

A Bayesian Method for Disease Cluster Detection

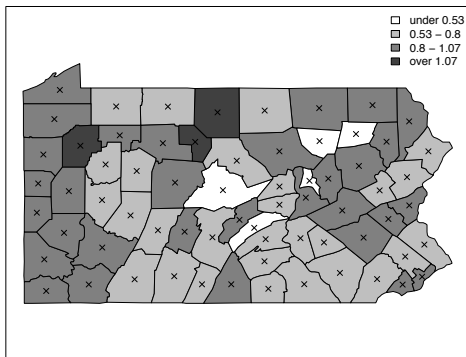
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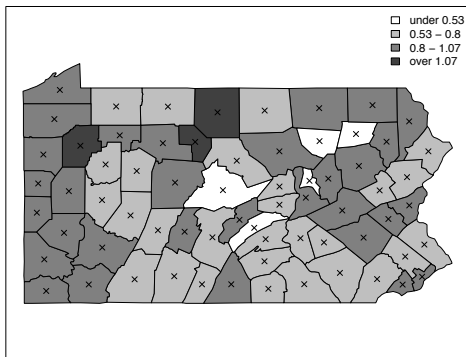
Scenario

Lung Cancer Incidence (per 1000) in Pennsylvania



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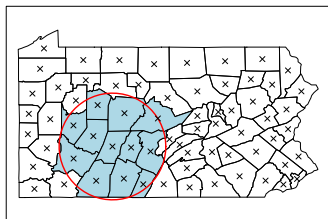


Disease counts Y_i are modeled via $\text{Poisson}(E_i \times \theta_i)$ where

- ▶ $E_i =$ *expected numbers of disease*
- ▶ $\theta_i =$ *relative risk of disease*. $\theta_i > 1$ suggests high risk.
- ▶ Point estimate $\hat{\theta}_i$ is the *standardized mortality ratio* $\frac{Y_i}{E_i}$

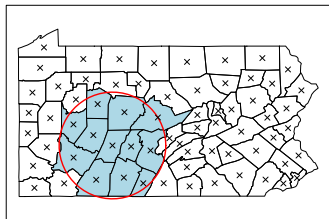
Geographic Units

We define a set of **single zones** z by placing circles on the map. Each z is a **candidate cluster**.

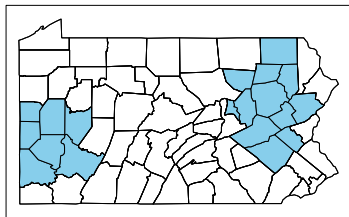


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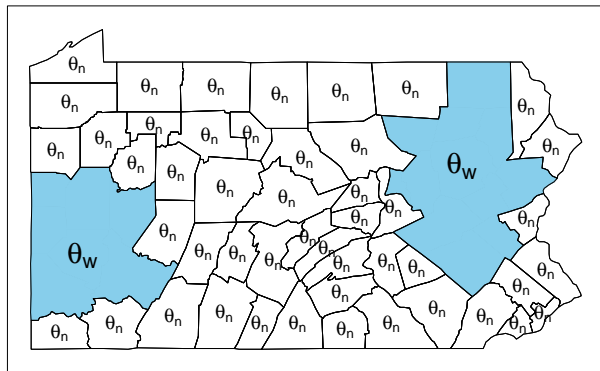


We define a set of **configurations** c : combinations of single zones. Each c is a **candidate explanatory model** for \vec{y}



Explanatory model for \vec{y}

Based on a particular c , we derive an explanatory model for \vec{y} :



Posterior Probability of a Configuration

The (discrete) parameter of interest is the configuration c with posterior probability:

$$\pi(c|\vec{y}) = \frac{\rho(\vec{y}|c)\pi(c)}{\sum_c \rho(\vec{y}|c)\pi(c)}$$

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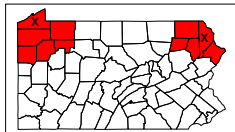
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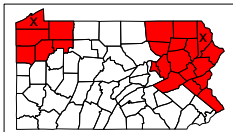
Solution: Use MCMC to estimate $\pi(c|\vec{y})$. Specifically a Metropolis-Hastings algorithm with proposal function Q .

Five Types of Moves in Proposal Function Q

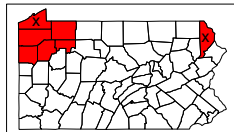
Current Configuration



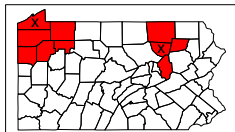
Move 1: Growth



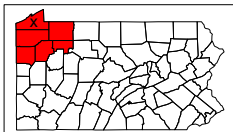
Move 2: Trim



Move 3: Replacement



Move 4: Death



Move 5: Birth

