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**Original Paper** 

# Psychophysical Threshold Estimates in Logistic Regression Using the Bootstrap Resampling

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#### Abstract

We propose the non-parametric bootstrap resampling algorithm for the problem of psychophysical threshold estimates. We use the logistic regression with guessing rate and the log-likelihood ratio test statistics of two samples for testing the hypothesis by using the bootstrap resampling. We apply our algorithm to the visual acuity test, and show that the bootstrap resampling is useful for the problem of the two-sample test when the numbers of observations are not identical between the two samples. We also propose the bootstrap algorithm for one-sample testing to certify the values of parameters and threshold obtained by logistic regression.

# Introduction

The bootstrap resampling method provides a powerful procedure for estimating the variance of a parameter of a function. For this computer-based method we can refer to Efron et al. [1], Davison et al. [2], Foster et al. [3, 4], Joy et al. [5] and Good [6].

For the psychophysical experiment by constant stimuli method, Nagai et al. [7] proposed the statistical significance testing of difference between multiple thresholds. Bach [8], Beck et al. [9] and Schulze-Bonse et al. [10] developed the automated procedures on the personal computer for the measurements of visual acuity.

Mita et al. [11] developed a statistical method for evaluating the logarithmic visual acuity (LogVA) changes in an individual, and calculated LogVA  $\pm$  SD (SD : standard deviation) by logistic regression, and also evaluated it using Nagai's test of significant difference.

The categorial data analysis and the logistic regression have been studied by McCullagh et al. [12], Christensen [13], Harrell [14] and Agresti [15].

In the present paper we propose the non-parametric bootstrap resampling for the problem of psychophysical threshold estimates. We propose the logistic regression with guessing rate and formulation of deviance residuals in sections 2 and 3. We show the log-likelihood ratio test statistics in section 4, and

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the non-parametric bootstrap resampling and testing of hypothesis in sections 5 and 6. Finally, in section 7 we present an application of our algorithm to psychophysical threshold estimates in the visual acuity test.

# Logistic regression with guessing rate

We assume that the logit function is expressed in the form:

$$\operatorname{logit}(p_0) \equiv \operatorname{log} \frac{p_0}{1 - p_0} = \alpha + \beta x,$$

where  $p_0$  is the (primitive) probability, x is the explanatory variable and  $\alpha$ ,  $\beta$  are constants. Then  $p_0$  is given by

$$p_0(x; \alpha, \beta) = (1 + \exp(-\alpha - \beta x))^{-1}$$

We introduce the third parameter  $\gamma$  (  $0 \le \gamma < 1$  ) for including the guessing rate. Then we have the probability p such that

$$p(x; \alpha, \beta, \gamma) = p_0(x; \alpha, \beta) + \gamma(1 - p_0(x; \alpha, \beta)).$$

Let

$$X = \{x_j, \mu_j (j = 1, 2, \dots, N)\}$$

be the set of binomial observations where  $x_j$  ( $j = 1, 2, \dots, N$ ) are the explanatory variables for *j*-th ( $j = 1, 2, \dots, N$ ) observations respectively and  $\mu_j$  ( $j = 1, 2, \dots, N$ ) are outcome data:

$$\mu_j = \begin{cases} 1 & \text{if } j\text{-th outcome is "success",} \\ 0 & \text{if } j\text{-th outcome is "failure".} \end{cases}$$

Then the logarithmic binomial likelihood *L* ( $\alpha, \beta, \gamma$ ) is given by

$$L(\alpha, \beta, \gamma) = \log \prod_{j=1}^{N} p_{j}^{\mu j} (1-p_{j})^{1-\mu_{j}}$$
$$= \sum_{j=1}^{N} (\mu_{j} \log p_{j} + (1-\mu_{j}) \log(1-p_{j})),$$

where  $p_j = p(x_j; \alpha, \beta, \gamma)$  ( $j = 1, 2, \dots, N$ ). We assume that  $\gamma$  is a known constant  $\gamma_0$  ( $0 \le \gamma_0 < 1$ ). Then the partial derivatives of  $L(\alpha, \beta, \gamma_0)$  with respect to  $\alpha$  and  $\beta$  are given by

$$\frac{\partial L}{\partial \alpha} = \frac{1}{1 - \gamma_0} \sum_{j=1}^N (\mu_j - p_j) \frac{p_j - \gamma_0}{p_j},$$
$$\frac{\partial L}{\partial \beta} = \frac{1}{1 - \gamma_0} \sum_{j=1}^N (\mu_j - p_j) \frac{p_j - \gamma_0}{p_j} x_j,$$
$$E\left[\frac{\partial^2 L}{\partial \alpha^2}\right] = -\sum_{j=1}^N \omega_j,$$
$$E\left[\frac{\partial^2 L}{\partial \alpha \partial \beta}\right] = E\left[\frac{\partial^2 L}{\partial \beta \partial \alpha}\right] = -\sum_{j=1}^N \omega_j x_j,$$
$$E\left[\frac{\partial^2 L}{\partial \beta^2}\right] = -\sum_{j=1}^N \omega_j x_j^2,$$

where *E* [X] is the expected value of *X*, and  $\omega_j$  is defined by

$$\omega_j \equiv \frac{1-p_j}{p_j} \left(\frac{p_j-\gamma_0}{1-\gamma_0}\right)^2.$$

We define the following notations for easy description:

$$f \equiv \frac{\partial L}{\partial \alpha}, \quad g \equiv \frac{\partial L}{\partial \beta}$$

We shall obtain  $\alpha$  and  $\beta$  by adopting the Fisher score method. Let  $\alpha^t$ ,  $\beta^t$ ,  $f^t$ ,  $g^t$  ( $t = 0, 1, 2, \cdots$ ) be the values of  $\alpha$ ,  $\beta$ , f, g at iterative step t ( $t = 0, 1, 2, \cdots$ ) and let  $\alpha^0 = \beta^0 = 0$ . Then we can write the algorithm for determining  $\alpha$  and  $\beta$  such that

$$\begin{split} v^{t+1} &= v^t + (F^t)^{-1} s^t (t=0, \ 1, \ 2, \ \cdots); \\ v^t &\equiv \begin{pmatrix} \alpha^t \\ \beta^t \end{pmatrix}, \\ s^t &\equiv \begin{pmatrix} f^t \\ g^t \end{pmatrix}, \\ F^t &\equiv -E \Big[ \Big( \frac{\partial \ (f^t, \ g^t)}{\partial \ (\alpha^t, \ \beta^t)} \Big) \Big], \end{split}$$

where ( $\partial$  (, ) /  $\partial$  (, )) is a Jacobian matrix. We stop the above iterative procedure if

Norm 
$$\equiv (v^{t+1} - v^t)^T (v^{t+1} - v^t) < \varepsilon$$

is satisfied for sufficiently small positive number  $\varepsilon$ .

Let  $\hat{\alpha}$ ,  $\hat{\beta}$ , and  $\hat{F}$  be the optimal values of  $\alpha$ ,  $\beta$  and F respectively. Then by the Cramér-Rao lower bound, we can obtain variances such that

$$(\hat{F})^{-1} = \begin{pmatrix} \operatorname{var}(\hat{\alpha}) & \operatorname{cov}(\hat{\alpha}, \ \hat{\beta}) \\ \operatorname{cov}(\hat{\beta}, \ \hat{\alpha}) & \operatorname{var}(\hat{\beta}) \end{pmatrix} \\ = \begin{pmatrix} (\operatorname{se}(\hat{\alpha}))^2 & r_{a\beta} \operatorname{se}(\hat{\alpha}) \operatorname{se}(\hat{\beta}) \\ r_{\rho\alpha} \operatorname{se}(\hat{\beta}) \operatorname{se}(\hat{\alpha}) & (\operatorname{se}(\hat{\beta}))^2 \end{pmatrix},$$

where var  $(\hat{\alpha})$  and var  $(\hat{\beta})$  are variances of  $\hat{\alpha}$  and  $\hat{\beta}$  respectively,  $\operatorname{cov}(\hat{\alpha}, \hat{\beta})(=\operatorname{cov}(\hat{\beta}, \hat{\alpha}))$  is the covariance of  $\hat{\alpha}$  and  $\hat{\beta}$ ,  $\operatorname{se}(\hat{\alpha})$  and  $\operatorname{se}(\hat{\beta})$  are standard errors of  $\hat{\alpha}$  and  $\hat{\beta}$  respectively,  $r_{\alpha\beta}(=r_{\beta\alpha})$  is the correlation factor between  $\hat{\alpha}$  and  $\hat{\beta}$ .

# Deviance and deviance residual

Let  $\hat{\ell}$  be the maximum binomial likelihood:

$$\hat{\ell} = \prod_{j=1}^{N} \hat{p}_{j}^{\mu_{j}} (1 - \hat{p}_{j})^{1 - \mu_{j}},$$

where  $\hat{p}_{j}(j=1, 2, \dots, N)$  is the probability given by optimal parameters  $\hat{\alpha}, \hat{\beta}$  and  $\gamma_{0}$ 

$$\hat{p}_{j} = \hat{p}_{0}(x_{j}) + \gamma_{0}(1 - \hat{p}_{0}(x_{j})),$$
  
 $\hat{p}_{0}(x_{j}) = (1 + \exp(-\hat{\alpha} - \hat{\beta}x_{j}))^{-1}$   
 $(j = 1, 2, \dots, N).$ 

Then we can obtain the deviance D of logistic regression:

$$D = -2\log\hat{\ell}.$$

If we adopt the following notation:

$$d_{j} \equiv 2\mu_{j}\log\frac{1}{\hat{p}_{j}} + 2(1-\mu_{j})\log\frac{1}{1-\hat{p}_{j}}$$

$$(j = 1, 2, \dots, N),$$

the deviance D is given by

$$D=\sum_{j=1}^N d_j.$$

The deviance residual  $\varepsilon_j$  is given by

$$\varepsilon_{j} = \operatorname{sgn}(\mu_{j} - \hat{p}_{j}) / d_{j}$$

$$(j = 1, 2, \dots, N),$$

where sgn(y) is the sign function:

$$\operatorname{sgn}(y) = \begin{cases} 1 & \text{if } y > 0, \\ 0 & \text{if } y = 0, \\ -1 & \text{if } y < 0. \end{cases}$$

Then we can write the deviance residual  $\varepsilon_j$  explicitly such that

$$\varepsilon_{j} = \begin{cases} -\sqrt{2\log\frac{1}{1-\hat{p}_{j}}} & \text{if } \mu_{j} = 0, \\ \sqrt{2\log\frac{1}{\hat{p}_{j}}} & \text{if } \mu_{j} = 1 \end{cases}$$
$$(j = 1, 2, \cdots, N).$$

Log-likelihood ratio test statistics of two-sample problem

Let  $X_1$  and  $X_2$  be two samples from the populations which have possibly different probability distributions  $\Phi_1$  and  $\Phi_2$  respectively. We shall test the following hypothesis:

null hypothesis 
$$H_0: \Phi_1 = \Phi_2$$
,

alternative hypothesis  $H_1: \Phi_1 \neq \Phi_2$ .

Let  $\hat{\ell}_k(k=1, 2)$  be the maximum binomial likelihood of samples  $X_k(k=1, 2)$  respectively. Let  $X_3$  be the combined sample of  $X_1$  and  $X_2$ :

$$X_3 = X_1 \bigcup X_2.$$

Let  $\hat{\ell}_3$  be the maximum binomial likelihood of sample  $X_3$ . Then we can define the log-likelihood ratio test statistics *G* such that:

$$G = -2\log\frac{\hat{\ell}(H_0)}{\hat{\ell}(H_1)} = -2\log\frac{\hat{\ell}_3}{\hat{\ell}_1\hat{\ell}_2},$$

where  $\hat{\ell}(H_0)$  is the maximum binomial likelihood if  $H_0$  is satisfied, and  $\hat{\ell}(H_1)$  is the maximum binomial likelihood if  $H_1$  is satisfied. Let  $D_k$  (k = 1, 2, 3) be the deviances which are obtained by logistic regression analysis for samples  $X_k$  (k = 1, 2, 3) respectively.  $D_k$  (k = 1, 2, 3) are given by

$$D_k = -2\log \hat{\ell}_k (k = 1, 2, 3)$$

Then we have the log-likelihood ratio statistics G for the two-sample test:

$$G = D_3 - (D_1 + D_2).$$

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Non-parametric bootstrap resampling

# (i) Bootstrap samples $X_1^{*b}$ and $X_2^{*b}$

Let  $X_1$  be the set of binomial observations, and  $\varepsilon_1$  be the set of deviance residuals of sample 1. Let B be the number of bootstrap samples. By adopting uniform random numbers, we draw B samples of size  $N_1$ with replacements from  $\varepsilon_1$  and we call them the bootstrap deviance residuals of sample 1:

$$\varepsilon_1^{*b} = \left\{ \varepsilon_j^b \mid \varepsilon_j^b \in \varepsilon_1 (j = 1, 2, \dots, N_1) \right\} \quad (b = 1, 2, \dots, B)$$

Then we obtain the bootstrap sample  $X_1^{b}$  for sample 1 such that

$$X_1^{*b} = \{x_j, \ \mu_j^b (j=1, \ 2, \ \cdots, \ N_1)\} \ (b=1, \ 2, \ \cdots, \ B),$$

where

$$\mu_j^b = \hat{p}_0^b(x_j) + \gamma_0 (1 - \hat{p}_0^b(x_j)),$$
$$\hat{p}_0^b(x_j) = (1 + \exp(-\hat{\alpha}_1 - \hat{\beta}_1 x_j - \varepsilon_j^b))^{-1}$$
$$(j = 1, 2, \cdots, N_1; b = 1, 2, \cdots, B).$$

By adopting the similar method described above, we can also obtain bootstrap deviance residuals  $\varepsilon_2^{`b}$  ( $b = 1, 2, \dots, B$ ) and bootstrap samples  $X_2^{`b}$  ( $b = 1, 2, \dots, B$ ) from the set of binomial observations  $X_2$  of sample 2.

(ii) Bootstrap sample  $X_3^{*b}$ 

The bootstrap sample  $X_3^{*b}$  ( $b = 1, 2, \dots, B$ ) for sample 3 is obtained by the following Steps 1, 2 and 3.

#### Step 1:

Let  $\varepsilon_1^{b}$  be the bootstrap deviance residuals of sample 1:

$$\varepsilon_1^{*b} = \{ \varepsilon_j^b \mid \varepsilon_j^b \in \varepsilon_1 (j=1, 2, \dots, N_1) \} (b=1, 2, \dots, B).$$

We obtain the bootstrap sample  $X_{31}^{*b}$  ( $b = 1, 2, \dots, B$ ) by using  $\varepsilon_1^{*b}$  ( $b = 1, 2, \dots, B$ ) for sample 1 and the optimal parameters  $\hat{\alpha}_3$ ,  $\hat{\beta}_3$  and  $\gamma_0$  for sample 3 such that

$$X_{31}^{*b} = \{x_j, \ \mu_j^b (j=1, \ 2, \ \cdots, \ N_1)\} \ (b=1, \ 2, \ \cdots, \ B),$$

where

$$\mu_{j}^{b} = \hat{p}_{0}^{b}(x_{j}) + \gamma_{0} (1 - \hat{p}_{0}^{b}(x_{j})),$$
$$\hat{p}_{0}^{b}(x_{j}) = (1 + \exp(-\hat{\alpha}_{3} - \hat{\beta}_{3}x_{j} - \varepsilon_{j}^{b}))^{-1}$$
$$(j = 1, 2, \cdots, N_{1}; b = 1, 2, \cdots, B).$$

Step 2:

By adopting the similar method in step 1, we obtain the bootstrap sample  $X_{32}^{`b}$  ( $b = 1, 2, \dots, B$ ) by using  $\varepsilon_2^{`b}$  ( $b = 1, 2, \dots, B$ ) for sample 2 and the optimal parameters  $\hat{\alpha}_3$ ,  $\hat{\beta}_3$  and  $\gamma_0$  for sample 3.

Step 3:

By combining  $X_{31}^{'b}$  and  $X_{32}^{'b}$  ( $b = 1, 2, \dots, B$ ), we obtain the bootstrap sample  $X_3^{'b}$  ( $b = 1, 2, \dots, B$ ) for sample 3 such that

$$X_3^{*b} = X_{31}^{*b} \bigcup X_{32}^{*b} \quad (b = 1, 2, \dots, B).$$

# Hypothesis testing with the bootstrap resampling

Let  $D_k$  (k = 1, 2, 3) be the deviances obtained from the sets of binomial observations  $X_k$  (k = 1, 2, 3) respectively. Let  $D_k^b$  (k = 1, 2, 3;  $b = 1, 2, \dots, B$ ) be the bootstrap deviances obtained from the bootstrap samples  $X_k^{,b}$  (k = 1, 2, 3;  $b = 1, 2, \dots, B$ ) respectively. Let G and  $G^b$  ( $b = 1, 2, \dots, B$ ) be the log-likelihood ratio test statistics defined by

$$G = D_3 - (D_1 + D_2),$$
  

$$G^b = D_3^b - (D_1^b + D_2^b)$$
  
( b = 1, 2, ..., B).

Then we have the achieved significance level *ASL*:

$$ASL = \frac{\sum_{b=1}^{B} \lambda^{b}}{B},$$

where  $\lambda^{b}$  ( $b = 1, 2, \dots, B$ ) are the notations defined by

$$\lambda^{b} = \begin{cases} 1 & \text{if } G^{b} \ge G, \\ 0 & \text{if } G^{b} \le G \\ (b = 1, 2, \dots, B). \end{cases}$$

For avoiding ASL = 0, ASL is also defined by

$$ASL = \frac{\sum_{b=1}^{B} \lambda^b + 1}{B+1}$$

when  $\sum_{b=1}^{B} \lambda^b \leq \varepsilon$  ( $\varepsilon$  is a sufficiently small positive number). We can say that the null hypothesis  $H_0$  (two samples  $X_1$  and  $X_2$  have common probability distributions:  $\Phi_1 = \Phi_2$ ) is rejected if ASL is less than or equal to the significance level.

# Application to psychophysical threshold estimates

# (i) Mathematical notations and definitions

Let *X* be the set of binomial observations:

$$X = \{x_j, \mu_j (j = 1, 2, \dots, N)\}.$$

Let  $\Omega_i$  ( $i = 1, 2, \dots, n$ ) be the properly chosen intervals of the explanatory variable and let  $\bar{x}_i$  ( $i = 1, 2, \dots, n$ ) be the mid-point of  $\Omega_i$ . We assume that  $n \leq N$ . Then we define the following notations:

$$\delta_{ij} = \begin{cases} 1 & \text{if } x_j \in \Omega_i, \\ 0 & \text{if } x_j \notin \Omega_i \end{cases}$$
$$(i = 1, 2, \cdots, n; j = 1, 2, \cdots, N),$$

and

$$n_i = \sum_{j=1}^N \delta_{ij}, \quad m_i = \sum_{j=1}^N \delta_{ij} \mu_j \quad (i = 1, 2, \dots, n).$$

We note that  $N = \sum_{i=1}^{n} n_i$ .

Let  $\hat{p}(x) = p(x; \hat{\alpha}, \hat{\beta}, \gamma_0)$  be the probability given by optimal parameters  $\hat{\alpha}, \hat{\beta}$  and  $\gamma_0$  such that

$$\hat{p}(x) = \hat{p}_0(x) + \gamma_0(1 - \hat{p}_0(x)),$$
  
 $\hat{p}_0(x) = (1 + \exp(-\hat{\alpha} - \hat{\beta}x))^{-1}.$ 

We define the psychophysical threshold  $\xi$  with guessing rate  $\gamma_0$ 

$$\xi = \hat{p}^{-1} \left( \frac{1+\gamma_0}{2} \right).$$

(ii) The visual acuity test of the two-sample problem

Since we adopt the Landolt-C of four different orientations in our visual acuity test, the guessing rate  $\gamma_0$  is chosen as

$$\gamma_0 = 0.25.$$

The explanatory variable x in our measurement is the logarithmic visual acuity.

Let  $X_1$  (sample 1) and  $X_2$  (sample 2) be samples from the populations which have possibly different probability distributions  $\Phi_1$  and  $\Phi_2$  respectively. We shall test the following hypothesis:

null hypothesis  $H_0: \Phi_1 = \Phi_2$ ,

alternative hypothesis  $H_1: \Phi_1 \neq \Phi_2$ .

We took the data from 1 individual with no visual abnormalities in order to assess our bootstrap algorithm. The LogVA (Logarithmic Visual Acuity) is  $0.3681 \pm 0.0209$  in complete refractive correction and we adopt this data set as sample 1. The data of sample 2 is taken in +0.50*D* incomplete refractive correction from the same individual of sample 1.

Table 1 and Table 2 show the observed data of sample 1 ( $N_1 = 120$ ) and sample 2 ( $N_2 = 80$ ) respectively. Table 3 shows sample 3 ( $N_3 = 200$ ) which is constructed by the combined data of samples 1 and 2.

The logistic regression results of samples 1, 2 and 3 are shown in Table 4. Figures 1, 2, and 3 show the observed data and  $\hat{p}(x) = p(x; \hat{\alpha}, \hat{\beta}, \gamma_0)$  of samples 1, 2 and 3 respectively. Psychophysical thresholds  $\hat{\xi}_k$  (k = 1, 2, 3) at probability = (1 +  $\gamma_0$ ) / 2 = 0.625 are shown in Table 4.

Now we shall prove that the samples 1 and 2 are taken from the populations which have different distributions.

Table 1 Observed data of sar	nple 1
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i	ā	22	222	
ı	$\bar{x}_i$	$n_i$	$m_i$	$m_i/n_i$
1	0.156975	0	0	-
2	0.198368	20	19	0.95
3	0.244125	20	18	0.9
4	0.295278	20	18	0.9
5	0.353270	20	13	0.65
6	0.420216	20	8	0.4
7	0.499398	20	8	0.4
$N_1 = \sum_{i=1}^{7} n_i = 120$				

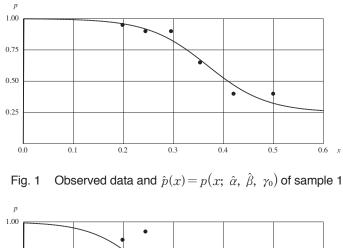
Table 2 Observed data of sample 2	2
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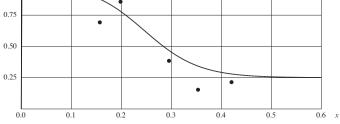
i	$\bar{x}_i$	$n_i$	$m_i$	$m_i/n_i$
1	0.156975	13	9	0.6923
2	0.198368	14	12	0.8571
3	0.244125	13	12	0.9231
4	0.295278	13	5	0.3846
5	0.353270	13	2	0.1538
6	0.420216	14	3	0.2143
7	0.499398	0	0	-
$N_2$	$N_2 = \sum_{i=1}^{7} n_i = 80$			

Table 3 Observed data of sample 3

i	$\bar{x}_i$	$n_i$	$m_i$	$m_i/n_i$	
1	0.156975	13	9	0.6923	
2	0.198368	34	31	0.9118	
3	0.244125	33	30	0.9091	
4	0.295278	33	23	0.6970	
5	0.353270	33	15	0.4545	
6	0.420216	34	11	0.3235	
7	0.499398	20	8	0.4	
$N_{i}$	$N_3 = \sum_{i=1}^7 n_i = 200$				

	5 5		1 ,
-	sample 1 $(k = 1)$	sample 2 $(k=2)$	sample 3 $(k=3)$
$N_k$	120	80	200
$\hat{\alpha}_k$	6.2105	4.5209	4.4349
$\hat{\beta}_k$	-16.8720	-18.3394	-14.1192
$\gamma_0$	0.25	0.25	0.25
$\operatorname{se}(\hat{\alpha}_k)$	1.4383	1.5872	0.9012
$\operatorname{se}(\hat{\beta}_k)$	4.2411	6.6365	3.0423
$se(\gamma_0)$	0.0	0.0	0.0
$\hat{\xi}_k$	0.3681	0.2465	0.3141
$\operatorname{se}(\hat{\xi}_k)$	0.0209	0.0214	0.0170
$D_k$	115.546	90.241	224.026
$G = D_{t}$	$(D_1 + D_2) = 18.$	.239	





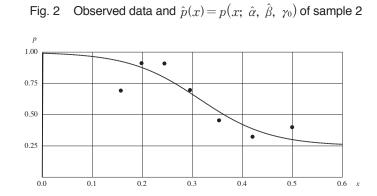


Fig. 3 Observed data and  $\hat{p}(x) = p(x; \hat{\alpha}, \hat{\beta}, \gamma_0)$  of sample 3

Table 4 Logistic regression results of samples 1, 2 and 3

The results of non-parametric bootstrap resampling are shown in Table 5. Since  $\Lambda(=\sum_{b=1}^{B}\lambda_b)$  is small for B = 2000, ASL is obtained as

$$ASL = \frac{\Lambda + 1}{B + 1} = 0.0005$$

This ASL shows that  $H_0$  is rejected at a very small significant level.

	sample 1 $(k=1)$	sample 2 $(k=2)$	sample 3 $(k=3)$	
$N_k$	120	80	200	
$B_k$	2000	2000	2000	
$\operatorname{mean}(D_k^b)$	8.962	6.612	17.862	
$G^{b} = D_{3}^{b} - (D_{1}^{b} + D_{2}^{b})  (b = 1, 2, \cdots, B)$				
$\Lambda = \sum_{b=1}^{B} \lambda^b = 0$				
$ASL = \frac{\Lambda}{B}$	$ASL = \frac{\Lambda + 1}{B + 1} = 0.0005$			

Table 5 Two-sample test by bootstrap resampling

#### (iii) The visual acuity test of one-sample problem

We use the same example described in (ii). We shall test here the parameters  $\alpha$ ,  $\beta$  and threshold  $\xi$  by using bootstrap resampling. Since the methods of one-sample test for  $\alpha$ ,  $\beta$  and  $\xi$  are the same, we show here only the case of  $\xi$ .

We adopt the following hypothesis:

null hypothesis  $H_0: \hat{\xi} = \hat{\xi}_c$ ,

alternative hypothesis  $H_1: \xi \neq \xi_c$ ,

where  $\xi_c$  is a prescribed value (which may be chosen from the threshold of control sample). Let  $\hat{\xi}$  and se  $(\hat{\xi})$  be the threshold and its standard error respectively of the (original) logistic regression. Let  $\hat{z}$  be the test statistics defined by

$$\hat{z} = \frac{\hat{\xi} - \xi_c}{\operatorname{se}(\hat{\xi})}.$$

Let  $\xi^b$  ( $b = 1, 2, \dots, B$ ) be the thresholds obtained by the logistic regression of each bootstrap resampling. Let  $\xi^{\hat{b}}$  be the mean of  $\xi^{\hat{b}}$  ( $b = 1, 2, \dots, B$ ):

$$\bar{\xi} = \frac{1}{B} \sum_{b=1}^{B} \xi^{b}.$$

Let  $\operatorname{se}(\hat{\xi})$  be the standard error of  $\hat{\xi}^b$  ( $b = 1, 2, \dots, B$ ):

$$\operatorname{se}(\hat{\xi}) = \sqrt{\frac{1}{B-2} \sum_{b=1}^{B} (\hat{\xi}^{b} - \bar{\xi})^{2}}.$$

Then we have the bootstrap test statistics  $z^b$  ( $b = 1, 2, \dots, B$ ) as

$$z^{b} = \frac{\xi^{b} - \bar{\xi}}{\operatorname{se}(\xi)}.$$

We have the achieved significance level ASL:

$$ASL = \frac{\sum_{b=1}^{B} \lambda^{b}}{B},$$

where  $\lambda^b$  ( $b = 1, 2, \dots, B$ ) are the notations defined by

$$\lambda^{b} = \begin{cases} 1 & \text{if } |z^{b}| \ge |\hat{z}|, \\ 0 & \text{if } |z^{b}| < |\hat{z}| \\ (b = 1, 2, \dots, B). \end{cases}$$

For avoiding ASL = 0, ASL is also defined by

$$ASL = \frac{\sum_{b=1}^{B} \lambda^b + 1}{B+1},$$

when  $\sum_{b=1}^{B} \lambda^b < \varepsilon$  ( $\varepsilon$  is a sufficiently small positive number). We can say that the null hypothesis  $H_0$  ( $\xi = \hat{\xi}_c$ ) is rejected if *ASL* is less than or equal to the significance level.

In the cases of one-sample tests of  $\alpha$  and  $\beta$ , we adopt the following hypothesis:

null hypothesis 
$$H_0: \alpha = 0$$
,

alternative hypothesis  $H_1: \alpha \neq 0$ ,

for  $\alpha$ , and

null hypothesis  $H_0: \beta = 0$ ,

alternative hypothesis  $H_1: \beta \neq 0$ ,

for  $\beta$ .

One-sample tests by bootstrap resampling for  $\alpha$ ,  $\beta$ ,  $\xi$  in samples 1, 2 are shown in Table 6.

Table 6 One-sample test by bootstrap resampling

	sample 1 $(k = 1)$	sample 2 $(k=2)$		
$N_k$	120	80		
$B_k$	2000	2000		
$\hat{\alpha}_k$	6.2105	4.5209		
$ASL_k^{*1}$	0.0005	0.0050		
$\hat{\beta}_k$	-16.8720	-18.3394		
$ASL_k^{*2}$	0.0005	0.0095		
$\hat{\xi}_k$	0.3681	0.2465		
$ASL_k^{*3}$	-	0.0005		
min $I_{0.95}$	0.355	0.231		
max $I_{0.95}$	0.380	0.262		
*1 $H_0$ : $\alpha_k$	$g_{2}=0, H_{1}: \alpha_{k}\neq 0$	(k = 1, 2)		
*2 $H_0$ : $\beta_k$	$=0, \ H_1: \ \beta_k \neq 0$	(k = 1, 2)		
$^{*3}H_0: \xi_2$	$=\xi_1, \ H_1: \ \xi_2 \neq \xi_1$			
$\Lambda_k = \sum_{b=1}^{B_k} \lambda^b  (k = 1, 2)$				
$ASL_k = \frac{\Lambda}{B}$	$\frac{k+1}{k+1}$			

#### (iv) Confidence interval of threshold

The symbols of  $\hat{\xi}$ , se $(\hat{\xi})$ ,  $\hat{\xi}^b$  ( $b = 1, 2, \dots, B$ ),  $\bar{\xi}$  and se $(\hat{\xi})$  are the same as in (iii). Let  $\psi(z)$  be the cumulative distribution function of bootstrap resampling defined by

$$\psi(z) = \frac{1}{B} \sum_{b=1}^{B} \varphi^b(z) \quad (-\infty < z < +\infty),$$

where  $\varphi^{b}(z)$  ( $b = 1, 2, \dots, B$ ) are functions of z:

$$\varphi^{b}(z) = \begin{cases} 1 & \text{if } z \ge z^{b}, \\ 0 & \text{if } z < z^{b} \end{cases}$$
$$(b = 1, 2, \dots, B)$$

We note that  $\psi(z)$  satisfies

$$\begin{split} \psi(z) &\to 0 \quad (z \to -\infty), \\ \psi(z) &\to 1 \quad (z \to +\infty). \end{split}$$

Then we can obtain the confidence interval  $I_{\rho}$  of confidence coefficient  $\rho$  ( $0 < \rho < 1$ ) such that

$$I_{\rho}: \hat{\xi} - \psi^{-1}\left(\frac{1-\rho}{2}\right) \cdot \operatorname{se}(\xi) \leq \xi \leq \hat{\xi} + \psi^{-1}\left(\frac{1+\rho}{2}\right) \cdot \operatorname{se}(\xi).$$

The confidence intervals  $I_{0.95}$  of threshold  $\xi$  for samples 1 and 2 are shown in Table 6.

#### Concluding remarks

We proposed the bootstrap resampling algorithm for the psychophysical threshold estimates. Main properties of our algorithm are summarized in the following:

(i) the logistic regression including the guessing rate,

(ii) the non-parametric bootstrap resampling with log-likelihood ratio statistics for two-sample testing,

(iii) the non-parametric bootstrap resampling for one-sample testing to certify the values of parameters and threshold obtained by logistic regression.

We applied our bootstrap algorithm to the visual acuity test problem. Our algorithm does not require the identity of the number of observations between two samples. We can say that the bootstrap resampling provides a useful tool which has the flexibility of sampling in actual visual acuity measurements.

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