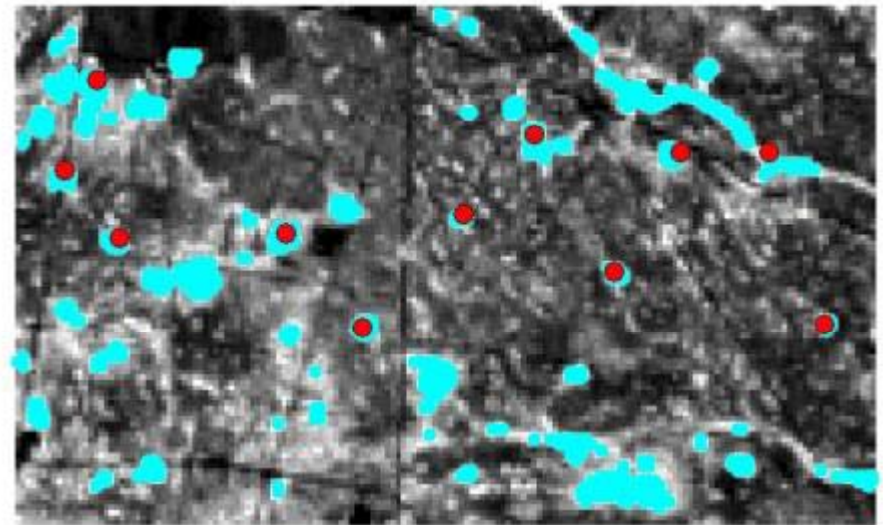


$$G_i^*(d) = \frac{\sum_j w_{ij}(d) \cdot x_j - w_i^* \cdot \bar{x}}{S \cdot \left\{ \left[(nS_{li}^*) - W_i^{*2} \right] / (n-1) \right\}^{1/2}}$$

Can we detect the spectral signature of potholes? Do those values cluster on the landscape?



February 14, 1999 (WINTER)



November 13, 1999

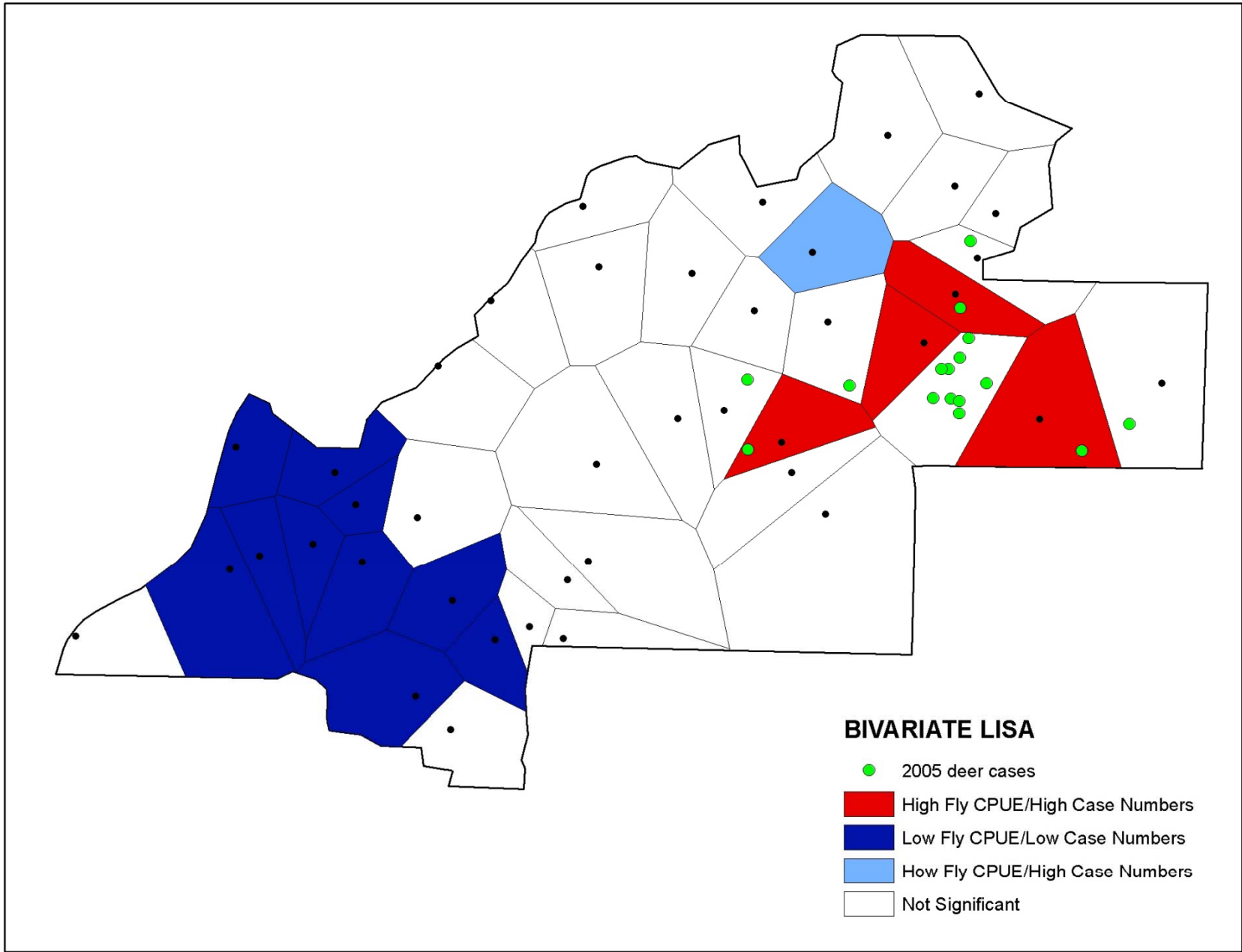
Geography of the *Enzootic Zone*

Q: What is the *Enzootic Zone* and how does it differ from the *Sporadic Zone*?

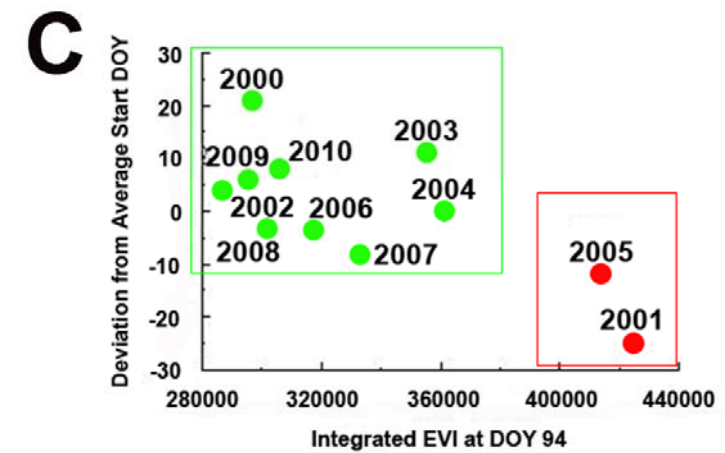
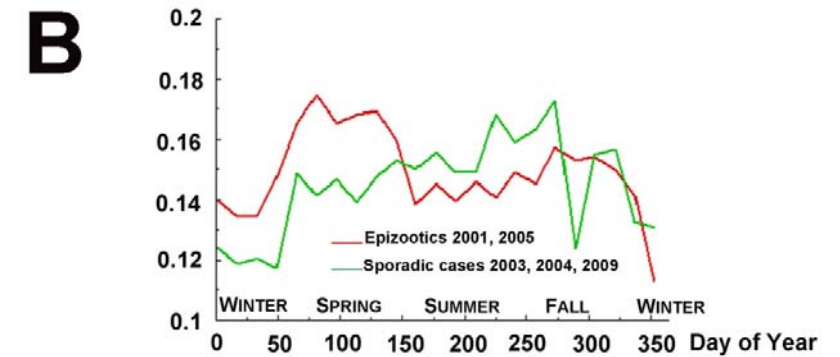
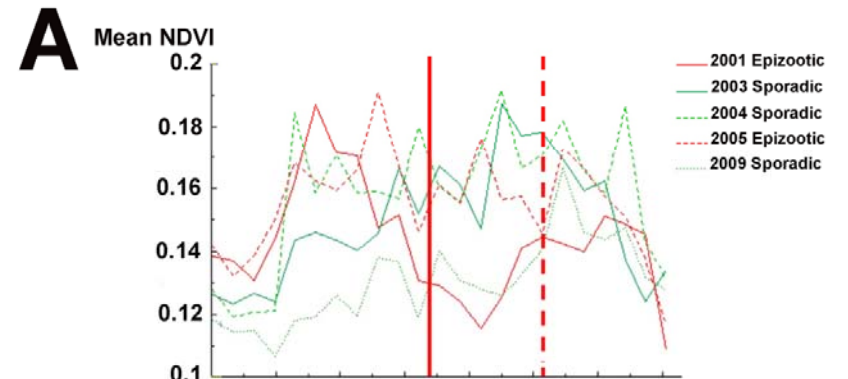
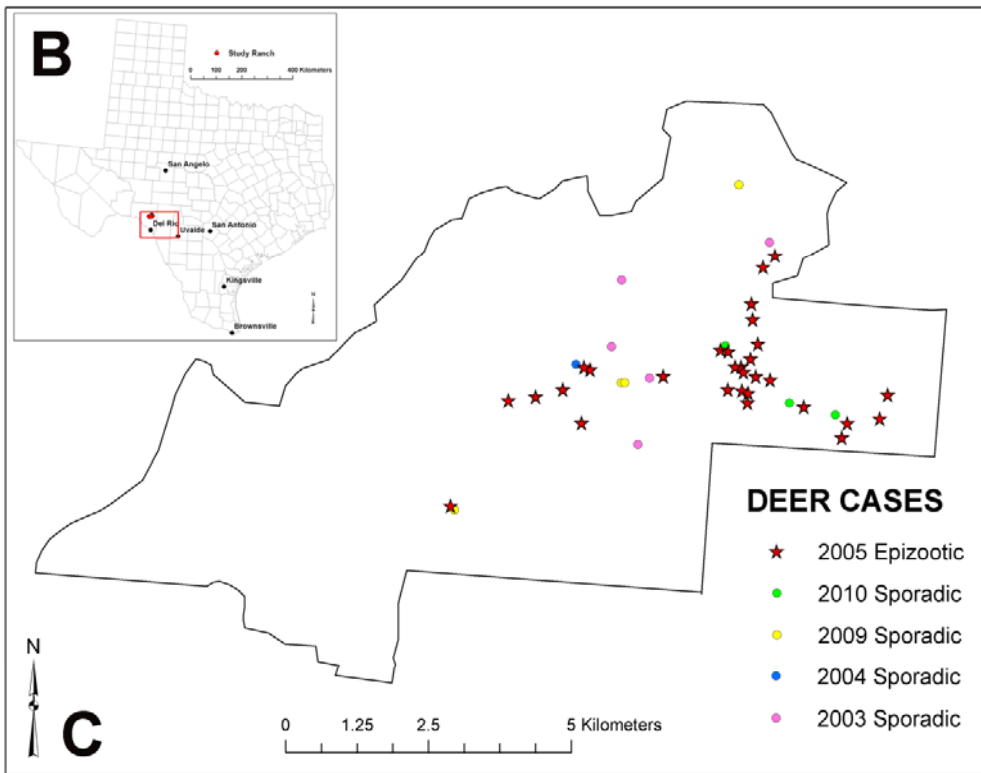
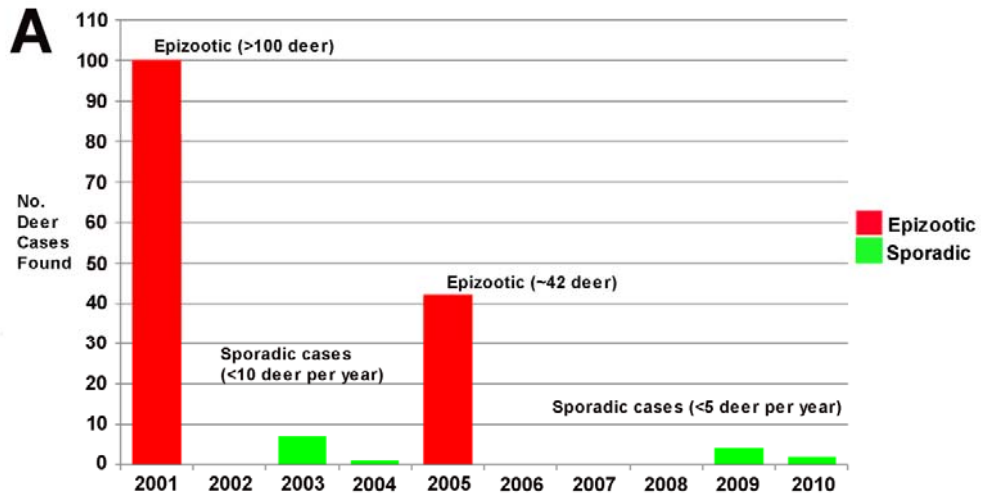




Epizootic	Mortality	Prevalence
<i>TEXAS 2009</i>		
White-tailed deer	30	~10.0
<i>TEXAS 2005</i>		
White-tailed deer	49	~3.2
<i>TEXAS 2001</i>		
White-tailed deer	~100	~7.7
<i>ITALY 2004</i>		
Red deer	8	~17.7



Blackburn et al. (In Review)



CASE MULTIPLIERS

NECROPHILIC FLIES IN OUTBREAKS

Q: Is there evidence for flesh eating flies moving contamination from carcass to the surrounding environment (browse, graze, etc)?





TRANSMISSION

Blow flies feed on carcasses and pick up *Bacillus anthracis* through contact with body fluids or through direct ingestion

2010 FIELD DATA



Blackburn et al. (2010)

Passing browsers may come into contact and become infected through ingesting contaminated browse.

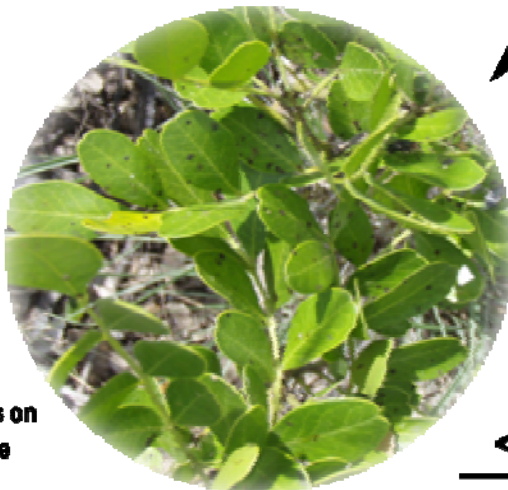


Blow flies serve as a localized mechanical vector moving bacteria from carcass to surrounding vegetation.



2010 FIELD DATA

2010 FIELD DATA



Flies deposit feces or emesis on leaves that may potentially be eaten by other browsers

< 3 m

While not fully understood, fly feeding ecology and movement studies suggest this to be a local transmission phenomenon

PERSISTENCE IN THE ENVIRONMENT

Q: What do we know about persistence?







The *(Re) Emerging Zone* in Western Montana

**Q: How long has the disease been here? Can we study “Spillover”?
What is the host system? What are the genetics?**



**YOU ARE ENTERING
A LIVESTOCK
QUARANTINE AREA.
DO NOT unload animals
until you reach
Forest Service boundary.**



Bison (sex)	Mortality	Prevalence
Mature cows 3+ (F)	208	10.25
2006 born cows (F)	1	0.18
herd bulls 6+ (M)	74	27.41
2007 stocker & born replc (M)	15	2.23
Calves (F, M)	0	0



FEMALE TOTALS PREVALENCE

8.11

MALE TOTAL PREVALENCE

9.44

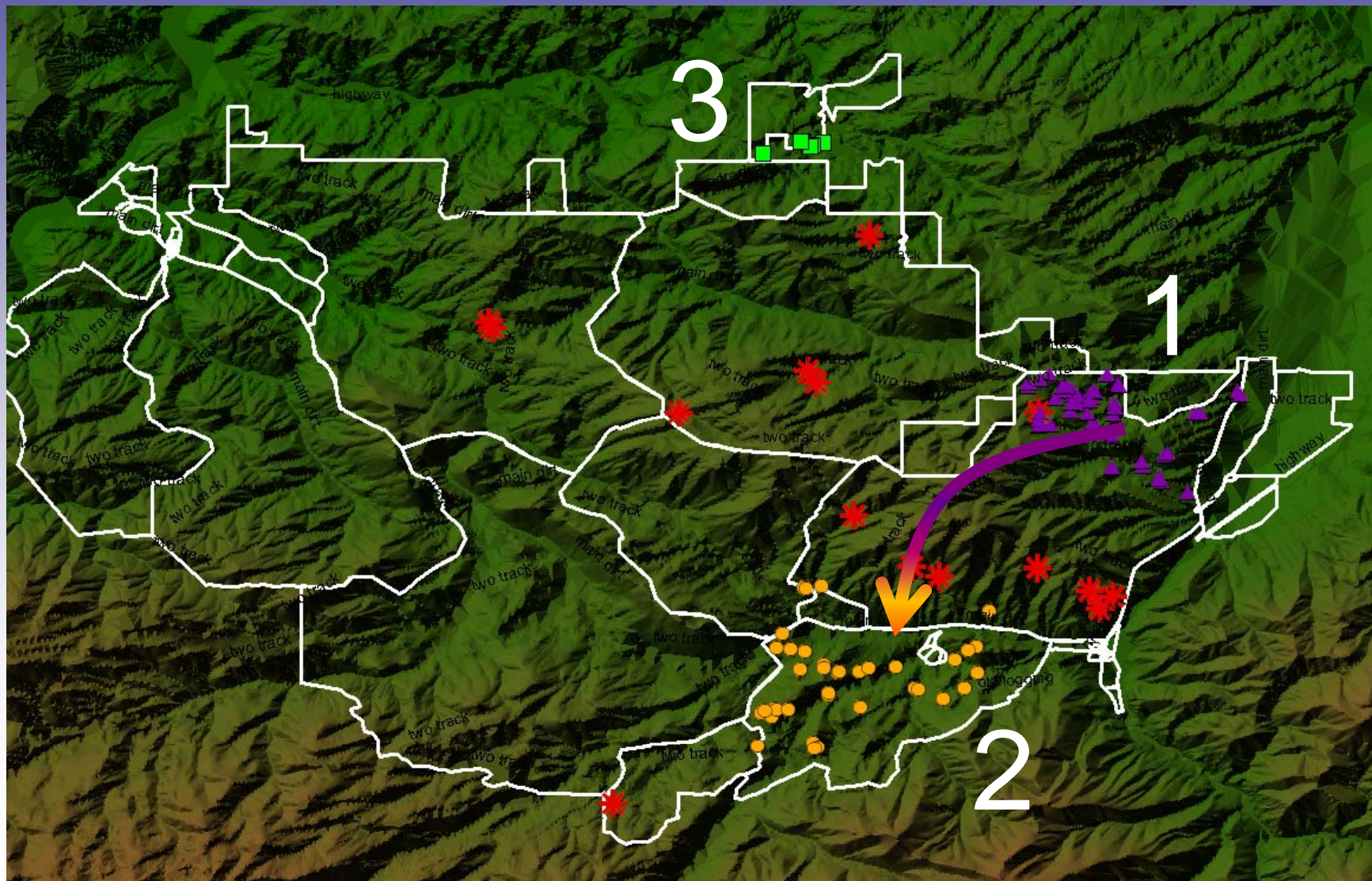
**BISON HERD TOTAL PREVALENCE
(excluding calves)**

8.47

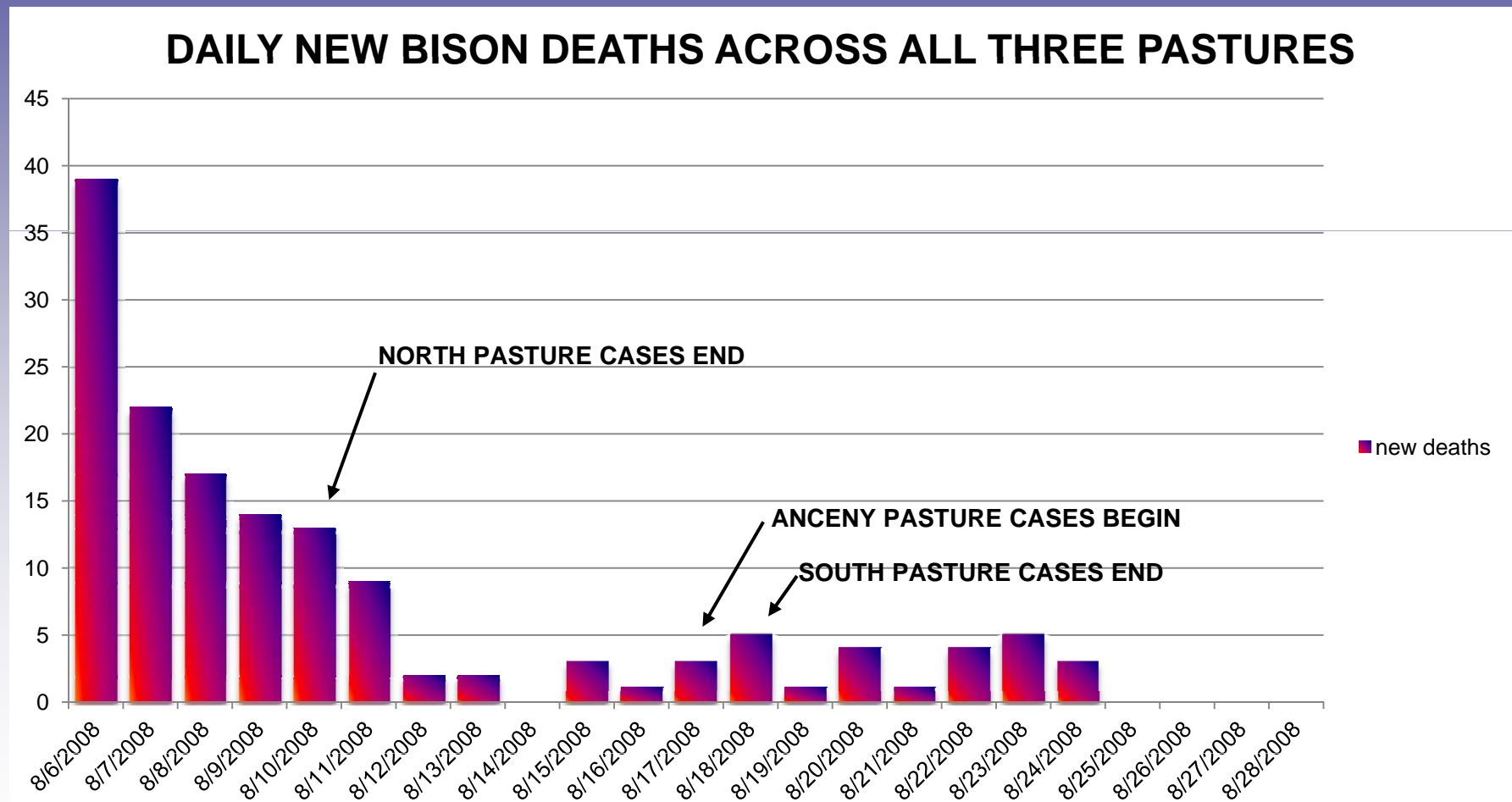


Elk	Mortality	Prevalence
Found during epizootic	18	13.95
Found during 2009-2010	27	
Total	45	34.88
Deer		
Buck	1	1.54
Doe	1	1.18
Total	2	1.33

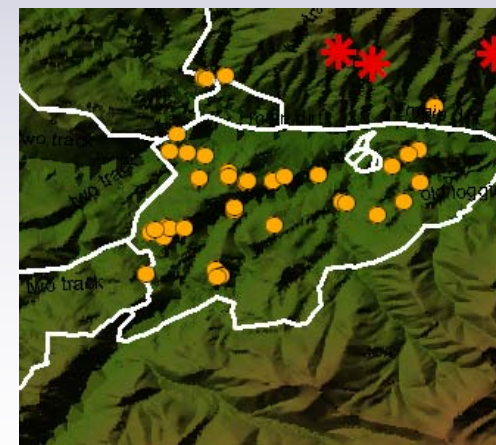
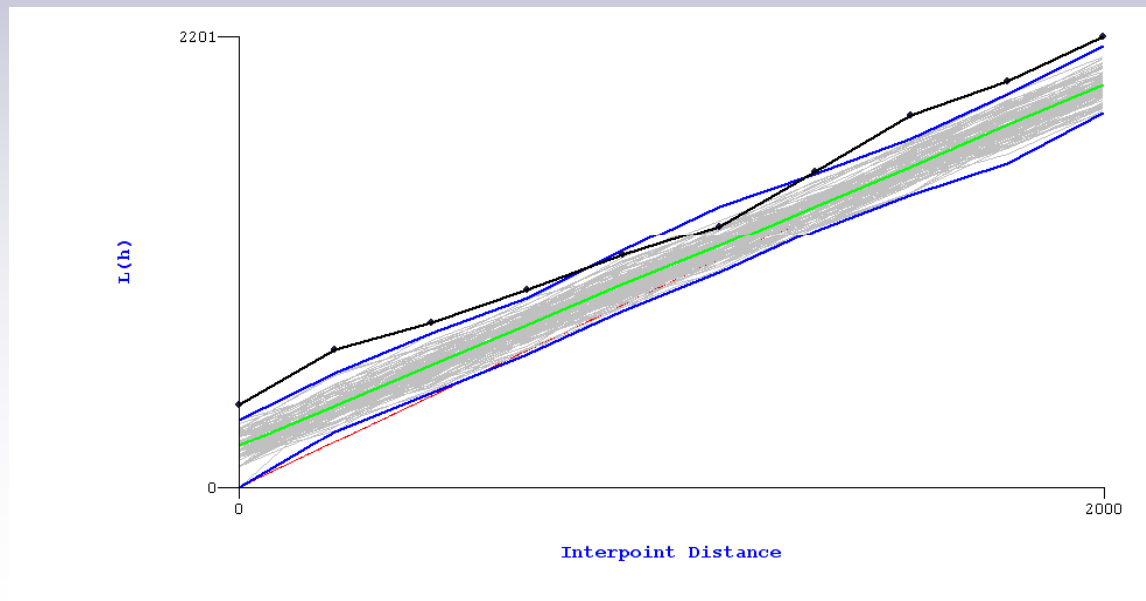
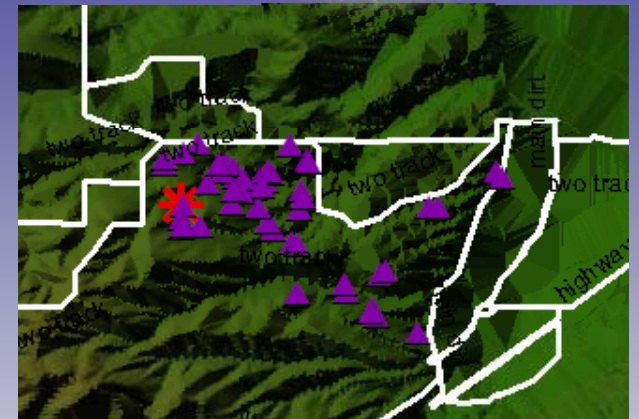
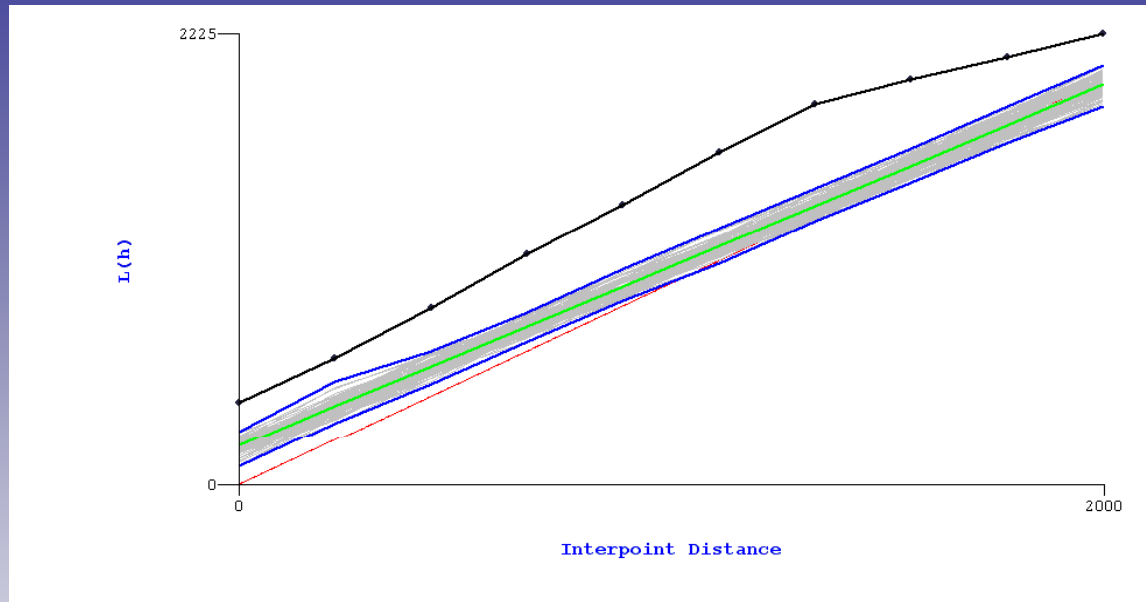
Spatial patterns of bison cases



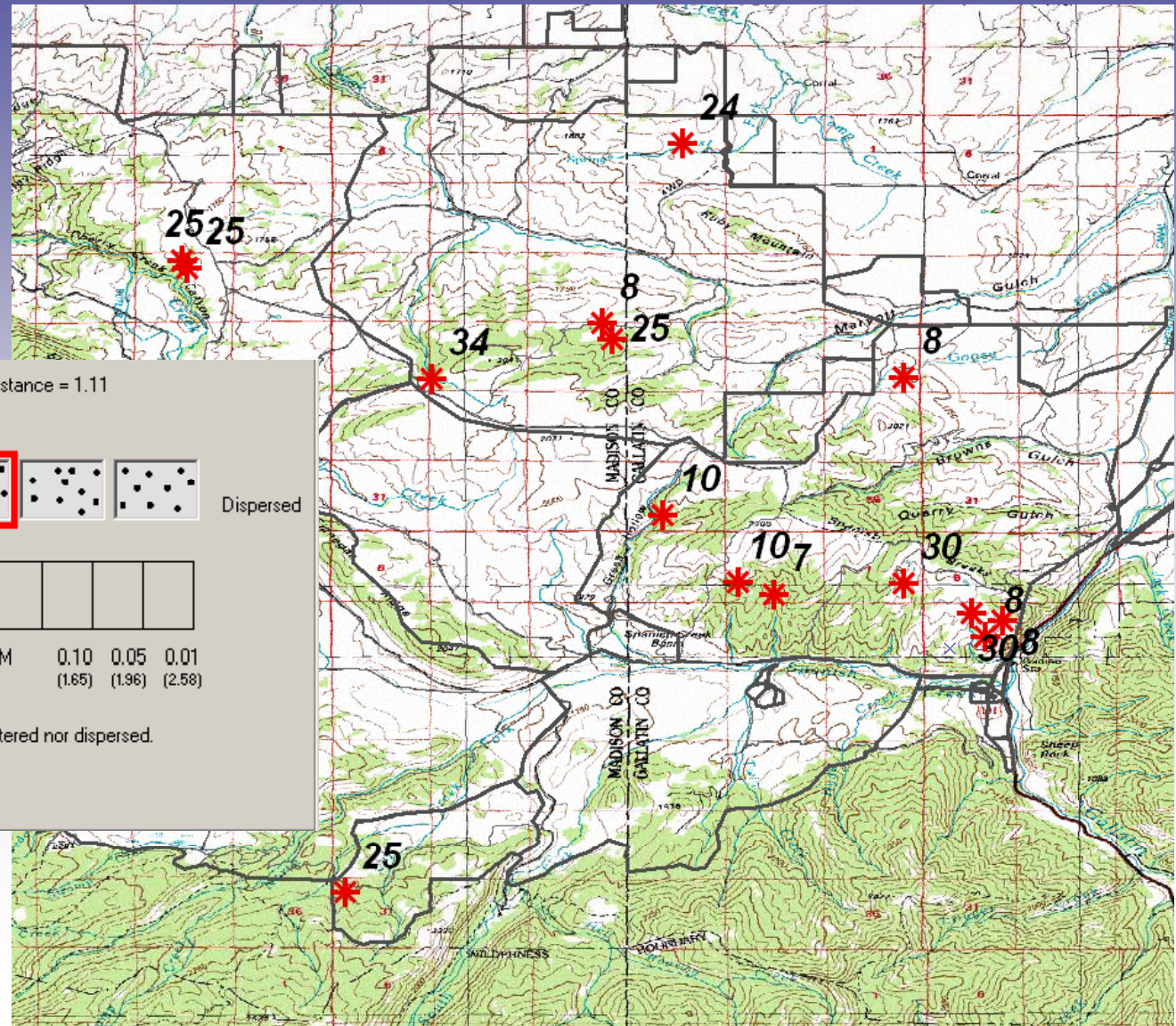
Daily deaths of bison across the pastures



Ripley's K Plots of Bison Cases



Elk spatial-temporal distribution and ANNI



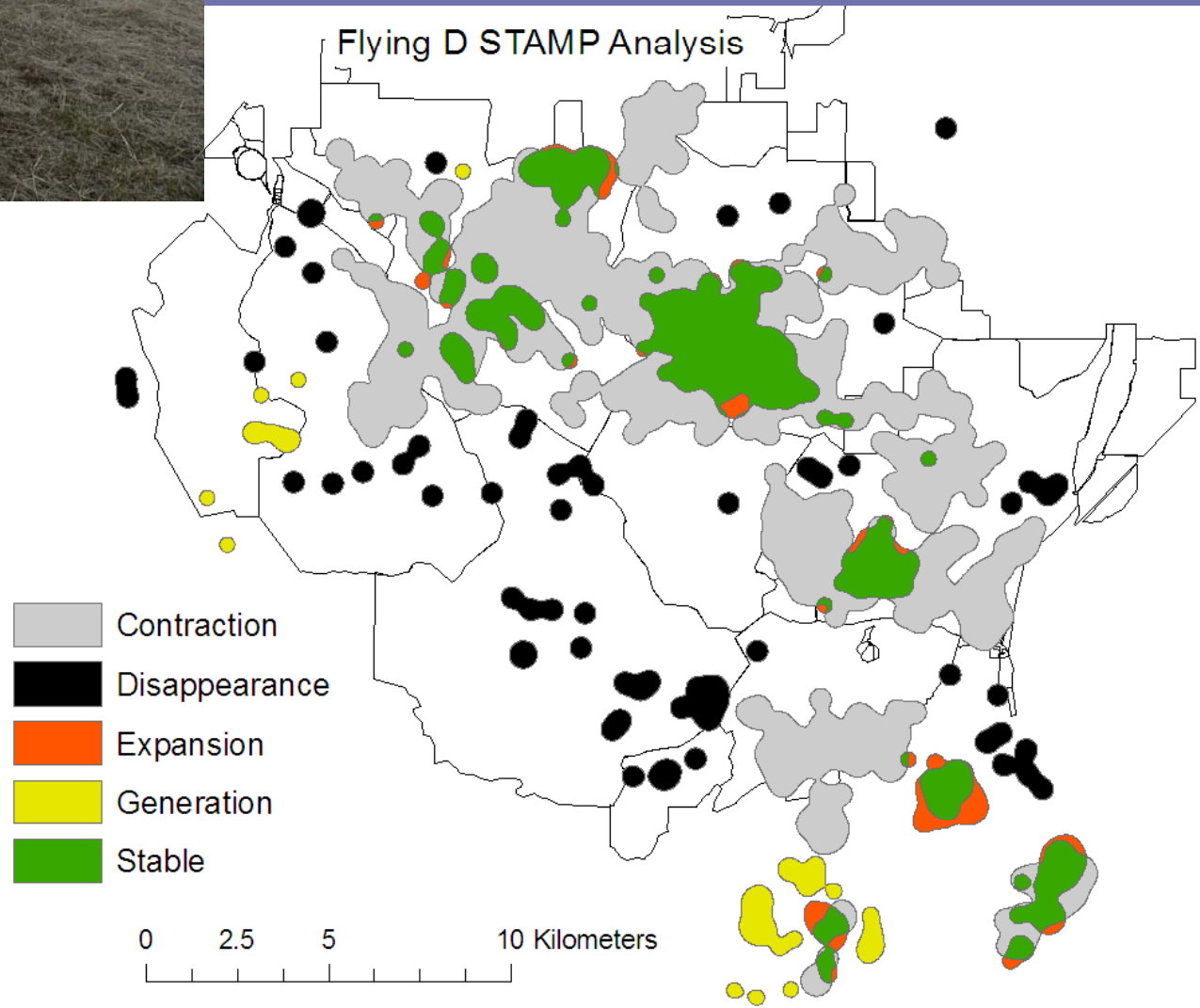
Observed Mean Distance / Expected Mean Distance = 1.11
Z Score = 0.8 standard deviations



Significance Level:	0.01	0.05	0.10	RANDOM	0.10	0.05	0.01
Critical Values:	(-2.58)	(-1.96)	(-1.65)		(1.65)	(1.96)	(2.58)

The pattern is neither clustered nor dispersed.

Elk spatial-temporal changes in home range during anthrax risk period



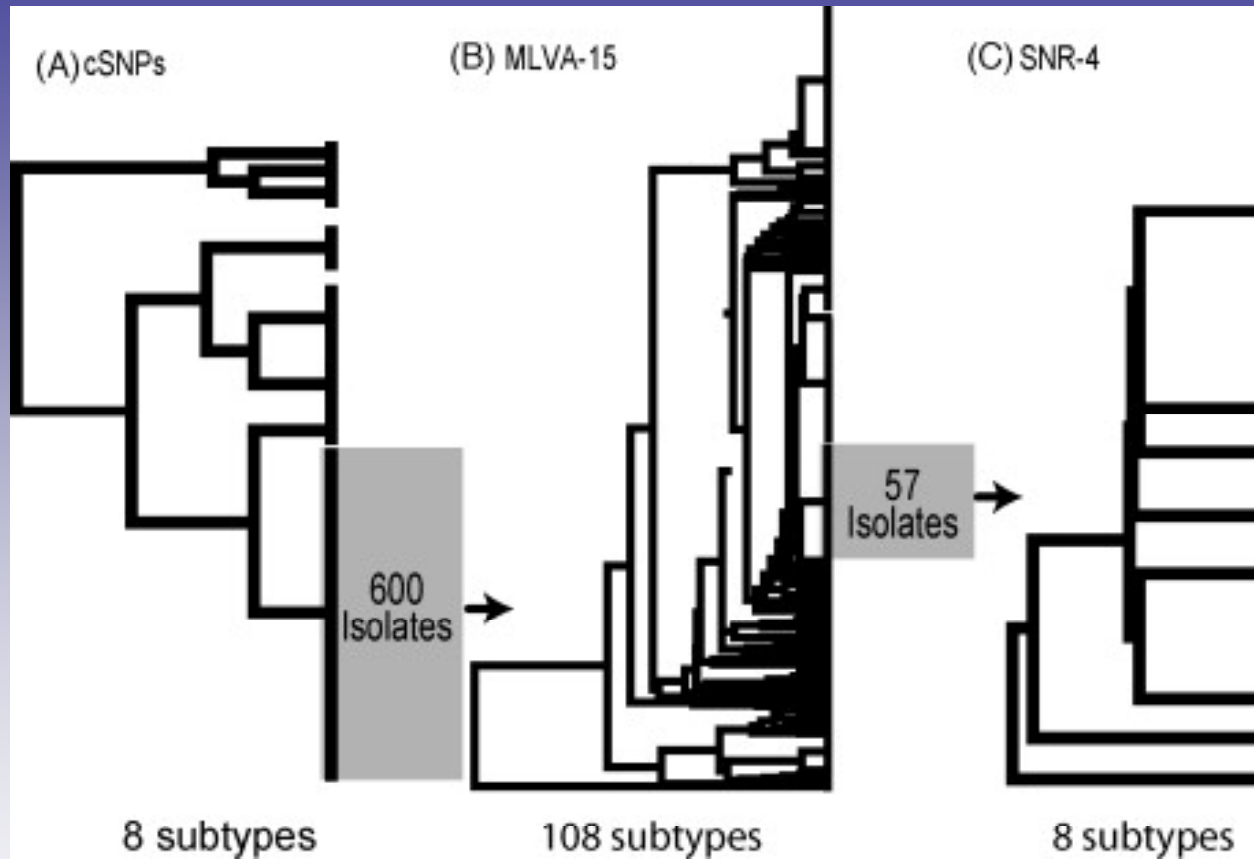
What about genetics? Do they matter?

Qs:

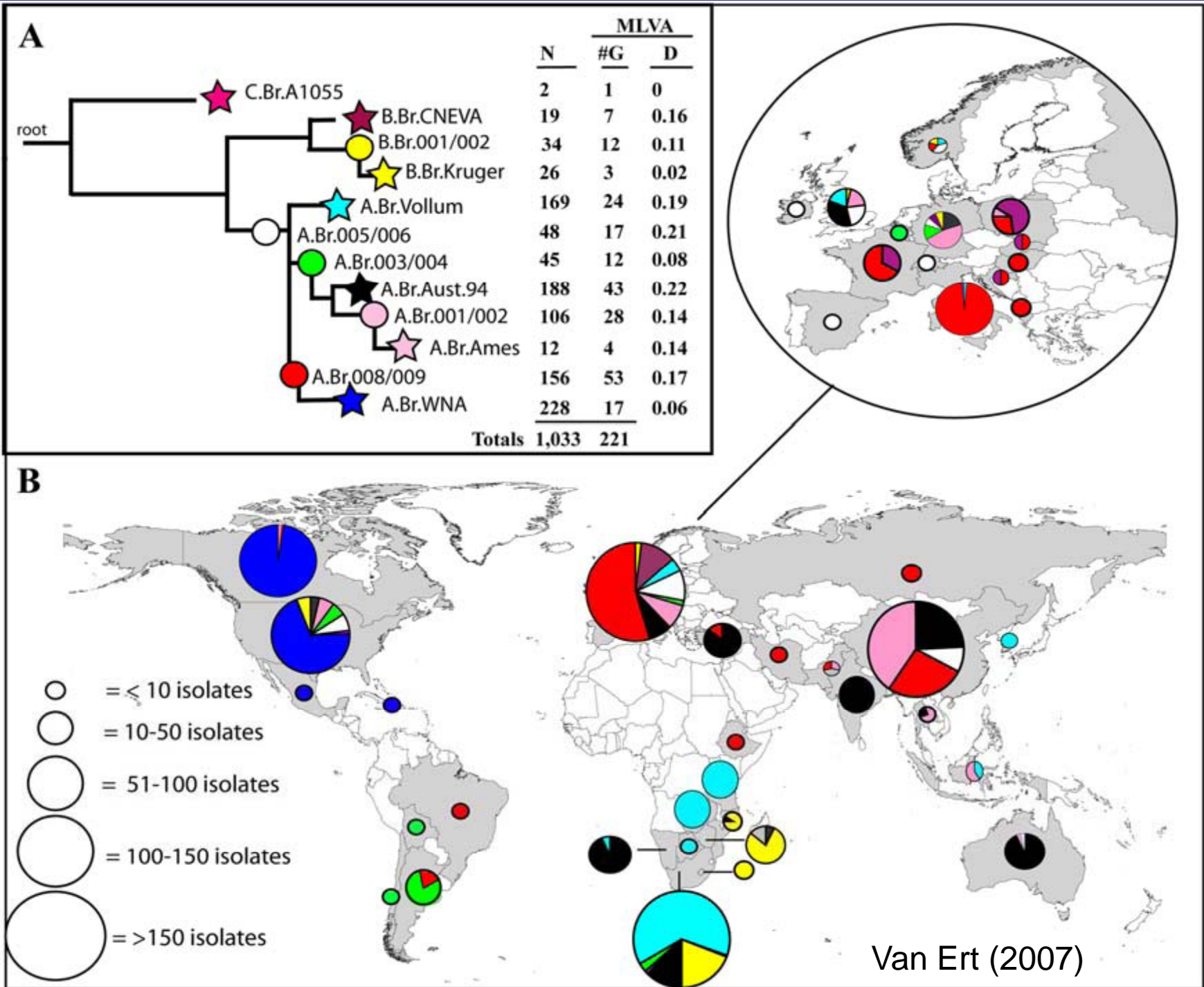
What strains are in the US?

Do the same strains always reoccur? Is there diversity per ranch, per outbreak, per situation?

Which genetic marker? When? Why?



Keim et al. (2004)



Acknowledgements

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