Statistical Issues Related to AVA Vaccine Evaluation

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Statistical Issues Related to AVA Vaccine Evaluation

Correlates of protection studies

Form statistical models that link
the results of animal studies to
those of human clinical trials

➤ Missing Data

Techniques which may be applied to resolve missing data

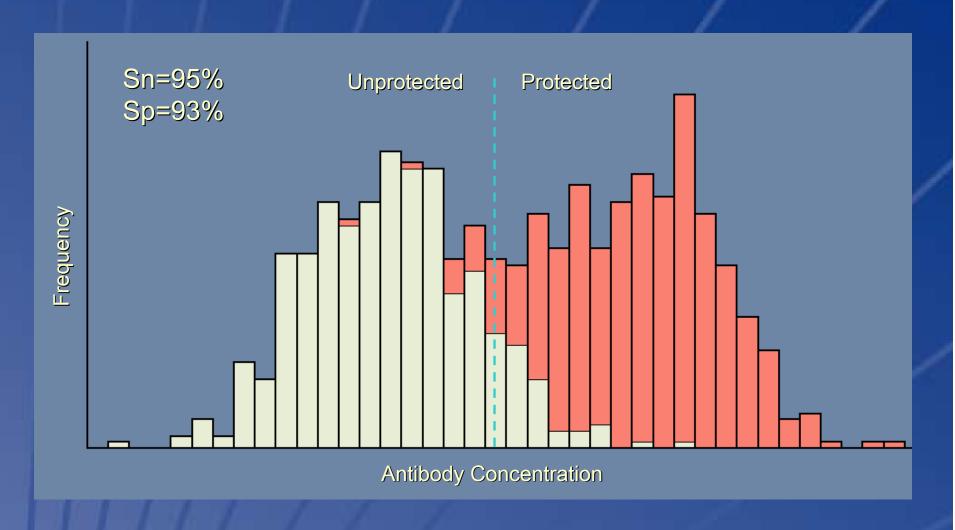


Predicting Protection (Survival of Challenge)

Given that we can measure more than one correlate of protection

How do we utilize them to predict protection more effectively than each test individually?



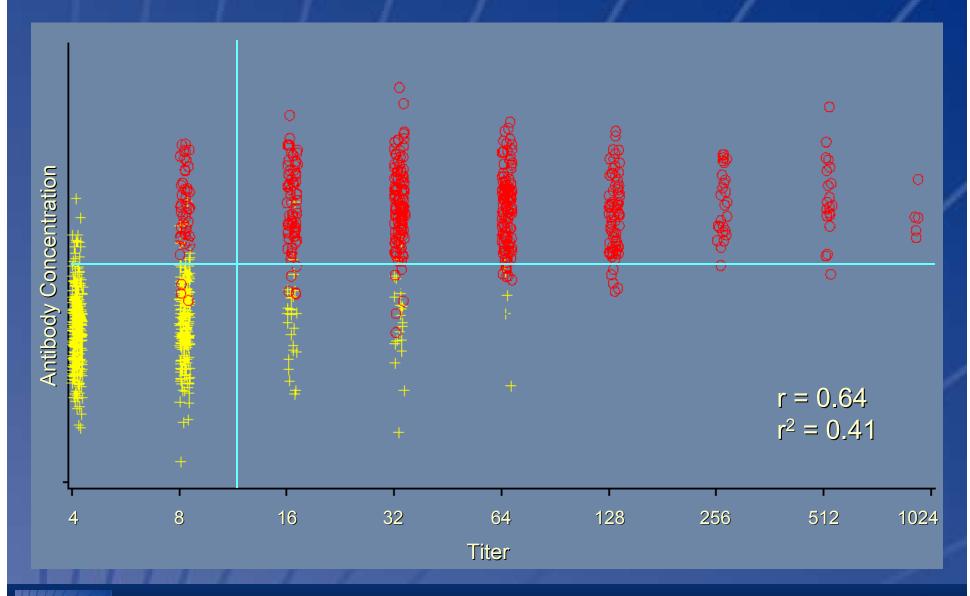








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Predictive Model Building

- Multiple approaches
 - logistic discriminant analysis, cluster analysis, other exploratory techniques (CART, etc.)
- Logistic Discriminant Analysis
 - Model relates a series of assay endpoints to protection
 - Construct a discriminant function from results:

$$IS = b_1 x_1 + b_2 x_2 + ... + b_p x_p$$

IS = immunologic 'score'

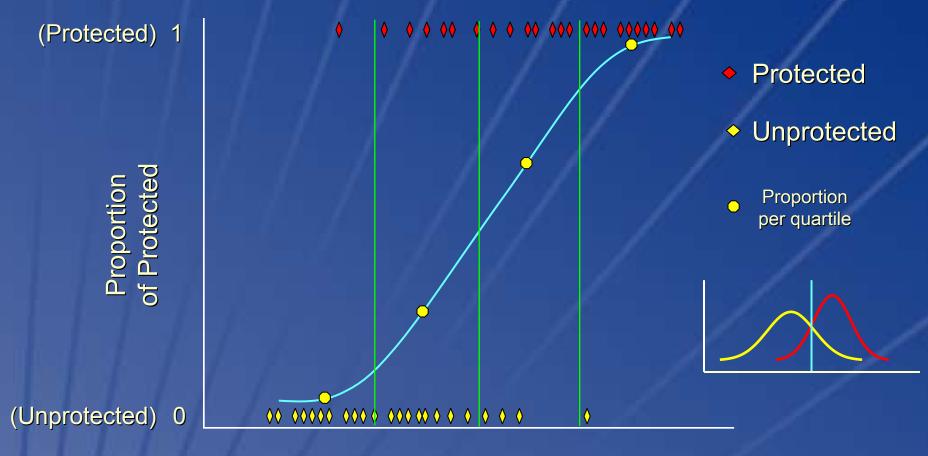
 $x_i = correlate data$

b_i = model coefficients

p = number of variables in the model



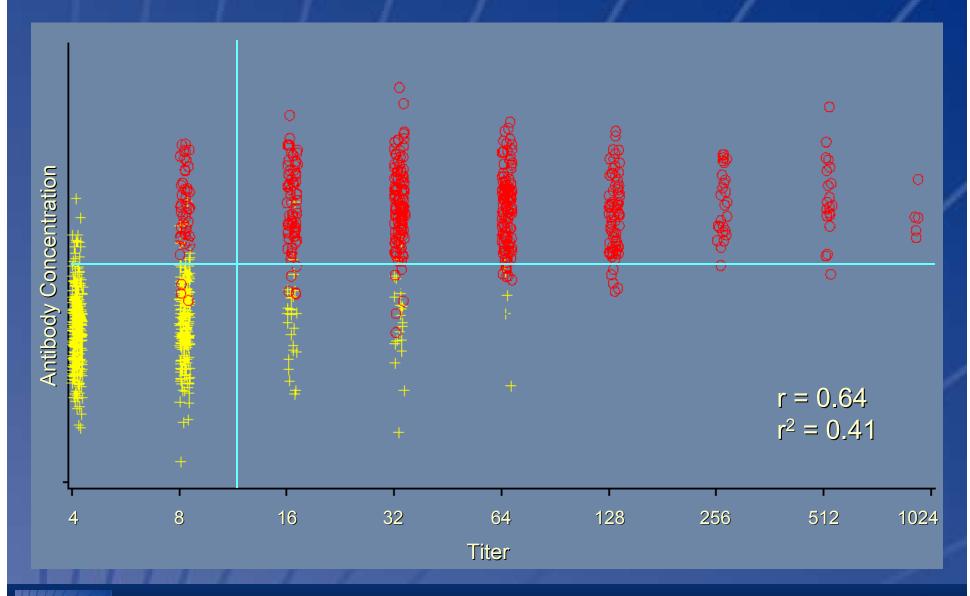
Immune Response vs. Protection Model Building



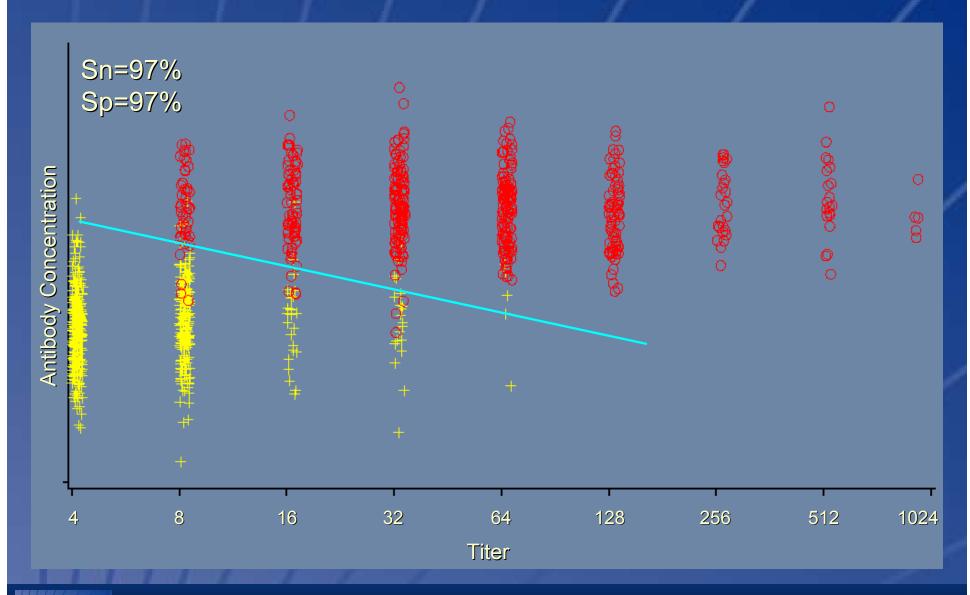
Antibody Concentration



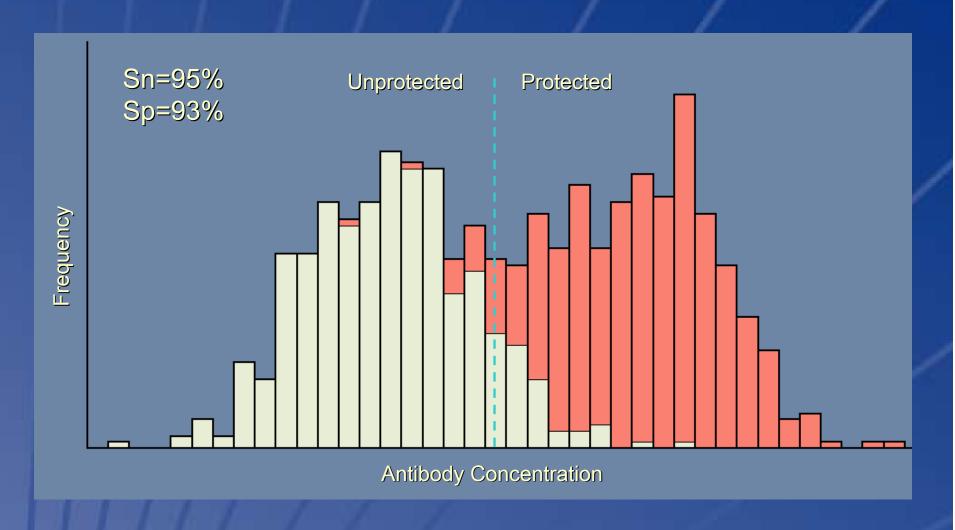
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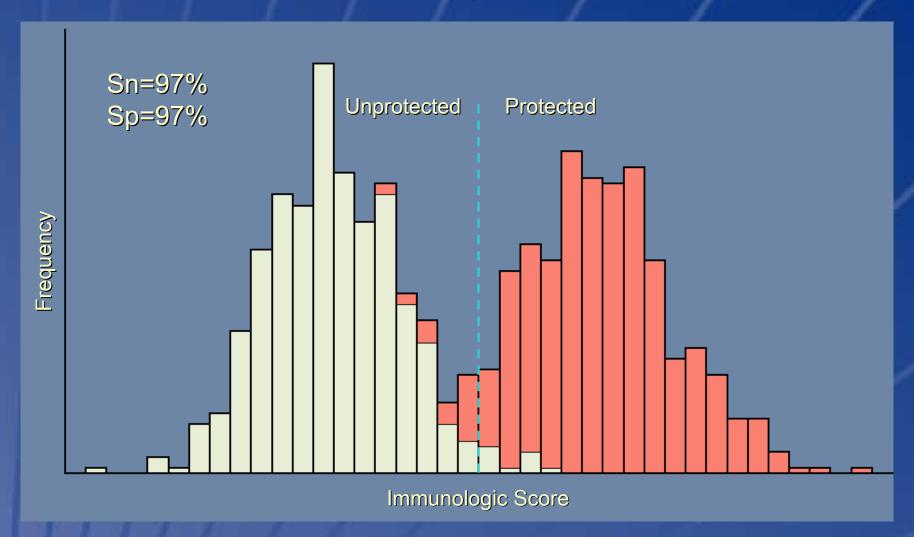






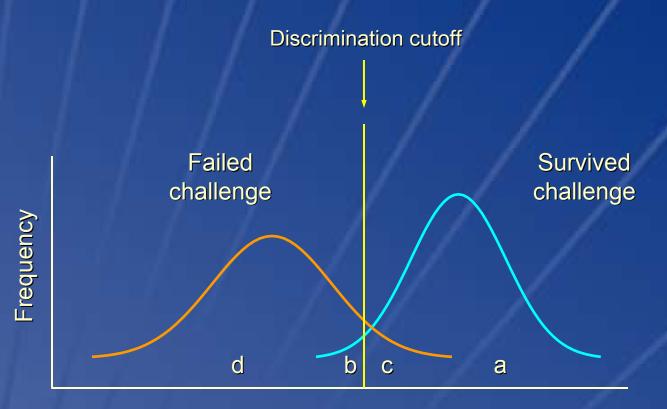








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Classification Index

b = survivors incorrectly classified as failed

a = survivors correctly classified as survived

c = failed, incorrectly classified as survived

d = failed, correctly classified as failed



| | | Known Outcome | | |
|--------------|-----------------------|-----------------------|---------------------|--|
| | | Survived Challenge | Failed Challenge | |
| Discriminant | Survived Challenge | а | С | |
| Analysis | Failed Challenge | b | d | |

- If survived challenge is defined as a 'positive' event then sensitivity = [a / (a+b)]
- If failed challenge is defined as a 'negative' event then specificity = [d / (c+d)]



- > Estimate model coefficients
 - Select variables with greatest discriminating power
- Calculate classification indexes for all subjects
- Discrminant cutoff may be placed to maximize sensitivity or specificity



- Apply discriminant function to subset of clinical trial participants
- Examine how vaccinated and unvaccinated individuals are classified
- Determine if these variables convey information about vaccine protection



Analytical Model

```
Outcome Animal # Dil. Assay 1 Assay 2 Assay_j
Survived / Died 1 1:40 1, 2, 3, ... A_{1k} 1, 2, 3, ... A_{2k} ··· 1, 2, 3, ... A_{jk}
```

```
j Assaysk Bleeds per animalj x k Assay data points per animal
```



Complete Data

| Animal | Assay 1 | Assay 2 | Assay 3 | Assay 4 | Assay 5 |
|--------|---------|---------|---------|---------|------------|
| 1 | / | / | | / | // |
| 2 | | | | | <i>-</i> [|
| 3 | / | -//- | | // | |
| 4 | - 1/- | | | | 11119 |
| 5 | | | | | |
| 6 | | | | | |
| 7 | 4-4 | | -1/ | | |
| 8 | 1-1/-) | | | | |
| 9 | 1-4-4-1 | / | | | |
| 10 | 1.1.1 | -//- | | (| |



| Animal | Assay 1 | Assay 2 | Assay 3 | Assay 4 | Assay 5 |
|--------|---------|---------|---------|------------|---------|
| 1 | X - | | //- | | - X |
| 2 | / | | X | 1,2111 | - |
| 3 | X | -//- | / | <i>[</i>] | |
| 4 | | X | X | | فيتتنا |
| 5 | X | | //- | X - | / . |
| 6 | | X | | | |
| 7 | | | / | | X |
| 8 | 4-7/- | 4 | | X - | |
| 9 | - X | X | | | |
| 10 | | | | | |



Imputation Strategies

- Recognized by the FDA
- Several methods
 - Last (worst) value carried forward
 - Hot deck
 - Listwise deletion complete case
 - Mean & regression substitution
 - Pairwise deletion
 - Multiple imputation

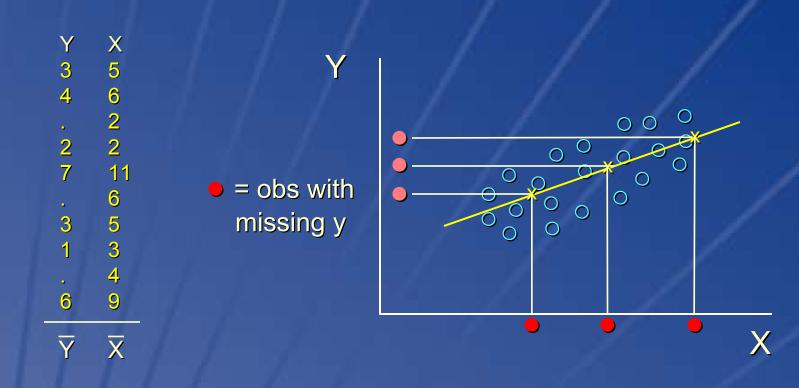


Listwise Deletion – Complete Case Analysis

- If data MCAR then probably OK
 - Complete data random sample
- If data not MCAR then
 - Point estimates biased
 - Inflated variances (smaller n)



Mean & Regression Substitution





Missing Data Mean Substitution

- Point estimates may be biased
- Underestimate standard errors



Pairwise Deletion - Available Case Analysis

| X | Y |
|-------------|--------|
| / | 2 |
| | 10 |
| | 4 |
| 5 | 6 |
| 2 | 2 |
| 5 2 7 | 2 6 |
| 3 | 2 |
| 1 | 2 7 |
| | 3 |
| 3 7 | |
| 7 | |
| 5 | |
| | |

For mean X, S_X²

For mean Y, S_Y^2

For S_{XY}², r_{XY}

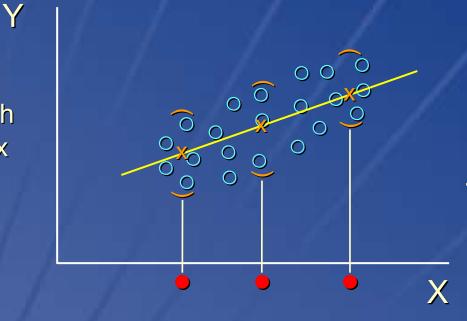
Pairwise Deletion - Available Case Analysis

- > Advantages:
 - Easy, but inferences problematic
 - Makes apparent better use of data
- Disadvantages:
 - Correlations may fall outside of (-1,1)
 - Cov. matrix has undesirable properties
 - May be even worse than compl. case



Multiple Imputation

= obs with missing x



$$y_i = a + bx_i + \varepsilon_i$$

$$y_i = a_j + b_j x_i + \varepsilon_i$$



Multiple Imputation – Example

X Y Y (1) Y (2)

N=37
?=113

N=75
?=75

N=150

Y values imputed under 3 correlation structures low, medium, high



N=150

Multiple Imputation – Example

50 imputations (Mean Y ± std. error)

| | 1 | Correlation | |
|--------------------|--------------|-------------------------------|--------------|
| Sample | Low | Medium | High |
| Full (n=150) | 29.53 ± 0.47 | 16.61 ± 0.17 | 19.48 ± 0.20 |
| 1/2 Missing (n=75) | 28.61 ± 0.71 | 16.75 ± 0.23 | 19.57 ± 0.24 |
| 3/4 Missing (n=37) | 30.79 ± 0.99 | 16.44 ± 0.26 | 19.35 ± 0.24 |



Multiple Imputation

- > Advantages:
 - Estimators consistent, asym. efficient and normal
 - Can be used with any kind of data or model
 - Analysis can be performed in SAS
- Disadvantages:
 - Easy to do in the wrong way
 - May get slightly different answers with each imputation



Missing Data Multiple Imputation

- Individual data points unimportant
- Primary goal unbiased estimates
 - Model parameters
 - Standard errors
 - Statistics
- Standard errors reflect degree of 'missingness'

