

Outline

- BSG
- Basics
- Estimation
- Identification
- Covariates
- NIH grant

Human or Cylon? Group testing on the Battlestar Galactica

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Battlestar Galactica

- Statistics and Battlestar Galactica
- The story so far...
 - [Video](#)

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Battlestar Galactica

- Statistics and Battlestar Galactica
- The story so far...
 - Video
- Cylons
 - Centurion
 - Humanoid form (new)
- How can you distinguish a human from a Cylon?

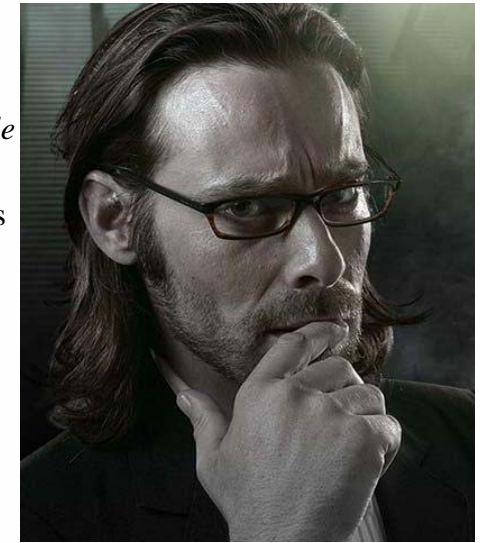


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Battlestar Galactica

- Dr. Gaius Baltar
 - Asked to develop a Cylon detector
 - Season 1's *Bastille Day* episode
 - # of Cylons in fleet is expected to be small
 - 47,905 individuals to test!



Battlestar Galactica

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- Dr. Gaius Baltar (continued)
 - Season 1's *Tigh me up and Tigh me down*



Tigh Me Up, Tigh Me Down

A frustrated Gaius Baltar contemplates the immense workload ahead of him as he prepares to test key fleet personnel with his Cylon-detector.

- [Video](#)

Battlestar Galactica

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 - Season 1's *Tigh me up and Tigh me down*



Tigh Me Up, Tigh Me Down

A frustrated Gaius Baltar contemplates the immense workload ahead of him as he prepares to test key fleet personnel with his Cylon-detector.

- Video
- $(47,905 \text{ blood tests}) * (11 \text{ hours each}) = 21,956 \text{ days}$

Battlestar Galactica

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- Individual testing



- Problems:
 - Time
 - Limited resources

Battlestar Galactica

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- Group testing



- If a GROUP is negative, then all 4 individuals are not Cylons
- If the GROUP is positive, then at least ONE of the 4 individuals is a Cylon
 - “Retesting” can be done to determine who is a Cylon

Battlestar Galactica

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- Group testing (continued)
 - Time savings
 - Save resources
 - Strategy works well when prevalence of a “trait” is small
 - If prevalence is large, all groups may test positive

Other examples

Outline

- BSG
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- Screening blood donations
 - American Red Cross uses groups of size 16
 - HIV, Hepatitis B, Hepatitis C, ...
 - Screen about 6 million a year
 - Source: Roger Dodd, Executive Director of Blood Services R & D at ARC
 - See Dodd et al. (*Transfusion*, 2002)
- Drug discovery experiments
 - Screen hundreds of thousands of chemical compounds to look for potentially good ones
 - Remlinger et al. (*Technometrics*, 2006)

Other examples

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- Multiple vector transfer design experiments
 - Estimate probability an insect vector transfers a pathogen to a plant
 - Swallow (*Phytopathology*, 1985, 1987)
- Veterinary
 - Bovine viral diarrhea in cattle (Peck, *Beef*, 2006)
 - Avian pneumovirus (APV) in turkeys (Maherchandani et al., *J. Veterinary Diagnostic Investigation*, 2004)
- Public health studies
 - Prevalence of HCV (Liu et al., *Transfusion*, 1997)
 - Prevalence of HIV (Verstraeten et al., *Trop. Med. & International Health*, 2000)



Notation

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- Individual responses
 - $Y_{ik} = 1$ if the i^{th} item in the k^{th} group has the “trait” (positive) and $Y_{ik} = 0$ otherwise (negative) for $i=1, \dots, I$ and $k=1, \dots, K$
 - Y_{ik} are independent Bernoulli(p) random variables
 - $p = P(Y_{ik} = 1)$
 - Homogenous population
 - p can be thought of as the “individual probability” or “prevalence in a population”
 - Y_{ik} 's are not directly observed (at least initially)

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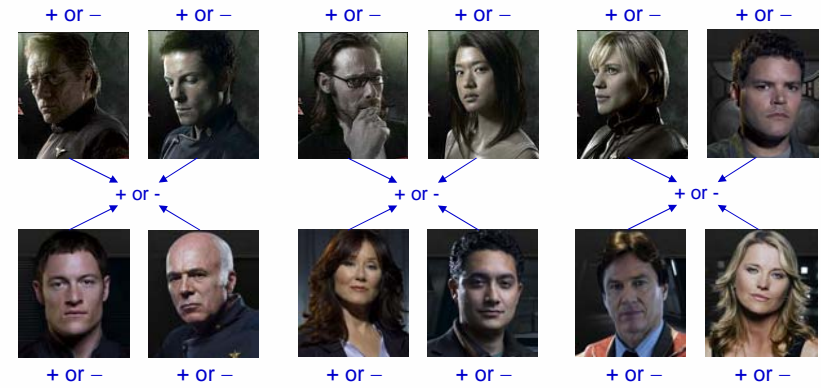
- Group responses
 - $Z_k = 1$ denotes a positive response
 - $Z_k = 0$ denotes a negative response for the k^{th} group
 - Z_k are independent Bernoulli(θ) random variables
 - $\theta = P(Z_k = 1)$
- Individual and group response relationship
 - $Z_k = 1$ if and only if $\sum_{i=1}^I Y_{ik} > 0$
 - $Z_k = 0$ if and only if $\sum_{i=1}^I Y_{ik} = 0$

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- Example random variables

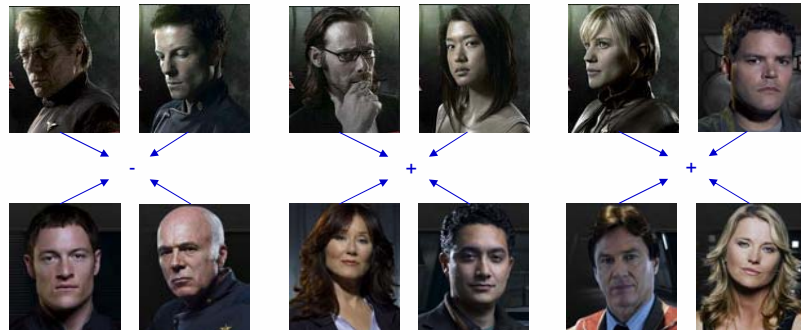


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- Example observed values

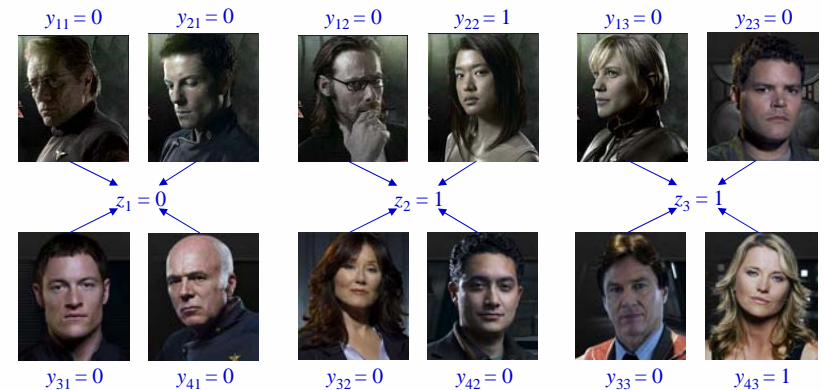


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- Example observed values



Purpose

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- Prevalence of a trait in a population (estimation problem)
- Which items are positive (identification problem)

Estimate p

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- How can we estimate $p = P(Y_{ik} = 1)$?
 - We observe information about the groups, not individuals!
 - $\theta = 1 - P(Y_{ik} = 0, \forall i) = 1 - (1 - p)^I$
 - Then $p = 1 - (1 - \theta)^{1/I}$
 - MLE for p : $\hat{p} = 1 - \left(1 - \sum_{k=1}^K z_k / K\right)^{1/I}$

- Unequal group sizes
 - Likelihood function

$$L(p) = \prod_{k=1}^K \theta_k^{z_k} (1 - \theta_k)^{I - z_k} = \prod_{k=1}^K [1 - (1 - p)^k]^{z_k} (1 - p)^{k(I - z_k)}$$

where

θ_k = positive probability for group k

I_k = size of group k

Testing error

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- What if there is testing error?
 - Can incorporate sensitivity (η) and specificity (δ)
 - $\theta_k = \eta + (1 - \delta - \eta)(1 - p)^{I_k}$

Identification

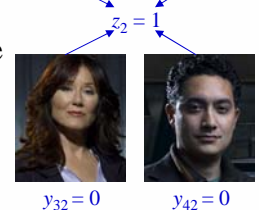
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- Dorfman (*Annals of Mathematical Statistics*, 1943)
 - Retest all items in a positive group
 - Often credited for the very first use of group testing



- Sterrett (*Annals of Mathematical Statistics*, 1957)
 - Individually retest until first positive is found
 - Re-group remaining items
 - If group is negative, STOP
 - If group is positive, repeat



- Expected number retests is smaller than Dorfman
- Gupta and Malina (*Statistics in Medicine*, 1999) provides a summary

Infertility Prevention Program

Outline

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- U.S. national program funded by Centers for Disease Control and Prevention
 - Assess and reduce prevalence of chlamydia and gonorrhea
- Nebraska
 - Swab or urine specimens are sent to the Nebraska Public Health Laboratory at U. of Nebraska Medical Center
 - NATs
 - About 30,000 **individual** tests done per year
- Group testing!

Infertility Prevention Program

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- Lindan et al. (*J. Clinical Microbiology*, 2005)
 - Estimates that 12% of the laboratories in the U.S. are already using group testing
 - Group testing has allowed “laboratories to achieve a significant increase in specimen loads.”
- Quarter #1 of 2006, chlamydia testing
 - Urine specimens – 1,384 total
 - Ignore sensitivity and specificity here
 - Individual data: $\hat{p} = 111/1,384 = 0.0802$
 - Group testing:
 - Randomly put known individual responses into groups of size $I = 2$
 - $\hat{p} = 1 - (1 - \sum_{k=1}^K z_k / K)^{1/I} = 1 - (1 - 105 / 692)^{1/2} = 0.0790$

Infertility Prevention Program

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- Quarter #1 of 2006 (continued)
 - Individual data: $111/1,384 = 0.0802$
 - Group testing:

	Group size				
	1	2	3	5	10
\hat{p}	0.0802	0.0790	0.0791	0.0776	0.0843
Dorfman		902	765	737	949
Sterrett		902	728	653	744

- Approximate cost per test
 - \$16 for urine
 - \$11 for swab

Heterogonous populations

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- Individual responses
 - Y_{ik} are independent Bernoulli(p_{ik}) random variables
 - $p_{ik} = P(Y_{ik} = 1)$ for item i in group k
- Group responses
 - Z_k are independent Bernoulli(θ_k) random variables
 - $\theta_k = P(Z_k = 1)$ for group k
- Covariates
 - $x_{ik1}, x_{ik2}, \dots, x_{ikp}$ for the i^{th} item in the k^{th} group
 - Incorporate factors which influence trait status
 - Not really done until recently in group testing!

Kenyan pregnant women study

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- Part of the data from Vansteelandt et al. (*Biometrics*, 2000)

Age	Marital Status	Education level	Parity	Syphilis	Hepatitis B	
						} $z_1 = 1$

Heterogeneous populations

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- Model
 - $\text{logit}(p_{ik}) = \beta_0 + \beta_1 x_{ik1} + \dots + \beta_p x_{ikp}$
- Estimation of $\beta_0, \beta_1, \beta_2, \dots, \beta_p$
 - Note that Y_{ik} are not directly observed
 - Vansteelandt et al. (*Biometrics*, 2000)
 - Likelihood function is written in terms of the Z_k

$$L = \prod_{k=1}^K \theta_k^{z_k} (1 - \theta_k)^{1-z_k}$$

$$= \prod_{k=1}^K \left[1 - \prod_{i=1}^{I_k} (1 - p_{ik}) \right]^{z_k} \left[\prod_{i=1}^{I_k} (1 - p_{ik}) \right]^{1-z_k}$$
 - Xie (*Statistics in Medicine*, 2001)
 - Likelihood function is written in terms of the Y_{ik}
 - EM algorithm used

Forming groups

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- Alike
 - Individuals with “similar” covariates are put into pools
 - Smallest variability in parameter estimates
 - How implement?
 - One covariate: Sort by covariate, then assign successive individuals to pools
 - Multiple covariates: ?
 - Usually requires one to have all individual testing specimens up front and available for testing at the same time

Forming groups

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- Random
 - Individuals are assigned to pools at random
 - Emulates chronological if no dependence over time
- Different
 - Pool individuals with covariates as different as possible
 - Emulates “worse case scenario” (?)

Forming groups

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- Simulate data from model fitted to the individual observations in Vansteelandt et al. (*Biometrics*, 2000)
 - $\text{logit}(p_{ik}) = \beta_0 + \beta_1 x_{ik} = -1.97 - 0.024x_{ik}$
 - Simulate the individual and group responses
 - $I = 7$ subjects per group
 - $K = 100$ groups
 - Overall sample size is $I*K = 700$

Forming groups

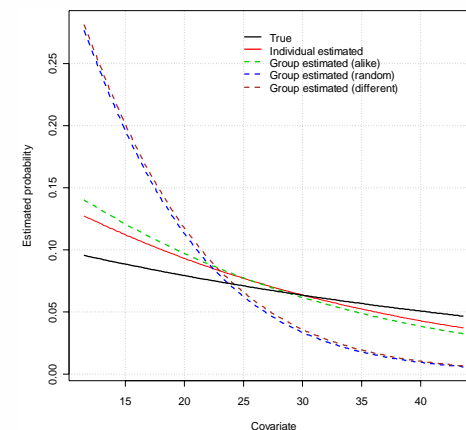
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- One simulated data set
- Relative efficiency

$$\widehat{RE}(\hat{\beta}_1) = \frac{\widehat{AsVar}(\hat{\beta}_1^{Individual})}{\widehat{AsVar}(\hat{\beta}_1^{Group})}$$

	$\widehat{RE}(\hat{\beta}_1)$
Alike	0.71
Random	0.12
Different	0.03

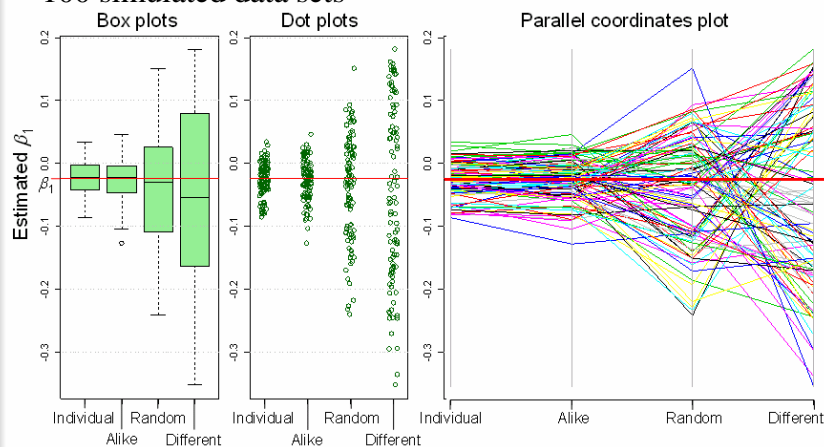


Forming groups

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- 100 simulated data sets



- Pearson correlations:

	Individual	Alike	Random
Alike	0.85		
Random	0.33	0.24	
Different	-0.05	-0.09	-0.13

Forming groups

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- Last slide examined a fixed $I*K$
- What if we fix the number of groups (tests), K , instead?

Settings

- $\text{logit}(p_{ik}) = -2 + 0.6931x_{ik}$
- $x_{ik} \sim \text{Uniform}(-70.079, 1.663)$
- $0.001 < p_{ik} < 0.3$
- Average value of p_{ik} is 0.02
- 500 simulated data sets for each simulation

Relative efficiency:

$K = 500$	I		
	2	5	10
Alike	2.20	4.62	6.72
Random	1.61	1.79	1.50
Different	1.16	0.51	0.22

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- Proposed

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