

Empirical Likelihood Estimation of a Diagnostic Test Likelihood Ratio

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Outline

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- An Introduction to Diagnostic Test Likelihood Ratios

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- An Empirical Likelihood Function for ρ_X

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- An Illustrative Example

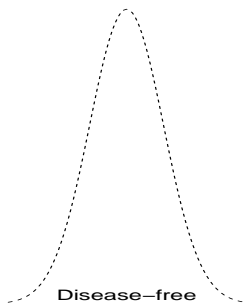
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- An Introduction to Diagnostic Test Likelihood Ratios
- An Empirical Likelihood Function for ρ_x
- An Illustrative Example
- Concluding Remarks

Introduction

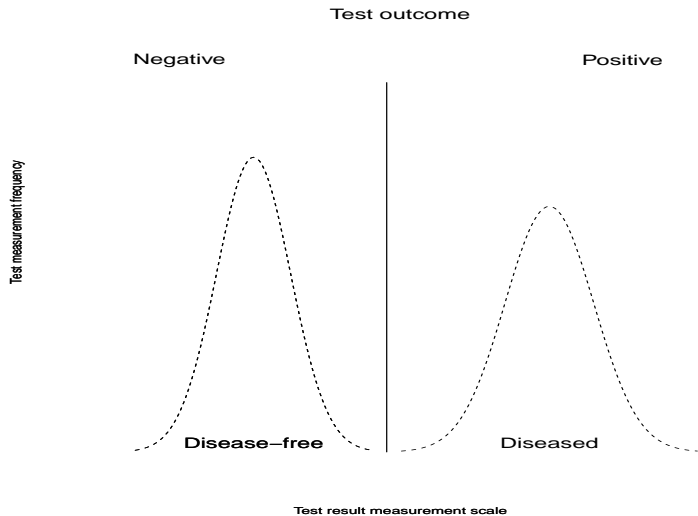
- Assume we have two subpopulations, diseased and disease-free individuals; label the former group 1 and the latter group 2

Test measurement frequency

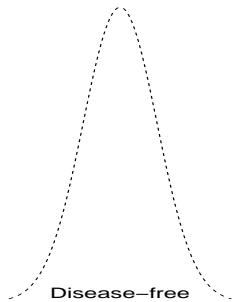


Disease-free

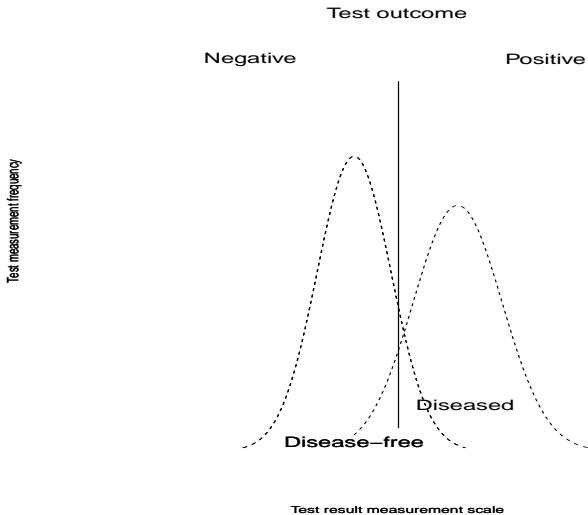
Test result measurement scale



Test measurement frequency



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- In the terminology of diagnostic testing, p_1 is the test sensitivity, and p_2 is the probability of a false positive test error, or 1 minus the test specificity
- Since 1975, the ratios

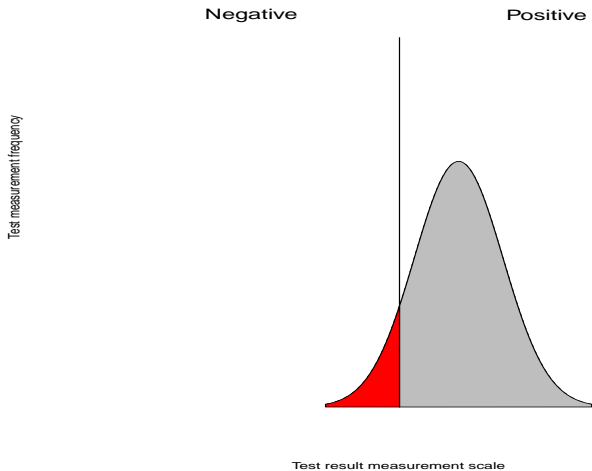
$$\rho_+ = p_1/p_2$$

and

$$\rho_- = (1 - p_1)/(1 - p_2)$$

have been of particular interest to advocates of evidence-based medicine

Test outcome in diseased group

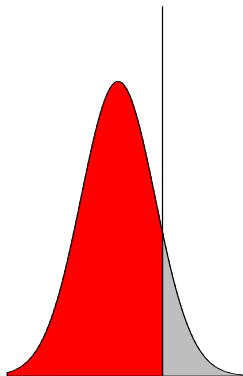


Test outcome in disease-free group

Negative

Positive

Test measurement frequency

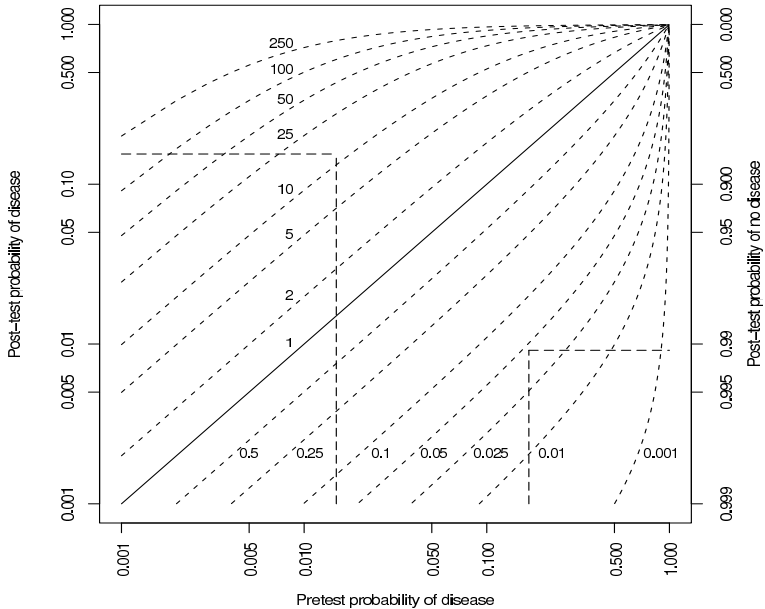


Test result measurement scale

- These functions of sensitivity and specificity have been called the “likelihood ratio of a positive test result” and the “likelihood ratio of a negative test result,” as a consequence of the books by Lusted (1968) and Sackett *et al.* (1991)

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$$\begin{aligned} & \frac{\text{Pr}(\text{disease}|\text{positive test})}{\text{Pr}(\text{no disease}|\text{positive test})} \\ &= \frac{\text{Pr}(\text{positive test}|\text{disease})}{\text{Pr}(\text{positive test}|\text{no disease})} \times \frac{\text{Pr}(\text{disease})}{\text{Pr}(\text{no disease})} \\ &= \rho_+ \frac{\text{Pr}(\text{disease})}{\text{Pr}(\text{no disease})} \end{aligned}$$

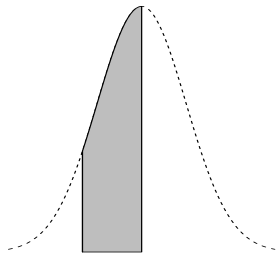


- Suppose the test result is classified into $K > 2$ categories, e.g., for iron-deficiency anemia, Guyatt *et al.* (1992) report

Group	Serum ferritin concentration ($\mu\text{gm/L}$)					
	[0, 15)	[15, 25)	[25, 35)	[35, 45)	[45, 100)	≥ 100
Diseased	474	117	58	36	76	48
Disease-free	20	29	50	43	398	1320

Test outcome in diseased group

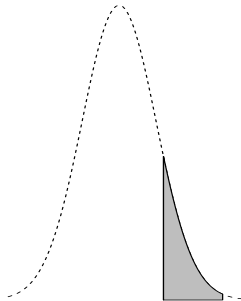
Test measurement frequency



Test result measurement scale

Test outcome in disease-free group

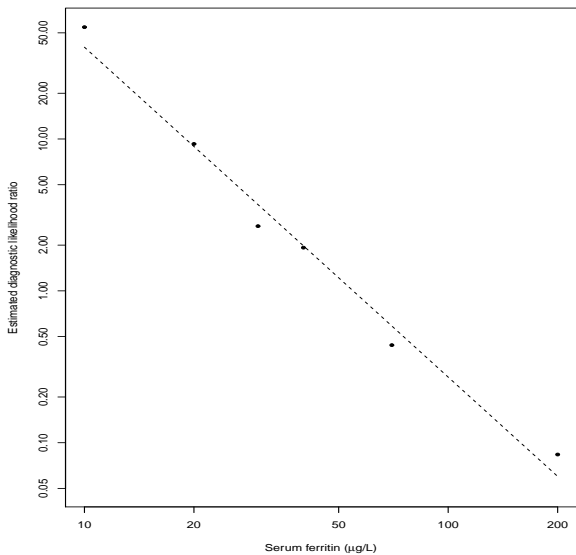
Test measurement frequency



Test result measurement scale

- By analogy with the case of $K = 2$ categories the corresponding table of estimated DLRs for each of the serum ferritin test result categories would be

Group	Serum ferritin concentration ($\mu\text{gm/L}$)					
	[0, 15)	[15, 25)	[25, 35)	[35, 45)	[45, 100)	≥ 100
Estimated DLR	54.5	9.3	2.7	1.9	0.4	0.1



- If we push the envelope for multiple categories to the limit, then the corresponding DLR for each category becomes

$$\lim_{h \rightarrow 0^+} \frac{\mathcal{F}_1(x) - \mathcal{F}_1(x+h)}{\mathcal{F}_2(x) - \mathcal{F}_2(x+h)} = \frac{f_1(x)}{f_2(x)} = \rho_x$$

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- Since each probability density function can be conveniently expressed in terms of the corresponding hazard function, i.e.,

$$f_i(x) = h_i(x) \exp \left\{ - \int_0^x h_i(s) ds \right\}$$

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- Formulate the estimation problem using the two-sample time-to-response framework of Kaplan-Meier (1958)

An Empirical Likelihood Function for ρ_x

- Denote the ordered, distinct response measurements in the two samples by

$$\text{Diseased} \quad x_{11} < x_{12} < \cdots < x_{1n}$$

$$\text{Disease-free} \quad x_{21} < x_{22} < \cdots < x_{2m}$$

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- Define d_{ij} and r_{ij} , the respective event and the risk sets in sample i at response measurement x_{ij} .
- The nonparametric log-likelihood function for $\mathbf{h} = \{h_{ij}\}$, based on these data, is

$$\begin{aligned} \ell(\mathbf{h}) = & \sum_{j=1}^n \{d_{1j} \log h_{1j} + (r_{1j} - d_{1j}) \log(1 - h_{1j})\} \\ & + \sum_{k=1}^m \{d_{2k} \log h_{2k} + (r_{2k} - d_{2k}) \log(1 - h_{2k})\} \end{aligned}$$

- Let t denote a fixed value of the response measurement; represent the corresponding value of the DLR at $x = t$ by ρ_t (but suppress the dependence on t subsequently); then

$$\log \rho_t = \log h_1(t) - \int_0^t h_1(s) ds - \log h_2(t) + \int_0^t h_2(s) ds ,$$

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- Then $\ell(\rho_t)$, the profile log-likelihood for ρ_t , can be obtained by evaluating the constrained MLEs, \tilde{h}_{ij} , that maximize

$$\begin{aligned} \ell_{\xi}(\rho_t) = & \ell + \xi \left\{ \log h_{1t} - \sum^{(t)} \log(1 - h_{1j}) \right. \\ & \left. - \log h_{2t} + \sum^{(t)} \log(1 - h_{2k}) - \log \rho_t \right\} \end{aligned}$$

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- The score equations for $\mathbf{h} = \{h_{ir}\}$ that lead to the constrained MLEs, $\tilde{\mathbf{h}} = \{\tilde{h}_{ir}\}$, are

$$\begin{aligned} \partial \ell_{\xi} / \partial h_{1j} &= d_{1j} / h_{1j} - (r_{1j} - d_{1j} - \xi) / (1 - h_{1j}) = 0, \\ &\quad \text{if } x_{1j} < t, \\ &= (d_{1t} + \xi) / h_{1t} - (r_{1t} - d_{1t}) / (1 - h_{1t}) = 0 \\ &\quad \text{if } x_{1j} = t \\ &= d_{1j} / h_{1j} - (r_{1j} - d_{1j}) / (1 - h_{1j}) = 0 \\ &\quad \text{if } x_{1j} > t \end{aligned}$$

$$\begin{aligned}
\partial \ell_{\xi} / \partial h_{2k} &= d_{2k} / h_{2k} - (r_{2k} - d_{2k} + \xi) / (1 - h_{2k}) = 0 \\
&\quad \text{if } x_{2k} < t \\
&= (d_{2t} - \xi) / h_{2t} - (r_{2t} - d_{2t}) / (1 - h_{2t}) = 0 \\
&\quad \text{if } x_{2k} = t \\
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&\quad \text{if } x_{2k} > t
\end{aligned}$$

i.e.,

$$\begin{aligned}
\tilde{h}_{1j} &= d_{1j} / (r_{1j} - \xi), & \text{if } x_{1j} < t \\
&= (d_{1t} + \xi) / (r_{1t} + \xi), & \text{if } x_{1j} = t \\
&= d_{1j} / r_{1j}, & \text{if } x_{1j} > t \\
\tilde{h}_{2k} &= d_{2k} / (r_{2k} + \xi), & \text{if } x_{2k} < t \\
&= (d_{2k} - \xi) / (r_{2k} - \xi), & \text{if } x_{2k} = t \\
&= d_{2k} / r_{2k}, & \text{if } x_{2k} > t
\end{aligned}$$

- It follows that the LRS for $\log \rho_t$, and hence for ρ_t , is equal to

$$\begin{aligned}
 & 2\{\ell(\hat{\mathbf{h}}) - \ell(\tilde{\mathbf{h}})\} \\
 &= 2 \sum^{[t]} \left[d_{1j} \log(\hat{h}_{1j}/\tilde{h}_{1j}) + (r_{1j} - d_{1j}) \log \left\{ \frac{1 - \hat{h}_{1j}}{1 - \tilde{h}_{1j}} \right\} \right] \\
 & \quad + 2 \sum^{[t]} \left[d_{2k} \log(\hat{h}_{2k}/\tilde{h}_{2k}) + (r_{2k} - d_{2k}) \log \left\{ \frac{1 - \hat{h}_{2k}}{1 - \tilde{h}_{2k}} \right\} \right] \\
 &= 2 \sum^{(t)} \left[r_{1j} \log \left(1 - \frac{\xi}{r_{1j}} \right) - (r_{1j} - d_{1j}) \log \left\{ 1 - \frac{\xi}{r_{1j} - d_{1j}} \right\} \right] \\
 & \quad + 2 \sum^{(t)} \left[r_{2k} \log \left(1 + \frac{\xi}{r_{2k}} \right) - (r_{2k} - d_{2k}) \log \left\{ 1 + \frac{\xi}{r_{2k} - d_{2k}} \right\} \right] \\
 & \quad + 2 \left[r_{1t} \log \left(1 + \frac{\xi}{r_{1t}} \right) - d_{1t} \log \left(1 + \frac{\xi}{d_{1t}} \right) \right. \\
 & \quad \quad \left. + r_{2t} \log \left(1 - \frac{\xi}{r_{2t}} \right) - d_{2t} \log \left(1 - \frac{\xi}{d_{2t}} \right) \right],
 \end{aligned}$$

- A $100(1 - \alpha)\%$ CI for ρ_t is found by solving the inequality

$$-2r(\rho_t) \leq c_{1,\alpha}^*$$

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- In practice, solve the equation

$$-2r(\rho_t) = c_{1,\alpha}^*$$

for the two zeros, $\xi_- < 0$ and $\xi_+ > 0$; use these values to calculate the corresponding lower and upper confidence bounds for ρ_t

- Via linear and quadratic expansions of various log functions, we can show the LRS is approximately equal to

$$\frac{(\log \hat{\rho}_t - \log \tilde{\rho}_t)^2}{V_t},$$

where

$$V_t = \sum^{(t)} \{1/(r_{1j} - d_{1j}) - 1/r_{1j}\} + (1/d_{1t} - 1/r_{1t}) \\ + \sum^{(t)} \{1/(r_{2k} - d_{2j}) - 1/r_{2k}\} + (1/d_{2t} - 1/r_{2t})$$

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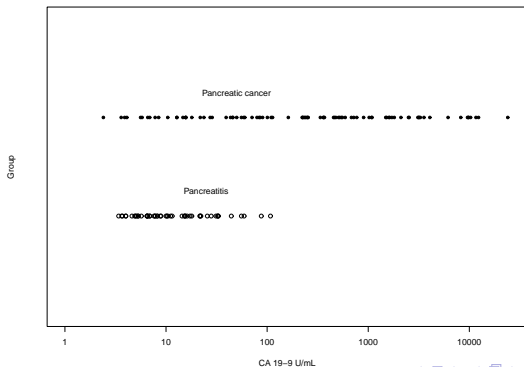
- This corresponds to the usual form of a Wald statistic, based on the MLE, used to test a hypothesis concerning log relative risk, i.e., $\log \rho_t$

An Illustrative Example

- Wieand *et al.* (1989) report results of CA 19-9 (cancer antigen) diagnostic test measurements. A total of 141 measurements were recorded, 51 from disease-free individuals (with pancreatitis) and 90 from subjects with confirmed pancreatic cancer.

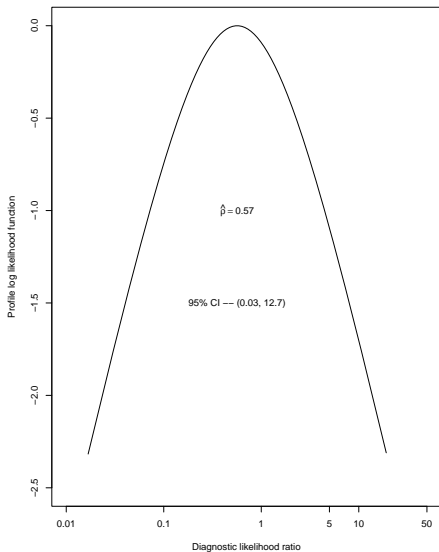
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- If we fix the value of t at 21.8 U/mL

the resulting profile log-likelihood is



Concluding Remarks

- In the absence of any distributional assumptions, empirical likelihood provides a convenient basis on which to estimate the DLR, ρ_x , for a continuous-scale test measurement

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- In the absence of any distributional assumptions, empirical likelihood provides a convenient basis on which to estimate the DLR, ρ_x , for a continuous-scale test measurement
- Empirical likelihood has the advantage that it is range-preserving, data-driven, and easy to construct; no variance estimate is required, and the resulting point or interval estimate is transformation-invariant
- Sensible estimates can only be derived at test measurements that are duplicated in both samples; additional assumptions, such as smoothness, should alleviate this drawback

Good medicine does not consist in the indiscriminate application of laboratory examinations to a patient, but rather in having so clear a comprehension of the probabilities of a case as to know what tests may be of value . . . it should be the duty of every hospital to see that no house officer receives his diploma unless he has demonstrated . . . a knowledge of how to use the results in the study of his patient.

Dr. George W. Peabody (1922)

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- 5 Wieand S, Gail MH, James BR, Jame KL: A family of nonparametric statistics for comparing diagnostic markers with paired or unpaired data. *Biometrika* 1989;76:585–592.