Eco-Epidemiology of Plague in the U.S. and Abroad

Kenneth L. Gage Bacterial Zoonoses Branch DVBID/CDC

The World's Big Three Plague Stars

- Yersinia pestis –
 Extremely virulent gramnegative bacterium
- Rattus rattus Widespread commensal rat; highly susceptible to plague
- Xenopsylla cheopis Unexcelled plague vector











Reported Human Plague Cases By Year-U.S.A., 1947-2005



Western US Counties With Plague Positive Data, 1970 - 2005







* UT,OR,NV,TX,ID,WY,MT,OK,WA











Flea Bite Direct Contact

Human Risks of Plague – U.S.

- Human risks vary by area (SW highest)
- Risks vary over time (partially climate-related)
- Risks highest during epizootics
- Peridomestic environment important risk factor
- Recreational areas also important exposure sites in California and surrounding areas







Areas at risk for Plague



Plague in the Southwest



• High risk areas (pinon-juniper and nearby areas)

• Peridomestic exposures (about 80% of exposures)

• Rock squirrels, other ground squirrels, prairie dogs, wood rats, deer mice and their relatives

• Acquired via:

a. Flea bite (~ 80%)

b. Direct contact with animals (~ 20%)

c. Inhalation (rare– cats with pneumonicplague)



Temporal Changes in Plague Risk

- Strong links between climatic variables and human plague and epizootic activity (Parmenter et al. 1999, Enscore et al 2002, Stapp et al. 2004, Collinge et al. 2005)
- Positive correlations with high winter precipitation
- Negative correlation with excessively high summer temperatures
- Effects most noticeable in areas with strong seasonal climate variations, especially late season winter precipitation peaks (Collinge et al. 2005)
- El Nino events correlated with plague in prairie dogs (Stapp et al. 2004)
- Results appear to have broad applicability but local variations in climatic variables need to be taken into account along with scale of area examine



Reported Human Plague Cases By Year-U.S.A., 1947-2004







Cumulative Reported Human Plague Cases By Month of Onset, US, 1970 - 2005



N = 390

How Could Climate influence Plague Activity?

- Seasonality of transmission
- Survival of fleas
- Ability of fleas to transmit and retain infection
- Rodent host and flea vector population dynamics

Spatial variations in Plague Risk

- Clear habitat
 associations
- Human risks highest near pinyon-juniper
- Cases not common on public land in NM, CO and AZ (mainly private or tribal lands)
- Human behaviors also contribute significantly to risk

Cases within 2.5 km of Selected Habitat Types (Pinyon-Juniper and Ponderosa Pine)

Human Plague Cases within Combined Rocky Mountain Great Basin Open and Closed Coniferous Woodlands

Human Cases and Rocky Mountain Great Basin Open and Closed Coniferous Woodlands (Private and Tribal Lands)

- Human cases within habitats
- Human cases outside habitats

County Boundaries

Private-Tribal Great Basin Open Coniferous Woodland Private-Tribal Great Basin Closed Coniferous Woodlands

Logistic Regression Analyses of Human Plague in New Mexico (Eisen et al. – unpublished)

- Used logistic regression to develop a predictive model of areas at risk for human plague
- Compared the distribution of case points with random points generated using human population density data (Census)
- Used NM GAP Analysis data as habitat layers and USGS land stewardship information

Random selection of control points, weighted by human population density in 1990

(R. Eisen et al. - unpublished)

Random control point selection, weighted by human population density in 1990

Peri-domestic plague risk increases with elevation (up to ~2,100 m), and is higher in close proximity to piñon-juniper ecotones and water

Covariates	k	neg_loglikelihood	AIC	delta AIC	AIC weight	lack of fit p-value	AUC
elevation + elevation ² + dist to 3121_3122 ecotone + dist2 water	5	202.75	415.5	0	100	0.105	0.8
elevation + elevation^2 + dist to 3121_3122 ecotone	4	211.92	431.84	16.34	0	0.04	0.76
elevation + elevation^2	3	218.75	443.5	28	0	0.0035	0.73
elevation + dist to 3121 or 3122	3	225.79	457.58	42.08	0	0.005	0.73
elevation + distkeyhab	3	231.56	469.12	53.62	0	0.0007	0.7
elevation + distance to water	3	238.18	482.36	66.86	0	0.0002	0.73
elevation	2	247.7	499.4	83.9	0	< 0.0001	0.68

Probability of plague

35.1% of the state is classified as high risk plague habitat

19.4% of the state is classified as high risk plague habitat and is privately or tribally owned.

Model evaluation based on buildset and a probability cut-off value > 0.4686

	Actual cl		
Model classification ^a	Human plague case	Control	% correct ^b
Human plague case	172	77	69.08
Control	23	118	83.69
% correct ^c	88.21	60.51	

Logit (P) = $-20.48 + (\text{elevation} * 0.020) - (\text{elevation}^2 * 0.0000047) - (\text{distance to } 3121_{3122} \text{ ecotone } * 0.000055) - (\text{distance to water } * 0.00029)$

^a Probability cut-off value used to classify a 30 m raster as suitable was based on the ROC

optimal cut-off probability ($P \ge 0.4686$).

^b User accuracy (commission error).

^c Producer accuracy (omission error).

Key habitat types

Pinyon-Juniper habitats

Habitat categories derived from the Southwest Regional Gap Analysis (USGS)

Cases and Controls, build set (recreational and peridomestic exposure)

Cases and Controls, evaluation set (recreational and peridomestic exposure)

Candidate Models peridomestic and recreational exposure

Covariates	model #	k	neg_loglikelihood	AIC	delta AIC	AIC weight	lack of fit p-value	AUC
elevation + elevation ² + dist to key habitat	1	4	252.15	512.3	0	0.9	0.3492	0.81
dist to key habitat	2	2	256.39	516.78	4.48	0.1	0.6657	0.81
elevation + elevation ² + dist to pjwoodlands	3	4	261.98	531.96	19.66	0	0.1135	0.8
elevation + elevation^2	4	3	263.79	533.58	21.28	0	0.1132	0.8
dist to pjwoodlands	5	2	285.1	574.2	61.9	0	0.062	0.8

Evaluation matrix, buildset

peridomestic and recreational exposure

	Actual cl		
Model classification ^a	Human plague case	Control	% correct ^b
High risk plague habitat	205	71	74.28
Not high risk	48	182	79.13
% correct ^c	81.03	71.94	

^a Probability cut-off value used to classify a 30 m raster as suitable was based on the ROC optimal cut-off probability ($P \ge 0.6409$). ^b User accuracy (commission error).

^c Producer accuracy (omission error).

Logit (P) = $-8.80 + (\text{elevation} * 0.008971) - (\text{elevation}^2 * 0.000002) - (\text{distance to key habitat} * 0.0003225)$

Evaluation matrix, evaluation set

peridomestic and recreational exposure

	Actual cl		
Model classification ^a	Human plague case	Control	% correct ^b
High risk plague habitat	52	19	73.24
Not high risk	11	44	80.0
% correct ^c	82.54	69.84	

^a Probability cut-off value used to classify a 30 m raster as suitable was based on the ROC optimal cut-off probability ($P \ge 0.6409$). ^b User accuracy (commission error).

^c Producer accuracy (omission error).

Logit (P) = $-8.80 + (\text{elevation} * 0.008971) - (\text{elevation}^2 * 0.000002) - (\text{distance to key habitat} * 0.0003225)$

Probability of plague (recreational and peridomestic exposure) within high risk plague habitat (P>0.6409)

Probability of plague within high risk, nonfederally owned plague habitat (P>0.6690)

Our model identified landscape features associated with human plague cases

- Plague risk increases up to an elevation of approximately 2,300 m and declines as elevation increase thereafter.
- Plague risk is highest in southern Rocky Mtn.
 PJ, Colorado PJ woodlands, Rocky Mtn.
 Ponderosa pine, or So. RM PJ woodlands or savanna habitats and decreases with distance away from these habitats.

Ecological Niche Modeling

- Often referred to as GARP (Genetic Algorithm for Rule set Prediction)
- Uses an iterative, AI-based approach with four types of algorithms (Atomic, logistic regression, bioclimatic envelope and negated bioclimatic envelope) to produce component rules in a broader rule
- Inputs are data layers providing spatial information on various environmental parameters (vegetation types, greenness indices, climatic variables, elevation, slope, aspect, etc.)
- Identifies non-random correlations between the environmental parameters
 and the presence or absence of the species or phenomenon of interest
- Model outputs represent environmental conditions where species should be able to maintain populations
- Check fit of models using test points (20-50% of points withheld at random from original set or use points from new region)
- Used successful to identify sites of likely species invasions, organism distributions and risk areas for disease (Chagas' disease, filovirus infections, etc.)

Ecological Niche Model for Human Plague in Colorado

Model Constructed Using

- Human case exposure site data (point occurrences in Colorado 1980-2004)
- •Elevation, slope, aspect
- NDVI values from (AVHRR)
- Similar environmental parameters used for modeling entire Four Corners region (NM, CO, AZ, UT)
- 50% of cases used as training points

🐂 Desktop Garp - G1.gxl - Finished		
File Datasets Model Results Help		
- Species Data Points		Environmental Layers
Species List: (2 selected)	Upload Data Points	Dataset:
Animals (663)		AVHRR and TOPO
☑ Humans_1980topresent (35)	Options:	Lavers to be used:
	Use 50 % for	▼feb95
	□ At least 20 training points	□ mar95
1		⊠ april95
- Optimization Parameters	- Bast Subset Selection Parameters	□may95
100 Punc	Active	⊠ june95
	Omission massura: 6 Extrinsia	□ july95
01 Convergence limit	C Intrinsic	aug95
1000 Max iterations	Omission threashold: C Hard	□ sepso Ø oct95
Rule types:	© Soft	□ nov95
☑ Atomic	20 % distribution	✓ dec95 🗸 🗸
☑ Range		
✓ Negated Range	Total models under hard 20	How layer will be used:
■ Lugistic Regression (Lugit)	omission threshold	All selected layers All selected layers
All combinations of the selected rules	Max models per spp. 100	C All combinations of the selected layers
(1 rule comb.) (100 total runs)	Commission threshold:	• All combinations of size 1 (1 comb.)
	50 % of distribution	Output
Projection Layers		Mana ast Madalat
Available datasets:	- Add	□ Bitmaps as. © All models
		ASCII Grids C Best subset
for projection:	TOPO 4 states	ARC/INFO Grids
(hesides the	Remove	Output directory:
training dataset)		D:\Plague\G1
,		

Four Corners model made with case exposure points from all 4 states (NM, UT, AZ, and CO)

- Highly significant results (p << 0.00001 or less for each of top 10 models)
- 97.5% of test points included in high risk area
- Approximately 68% of region in risk area

Plague in North America

- U.S. Commonly found in western states
- Mexico
 - Single report of infected Cynomys mexicanus
 - Evidence of *Y. pestis* infection identified in U.S. border counties

Canada –

- Previous reports in ground squirrels in Alberta
- Suspect human case in Alberta (1939)
- Wood rat positives in British Columbia (Morshed pers comm)
- Leighton et al. (2001) report of seropositive cats and dogs in southwestern Saskatchewan and Alberta
- Evidence of *Y. pestis* infection near U.S. border counties

Four Corners GARP model projected to South America and Africa

Comparison of World Plague Map with 4-Corners Model Projected to Africa and South America

PLAGUE PREVENTION

Fuge cito, vade longe, rede tarde

Flee quickly, go far, return slowly

