



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

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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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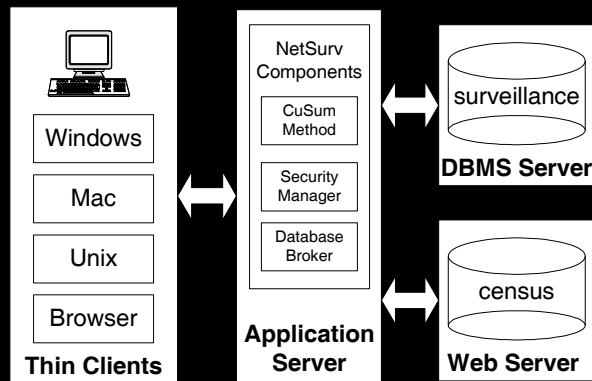
NetSurv: Phase I

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

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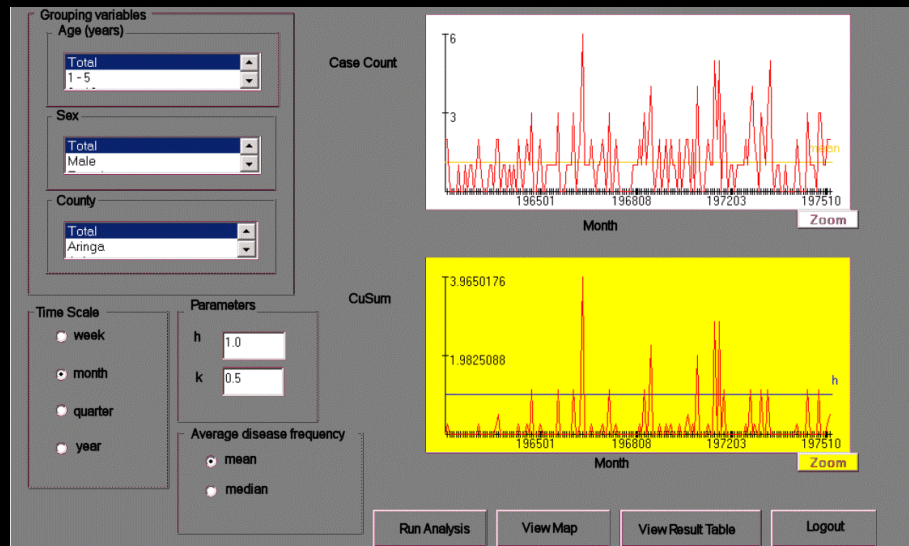
Multi-Tier Distributed Apps



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Interface screenshot





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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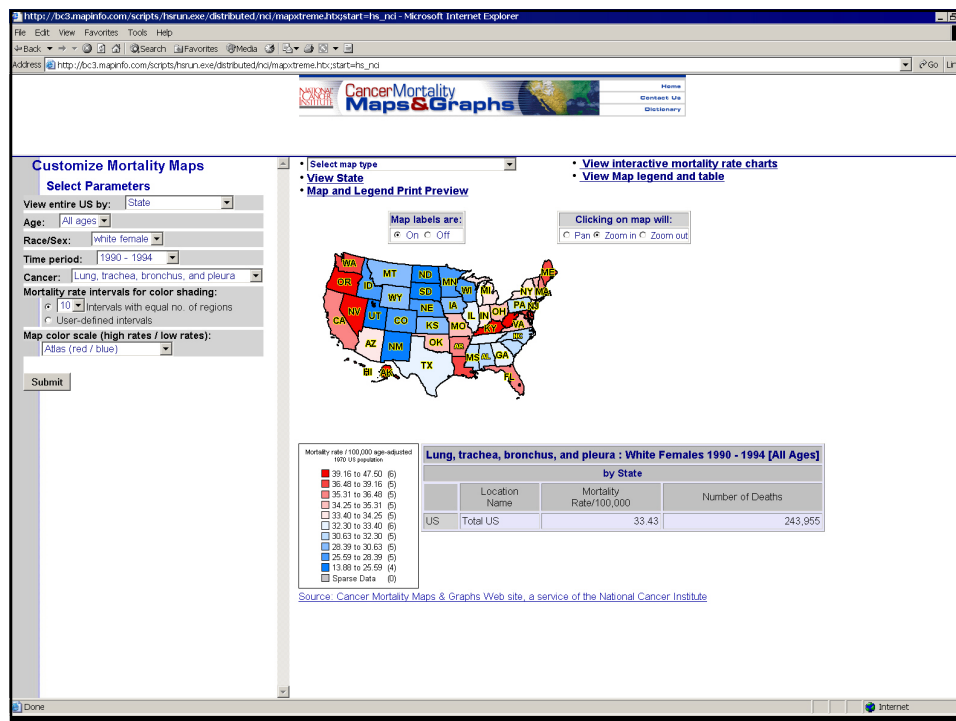
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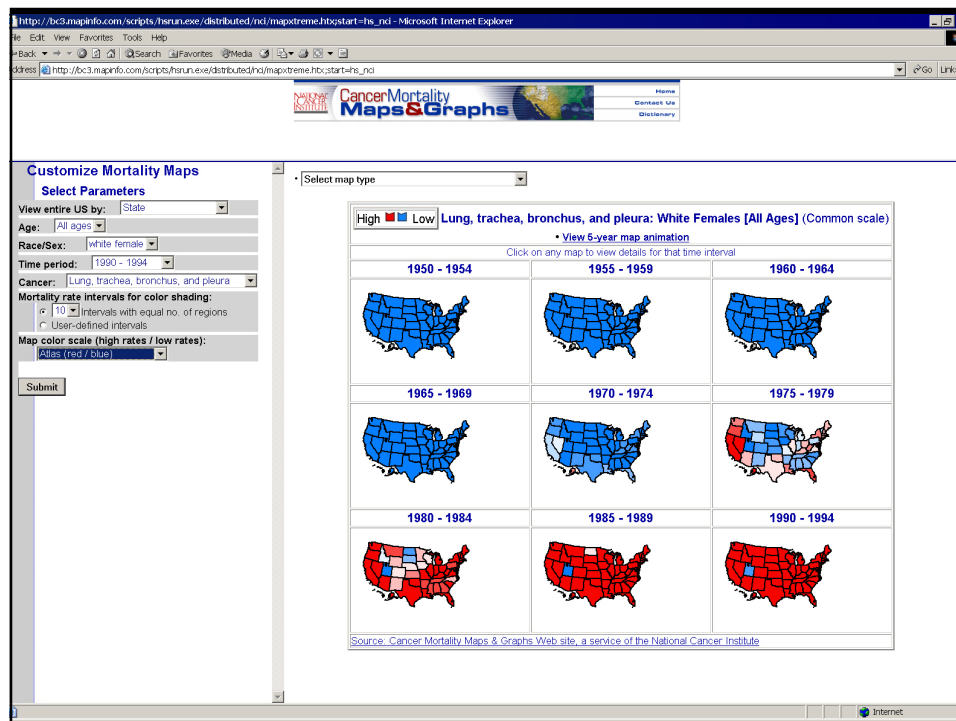


Motivation for Cancer Atlas Viewer

- 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

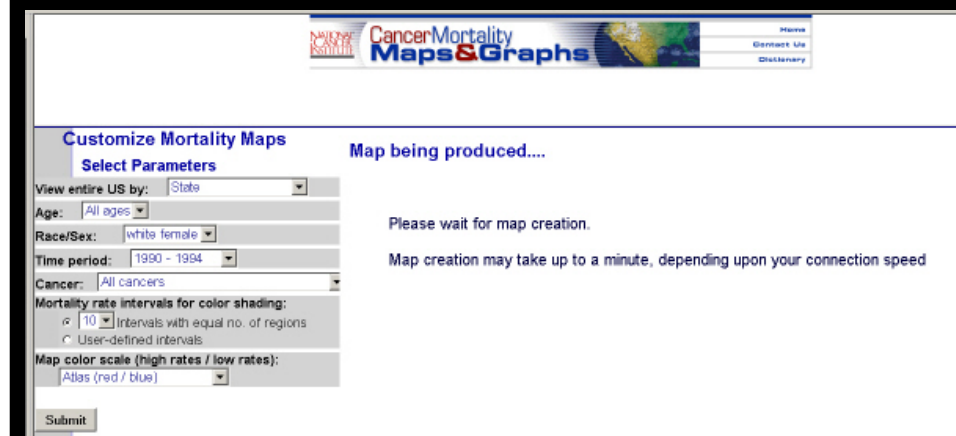
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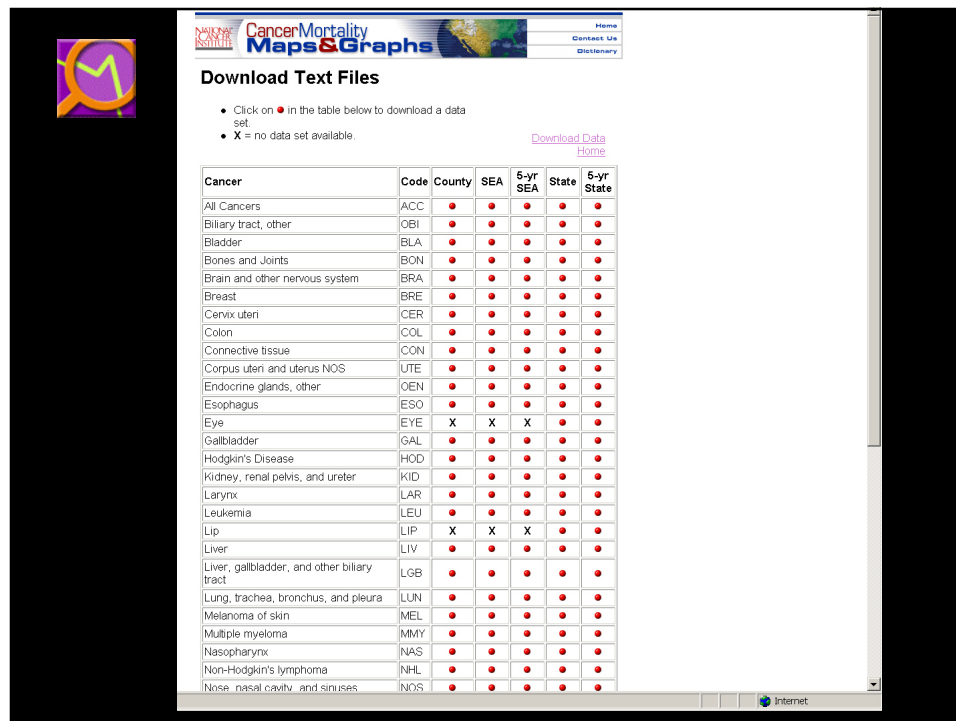




Real Time Interaction

- Avoid the “world wide wait”





Cancer Mortality Maps & Graphs

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

Import dataset for Counties

Database location: cecropia.biomedware.com

Database name: cancer_atlas Table name: Counties

Login name: atlas Password: password

Login

Cancer data:

Query keys

- Esophageal glands, other
- Esophagus
- Gallbladder
- Hodgkin's Disease
- Kidney, renal pelvis, and ureter
- Larynx
- Leukemia
- Liver
- Liver, gallbladder and other biliary tract
- Lung, trachea, bronchus, and pleura
- AGE
 - 0 to 19 Years
 - 20 to 49 Years
 - 50 to 74 Years
 - 75+ Years
 - All Ages
- RACE_SEX
 - Black Female
 - Black Male
 - White Female
 - White Male
- TIME_PERIOD
 - 1950 to 1969
 - 1950 to 1994
 - 1970 to 1994
- Melanoma of skin
- Multiple myeloma
- Nasopharynx
- Non-Hodgkin's lymphoma
- Nose, nasal cavity, and sinuses
- Oral cavity and pharynx, other
- Other and unspecified
- Ovary
- Pancreas

Data fields to import:

- ☒ C (Number of deaths)
- ☒ R (Mortality rate per 100,000 person-years)
- ☐ LBR (Lower bound of the 95% c.i. on the mortality rate)
- ☐ UBR (Upper bound of the 95% c.i. on the mortality rate)
- ☐ Treat zero as missing value

Help OK Cancel



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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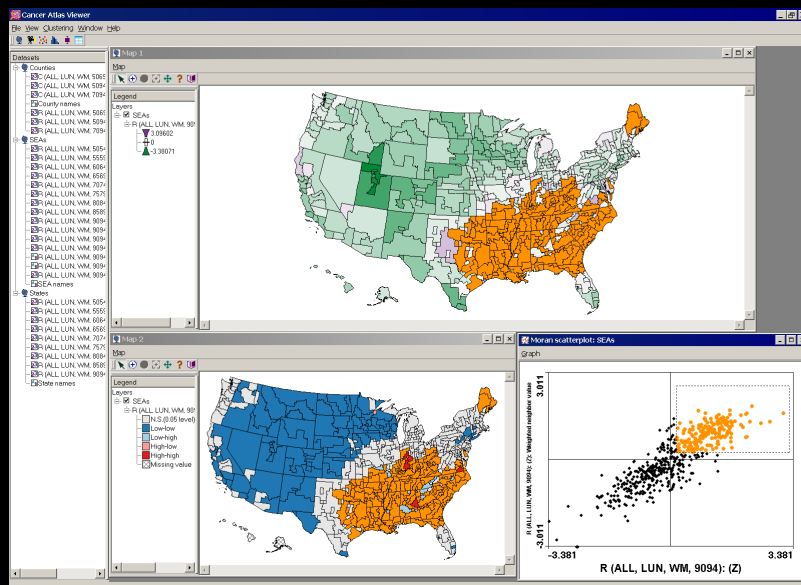
The Moran Scatter Plot

- 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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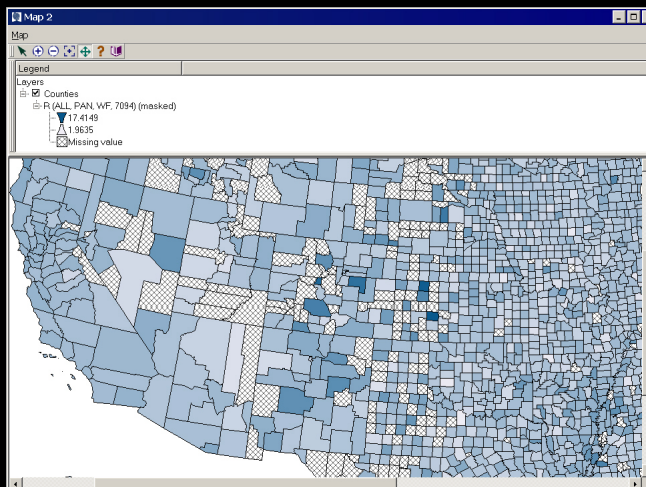
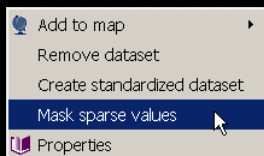
Local Clustering (LISA)



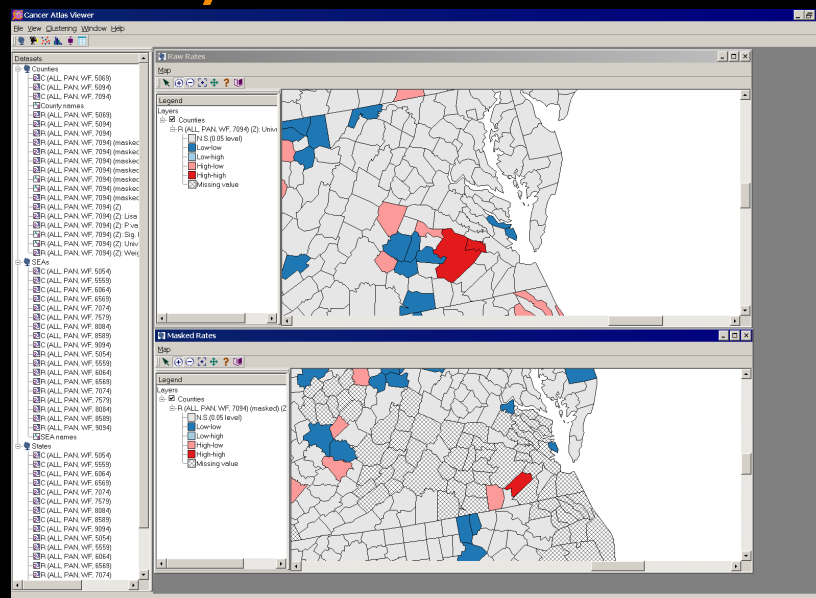


Mask Sparse Data

■ Count < 6



Analyze Masked Datasets





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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