



NetSurv & Data Viewer

Prototype space-time analysis and
visualization software from TerraSeer

Dunrie Greiling, TerraSeer Inc.



TerraSeer

- Software sales
 - BoundarySeer for boundary detection and analysis
 - ClusterSeer for disease cluster detection
 - SpaceStat for spatial regression modeling
- Training
 - Short courses
- Custom development

www.terraSeer.com



BioMedware

- TerraSeer's R&D partner
 - developed BoundarySeer and ClusterSeer
 - NIH/NCI SBIR funding
- Selection from current projects
 - NetSurv
 - distributed disease surveillance software
 - Cancer Atlas Viewer
 - spatio-temporal visualization of the National Cancer Mortality Atlas
 - DataViewer under construction

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NetSurv Project

- Provide decision support and monitoring tools that will enhance existing disease surveillance systems and support timely analysis, policy formulation, and public health actions

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Surveillance

- Continuous and systematic process of collection, *analysis*, and *interpretation* of information for monitoring health problems
- Ongoing monitoring of temporal and spatial disease trends

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NIH SBIR grants

- Small Business Innovation Research
 - Phase I
 - Evaluate scientific and technical merit and feasibility of an idea (6 months)
 - Phase II
 - Expand on the results and further pursue the development of Phase 1 (2 years)

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NetSurv: Phase I

- Provide CuSum technique (Hutwagner et al 1997) for monitoring temporal trends, providing direct access to a surveillance database and graphical display of results
 - access to single dataset

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CuSum Technique

- Cumulative sum over time, of the differences between observed case counts and a reference/baseline value
- Differences are added together and plotted on graph over time
- Magnifies small, abrupt change which are too small to be visible in conventional graphical plots of a fluctuating series of data

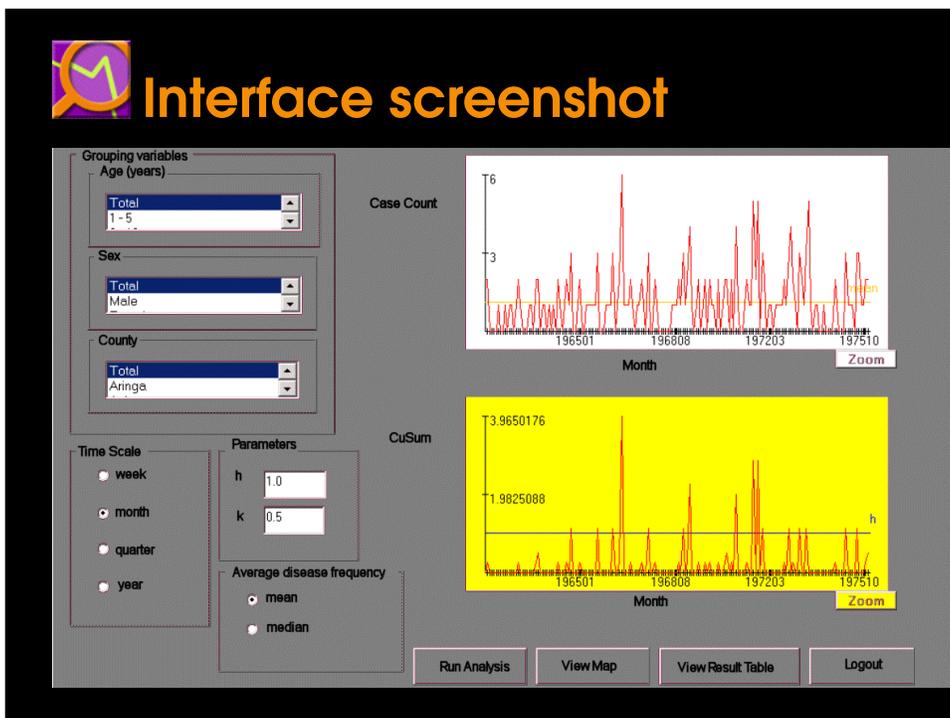
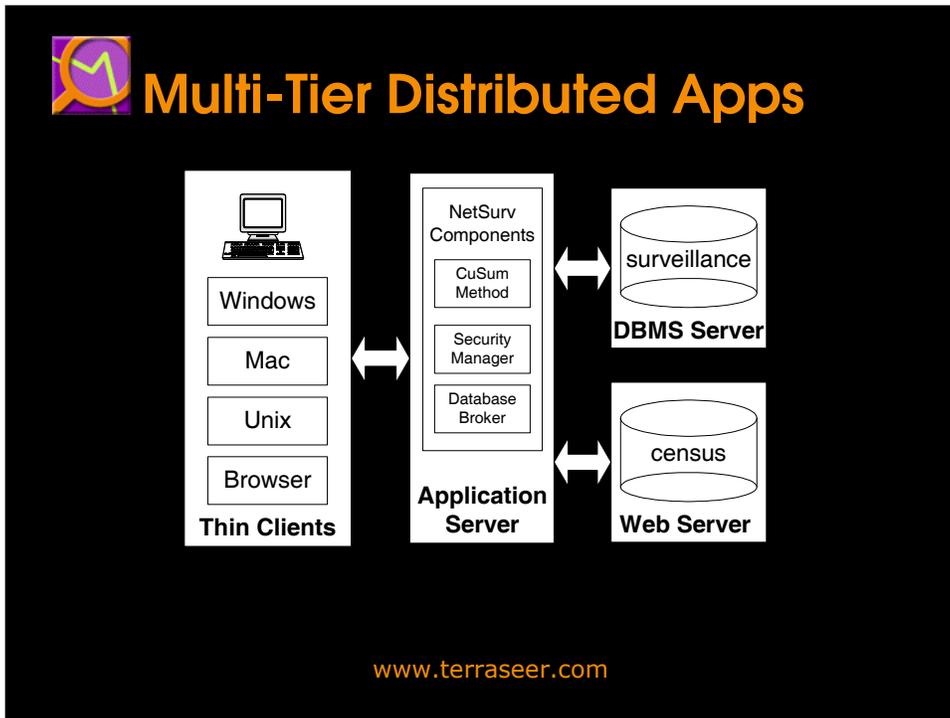
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NetSurv: Phase I

- CuSum technique (Hutwagner et al 1997)
- Distributed system
 - Web browser interface “thin client”

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NetSurv phase I results

- Web-based interface difficult, not user friendly
 - difficult: interface complex, difficult to implement
 - not user friendly: mapping, graphing slow, interface static not dynamic

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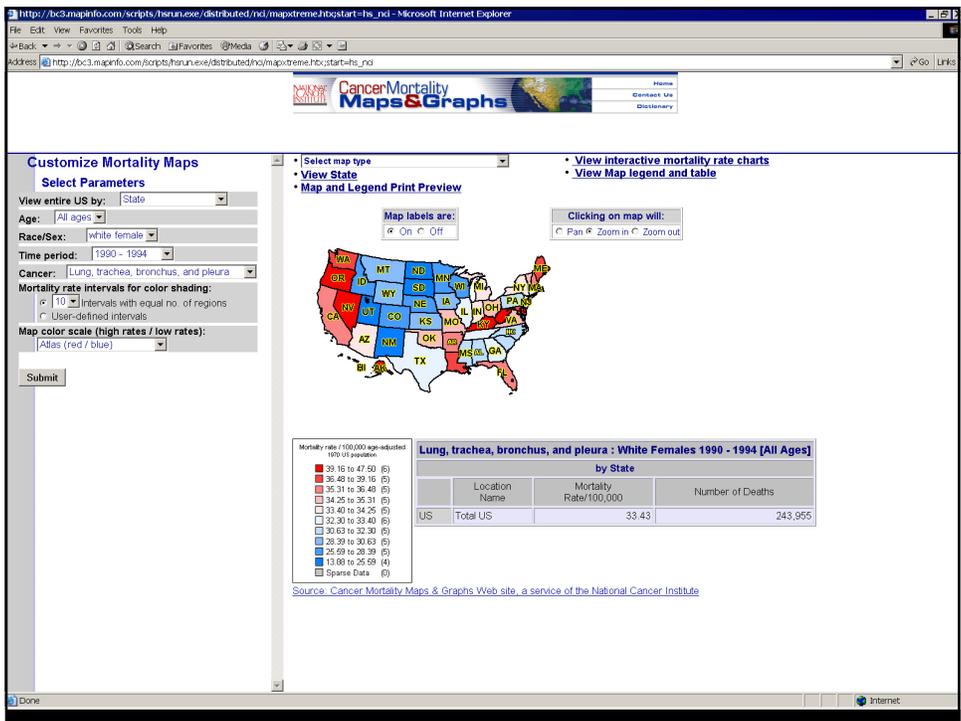
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Motivation for Cancer Atlas Viewer

- Provide real-time visualization of the National Cancer Mortality Atlas Data
- Provide statistics for spatial, temporal, and space-time evaluation of Atlas data
- Explore general STIS specifications with a specific example

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http://bc3.mapinfo.com/scripts/harun.exe/distributed/ncj/mapxtreme.hbgstart=hs_ncj - Microsoft Internet Explorer

Address: http://bc3.mapinfo.com/scripts/harun.exe/distributed/ncj/mapxtreme.hbgstart=hs_ncj

Cancer Mortality Maps & Graphs

Home
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Dictionary

Customize Mortality Maps

Select Parameters

View entire US by:

Age:

Race/Sex:

Time period:

Cancer:

Mortality rate intervals for color shading:
 10 intervals with equal no. of regions
 User-defined intervals

Map color scale (high rates / low rates):

Select map type
 • View State
 • Map and Legend Print Preview

View interactive mortality rate charts
 • View Map legend and table

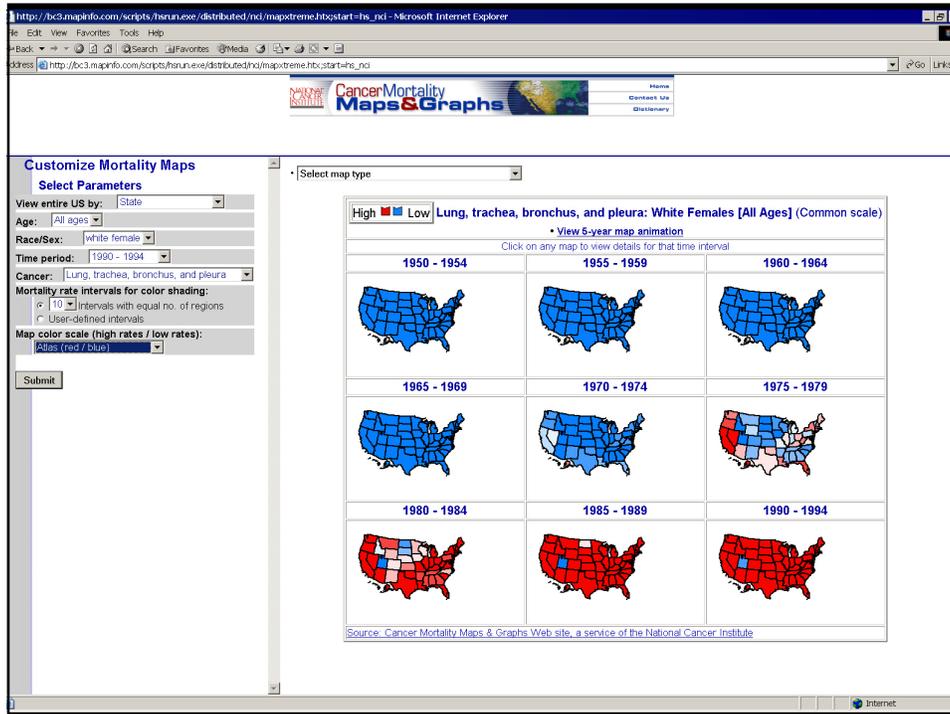
Map labels are:
 On Off

Clicking on map will:
 Pan Zoom in Zoom out

Lung, trachea, bronchus, and pleura : White Females 1990 - 1994 [All Ages]

by State			
	Location Name	Mortality Rate/100,000	Number of Deaths
US	Total US	33.43	243,955

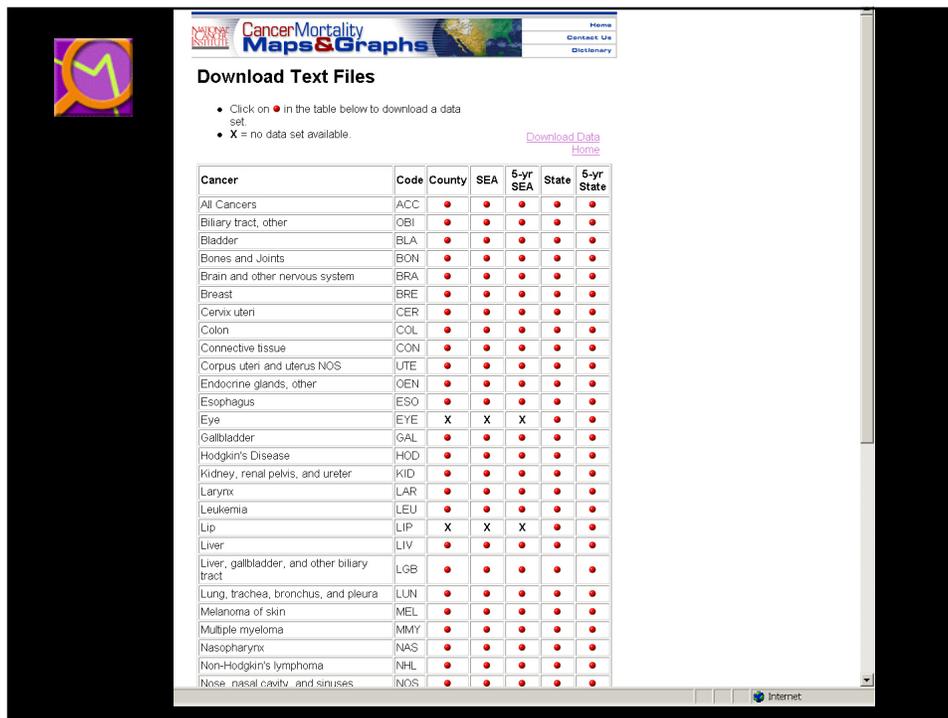
Source: Cancer Mortality Maps & Graphs Web site, a service of the National Cancer Institute



Real Time Interaction

- Avoid the “world wide wait”

The screenshot shows the same website interface as above, but with a loading screen overlay. The 'Submit' button has been clicked, and the page displays 'Map being produced....' and 'Please wait for map creation.' Below this, a message states: 'Map creation may take up to a minute, depending upon your connection speed'. The parameter section on the left remains visible, with the 'Submit' button highlighted.



Cancer Mortality Maps & Graphs

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History

Download Text Files

- Click on ● in the table below to download a data set.
- X = no data set available.

[Download Data](#)
[Home](#)

Cancer	Code	County	SEA	5-yr SEA	State	5-yr State
All Cancers	ACC	●	●	●	●	●
Biliary tract, other	OBI	●	●	●	●	●
Bladder	BLA	●	●	●	●	●
Bones and Joints	BON	●	●	●	●	●
Brain and other nervous system	BRA	●	●	●	●	●
Breast	BRE	●	●	●	●	●
Cervix uteri	CER	●	●	●	●	●
Colon	COL	●	●	●	●	●
Connective tissue	CON	●	●	●	●	●
Corpus uteri and uterus NOS	UTE	●	●	●	●	●
Endocrine glands, other	OEN	●	●	●	●	●
Esophagus	ESO	●	●	●	●	●
Eye	EYE	X	X	X	●	●
Gallbladder	GAL	●	●	●	●	●
Hodgkin's Disease	HOD	●	●	●	●	●
Kidney, renal pelvis, and ureter	KID	●	●	●	●	●
Larynx	LAR	●	●	●	●	●
Leukemia	LEU	●	●	●	●	●
Lip	LIP	X	X	X	●	●
Liver	LIV	●	●	●	●	●
Liver, gallbladder, and other biliary tract	LGB	●	●	●	●	●
Lung, trachea, bronchus, and pleura	LUN	●	●	●	●	●
Melanoma of skin	MEL	●	●	●	●	●
Multiple myeloma	MMY	●	●	●	●	●
Nasopharynx	NAS	●	●	●	●	●
Non-Hodgkin's lymphoma	NHL	●	●	●	●	●
Nose, nasal cavity, and sinuses	NOS	●	●	●	●	●



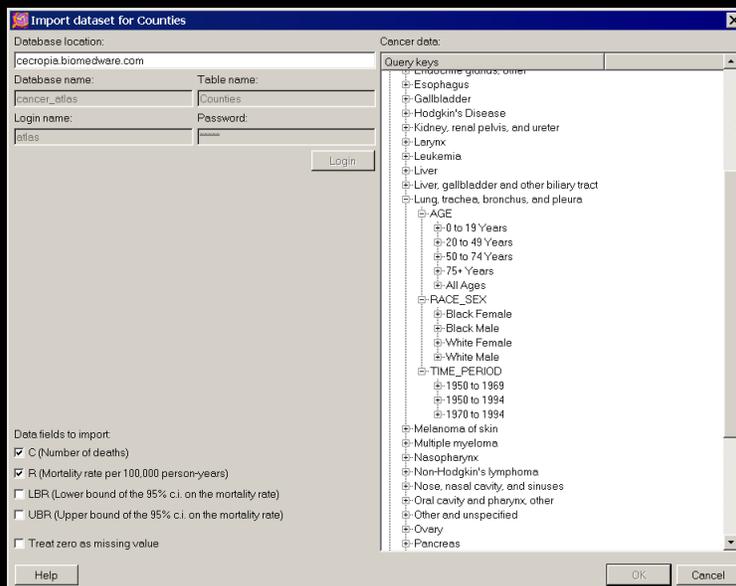
Real Time Interaction

- Provide more flexible access to the data.
- Concurrency issues

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Downloading Data



Real Time Interaction

- Provide linked views that you can brush for interactive data exploration
 - Map
 - Scatterplot
 - Box plot
 - Histogram
 - Table

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Space-Time Viz

- Slideshow
 - Group of maps with a common legend

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Provide Statistics

- Standardization
 - Z-score
- LISA
 - Univariate
 - spatial contagion
 - Bivariate
 - space-time contagion
 - Cluster persistence

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Moran's I

- Global statistic – 1 value for entire dataset
- Spatially weighted correlation coefficient
- Range ~ (-1, 1)
- Moran, P.A.P. 1950. Notes on continuous stochastic phenomena. Biometrika 37: 17-23.

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Calculation of LISA's

1. Standardize data as z-score
$$z_i = (x_i - \mu_x) / \text{var}(x)^{1/2}$$
2. Calculate LISA statistics (Anselin, 1995)
 - local statistic, 1 value for every location
$$I_i = z_i \sum w_{ij} z_j$$
3. Evaluate significance of LISA statistics via Monte Carlo randomization

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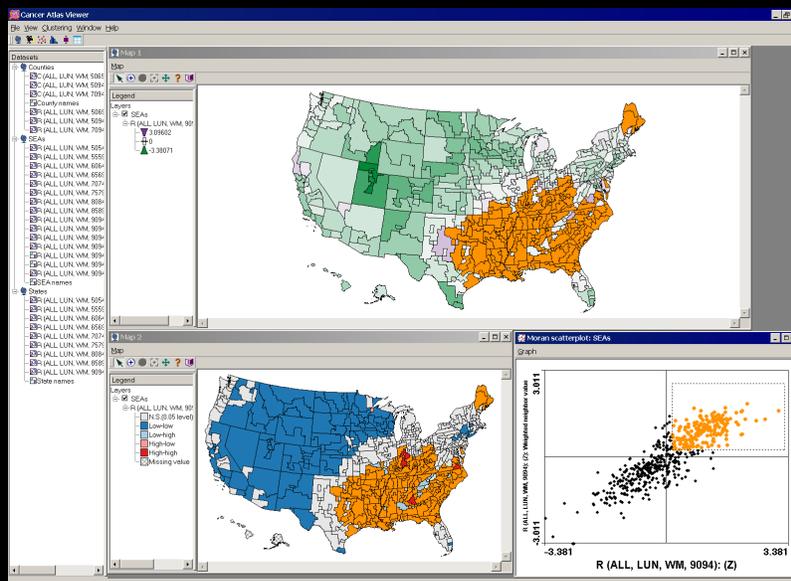
The Moran Scatter Plot

- Graphs the values (z_i) of each area versus the average of its neighbors
 - $\sum w_{ij} z_j$
- Has four quadrants that display high-high and low-low clusters, and high-low and low-high outliers

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Local Clustering (LISA)



Mask Sparse Data

- Count < 6

The screenshot shows a GIS application window titled 'Map 2'. On the left, a context menu is open with the following options: 'Add to map', 'Remove dataset', 'Create standardized dataset', 'Mask sparse values' (highlighted with a mouse cursor), and 'Properties'. The map area displays a grid of small polygons, with some shaded in light blue and others in a hatched pattern. A legend in the top right corner shows a scale from 1.9635 to 17.4149 and a 'Missing value' category.

Analyze Masked Datasets

The screenshot shows the 'Cancer Atlas Viewer' application. The left sidebar lists various datasets, including 'COUNTIES', 'SEAS', and 'RATES'. The main window is divided into two map panes. The top pane, titled 'Raw Rates', shows a map of a region with a legend for 'COUNTIES' and 'RATES (0.05 level)' with categories: Low/low (blue), High/low (red), High/high (red), and Missing value (hatched). The bottom pane, titled 'Masked Rates', shows the same region with a legend for 'RATES (0.05 level)' and 'R (ALL PAN WF 7094) (masked) (2)' with categories: Low/low (blue), High/low (red), High/high (red), and Missing value (hatched). The 'Masked Rates' map shows significantly more hatched areas, indicating where data has been masked.



Provide Statistics

- Standardization
 - Z-score
- LISA
 - Univariate
 - spatial contagion
 - Bivariate
 - space-time contagion
 - Cluster persistence

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Long Term

- Include other statistics
 - ClusterSeer
 - temporal, spatial, spatio-temp, & surveillance methods
 - BoundarySeer
 - edge detection (wombling), classification (fuzzy, spatially-constrained)
 - Other
 - change detection
- Provide open interface for user-scripted methods
 - Python

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Long Term

- Open to other data (more general product)
 - Currently - Adding visualization of points moving through time
 - modeling individuals' movements
 - Interested in applying to infectious disease spread
 - humans
 - plant pathogen
 - amphibians

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Back to NetSurv

- Replace static web-based interface with more interactive Atlas/Data Viewer like interface

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NetSurv phase II

- Retain attention to data concurrency
 - web access to download data
 - check for updates
- Retain attention to permissions/privacy concerns
- Pull down data and then do analysis on local machine
 - avoids world-wide-wait for mapping, graphing

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Long term plans for NetSurv

- Atlas-like interface
- Custom statistics for surveillance applications
 - User-programmed in Python
- Interact with existing web data repositories
 - DataWeb
 - Census
 - Geographic data
 - plus provide room for custom/non-public data repositories

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- The Cancer Atlas software was funded by a grant from the National Cancer Institute to BioMedware, Inc.

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