Details on Limited-information Goodness-of-fit Testing

To avoid notation clutter, we will spell out the details under dichotomous responses and mention how the theory can be extended to polytomous data. Recall the $C = 2^n$-cell multinominal table on which the IRT model likelihood is define. We shall first write the $C \times 1$ vector of cell probabilities as $\pi(\theta) = [\pi_1(\theta), \ldots, \pi_C(\theta)]'$. The true probabilities is given by $\pi_0 = (\pi_{01}, \ldots, \pi_{0C})'$. Correspondingly, the $C \times 1$ vector of observed proportions is $p = (p_1, \ldots, p_C)'$. Denote the $C \times 1$ vector of model-implied probabilities as $\hat{\pi} = \pi(\hat{\theta}) = (\hat{\pi}_1, \ldots, \hat{\pi}_C)'$ after obtaining $\hat{\theta}$ from the maximization of $L(\theta)$. The $C \times 1$ vector of cell residuals is given by $e = p - \hat{\pi}$. Standard discrete multivariate analysis results suggest that $e$ is asymptotically $C$-variate normally distributed under exactly correct model specification (see e.g., Bishop, Fienberg, & Holland, 1975)

$$
\sqrt{N}e = \sqrt{N}(p - \hat{\pi}) \overset{D}{\rightarrow} N_C(0, \Omega),
$$

where $\Omega = \Xi - \Delta F^{-1} \Delta'$. The multinominal covariance matrix $\Xi$ is equal to $\text{diag}(\pi) - \pi \pi'$, and the Jacobian matrix is

$$
\Delta = \frac{\partial \pi(\theta)}{\partial \theta'}.
$$

The Fisher information matrix is defined as $F = \Delta' [\text{diag}(\pi)]^{-1} \Delta$.

In $M_2$, the marginal residuals up to order 2 are used. For $n$ dichotomous items, there are $n$ first-order marginal residuals, and $n(n - 1)/2$ second-order marginal residuals. They are obtained via operator matrices. Let $\hat{L}$ be an $n \times C$ fixed matrix consisting of zeros and ones. The $(i, c)$th element in $\hat{L}$ is one if and only if item $i$ is endorsed in the $c$th response pattern. Similarly, let $\bar{L}$ be an $n(n - 1)/2 \times C$ fixed matrix of zeros and ones. Each row of this matrix corresponds to an item pair, and for each row, column $c$ is equal to one if and only if the pair of items is endorsed in the $c$th response pattern.
Pre-multiplying $\mathbf{e}$ by $\dot{\mathbf{L}}$ leads to $\dot{\mathbf{e}} = \dot{\mathbf{L}} \mathbf{e}$, the $n \times 1$ vector of first-order marginal residuals. Similarly, pre-multiplying $\mathbf{e}$ by $\ddot{\mathbf{L}}$ leads to $\ddot{\mathbf{e}} = \ddot{\mathbf{L}} \mathbf{e}$, the $n(n-1)/2 \times 1$ vector of second-order marginal residuals. Stacking $\dot{\mathbf{e}}$ and $\ddot{\mathbf{e}}$, the vector of marginal residuals up to order 2 is $\mathbf{e}_2 = (\dot{\mathbf{e}}, \ddot{\mathbf{e}})$. Because $\mathbf{e}_2$ is a linear function of $\mathbf{e}$, the asymptotic distribution of $\mathbf{e}_2$ is also normal

$$\sqrt{N} \mathbf{e}_2 \overset{D}{\rightarrow} N_{n(n+1)/2}(0, \mathbf{L}_2 \mathbf{\Omega}_2 \mathbf{L}_2')$$

where $\mathbf{L}_2 = (\dot{\mathbf{L}}, \ddot{\mathbf{L}})$, stacking $\dot{\mathbf{L}}$ and $\ddot{\mathbf{L}}$. Let us simplify the notation and let $\mathbf{\Omega}_2 = \mathbf{L}_2 \mathbf{\Omega}_2 \mathbf{L}_2' = \mathbf{\Xi}_2 - \Delta_2 \mathbf{F}^{-1} \Delta_2'$, where $\mathbf{\Xi}_2 = \mathbf{L}_2 \mathbf{\Xi}_2 \mathbf{L}_2'$ and $\Delta_2 = \mathbf{L}_2 \Delta$. Specifically, $\Delta_2$ is the matrix of derivatives of the first- and second order marginal probabilities with respect to $\theta$.

Now Maydeu-Olivares and Joe’s (2005) $M_2$ statistic can be defined:

$$M_2 = N \mathbf{e}_2' \left[ \hat{\mathbf{\Xi}}_2^{-1} - \hat{\mathbf{\Xi}}_2^{-1} \hat{\Delta}_2'(\hat{\Delta}_2' \hat{\Xi}_2^{-1} \hat{\Delta}_2)^{-1} \hat{\Delta}_2' \hat{\Xi}_2^{-1} \right] \mathbf{e}_2'$$

where $\hat{\Delta}_2$ and $\hat{\mathbf{\Xi}}_2$ denote the evaluation of $\Delta_2$ and $\mathbf{\Xi}_2$ at the maximum likelihood estimate $\hat{\theta}$. It follows from Proposition 4 in Browne’s (1984) that $M_2$ is asymptotically chi-square distributed with $n(n+1)/2 - \dim(\theta)$ degrees of freedom under the null hypothesis that the model fits exactly in the population.

For $n$ polytomous items with $K$ categories, the total number of possible response patterns becomes $C = K^n$. Consequently, the vectors of multinomial cell probabilities $\mathbf{\pi}(\theta)$ and $\hat{\mathbf{\pi}}$ is $C \times 1$ in size. In limited information goodness-of-fit testing, expanded operator matrices $\dot{\mathbf{L}}$ and $\ddot{\mathbf{L}}$ collapse cell probabilities into univariate and bivariate marginal probabilities or residual vectors, as detailed in Cai and Hansen (2013), among others. The asymptotic distribution theory of residuals under polytomous data can derived using along the same lines as Equation (2), enabling the formation of chi-square statistics similar to $M_2$ in Equation (3).
2 The Independence Model

Again, with no loss of generality, we illustrate the computations related to the full-independence model under dichotomous data. Recall that the 2PL model is

\[ P(U_i = 1|\eta) = \frac{1}{1 + \exp[-(\alpha + \beta'\eta)]}, \]

where \(\alpha\) is the intercept term and \(\beta\) a potentially vector-valued item slope parameter conformable with the dimensions of \(\eta\). For the complete-independence null model, no latent variable is present

\[ P(U_i = 1) = \frac{1}{1 + \exp(-\alpha)}. \] (4)

Note that \(P(U_i = 0) = 1 - P(U_i = 1)\).

Now let us turn to the computation of the \(M_2\) statistic for the independence model. Requirement components are the first- and second-order marginal probabilities, the Jacobian, and the multinomial covariance matrix. As a practical matter, the fit statistics for independence model are implemented in flexMIRT®(Cai, 2015). We note that \(M_2\) for independence model leads us to compute other relative fit indices such as the comparative fit index (CFI; Bentler, 1990), the normed-fit index (NFI; Bentler & Bonett, 1980), and the incremental fit index (IFI; Bollen, 1989) based on \(M_2\).

2.1 Marginal probabilities

Let us begin with the computation of the first- and second-order marginal probabilities. Again, without loss of generality, we only consider dichotomous items. Let there be \(n\) items scored \(U_i = 0\) or 1. Let \(\hat{\pi}_i\) denote the first order marginal probability for item \(i\) and \(\hat{\pi}_{ij}\) the second order marginal probability for item pair \((i, j)\). Accordingly, \(\hat{\pi}_i\) means the model-implied probability for item \(i\) when \(U_i = 1\). Likewise, \(\hat{\pi}_{ij}\) means the model-implied joint probability for items \(i\) and \(j\) when \(U_i = 1\) and \(U_j = 1\). We can compute \(\hat{\pi}_i\)
and \(\pi_{ij}\) directly as follows:

\[
\dot{\pi}_i = \int P(U_i = 1|\theta)g(\eta)\,d\eta, \tag{5}
\]

\[
\dot{\pi}_{ij} = \int P(U_i = 1|\theta)P(U_j = 1|\theta)g(\eta)\,d\eta. \tag{6}
\]

In our independence model, the first-order marginal probability when \(U_i = 1\) becomes

\[
\pi_i = \frac{1}{1 + \exp(-\alpha)} \int g(\eta)\,d\eta = \frac{1}{1 + \exp(-\alpha)}. \tag{7}
\]

It follows that \(\pi_i = \hat{p}_i\), where \(\hat{p}_i\) is the observed counterpart. This is due to the fact that intercepts perfectly fit the observed univariate proportions. Once this is established, the second-order marginal probabilities can be obtained simply as the products of two observed univariate proportions, i.e., \(\pi_{ij} = \hat{p}_i\hat{p}_j\).

2.2 The Jacobian

Next, the Jacobian, \(\Delta_2\), for the independence model can be expressed as

\[
\Delta_2 = \frac{\partial \pi(\hat{\theta})}{\partial \hat{\theta}'} - \frac{\partial \pi(\hat{\alpha})}{\partial \hat{\alpha}'} = \left(\frac{\partial \pi(\hat{\alpha})}{\partial \hat{\alpha}'} \right).
\]

Note that we only have to take the first-order derivatives of the model-implied marginal probabilities with respect to intercept. A typical elements in the upper block is

\[
\frac{\partial \pi_i}{\partial \alpha_i} = \frac{\partial P(U_i = 1)}{\partial \alpha_i} = \pi_i(1 - \pi_i), \tag{8}
\]

and those of the lower block are

\[
\frac{\partial \pi_{ij}}{\partial \alpha_i} = \frac{\partial P(U_i = 1)}{\partial \alpha_i}P(U_j = 1) = \pi_i(1 - \pi_i)\pi_j
\]

\[
\frac{\partial \pi_{ij}}{\partial \alpha_j} = \frac{\partial P(U_j = 1)}{\partial \alpha_j}P(U_i = 1) = \pi_j(1 - \pi_j)\pi_i. \tag{9}
\]
2.3 The multinomial covariance matrix

The multinominal covariance matrix $\Xi_2$ may be rewritten as (see Cai & Hansen, 2013):

$$\Xi_2 = \Sigma - \pi_2 \pi_2',$$

(10)

where $\Sigma = L_2 \text{diag}(\pi) L_2'$, and can be partitioned into $\Sigma_{11}$, $\Sigma_{21}$, and $\Sigma_{22}$ as follows:

$$\Sigma = \begin{pmatrix}
\Sigma_{11} \\
\Sigma_{21} \\
\Sigma_{22}
\end{pmatrix} = \begin{pmatrix}
L \text{diag}(\pi) L' \\
\bar{L} \text{diag}(\pi) \bar{L}' \\
\bar{L} \text{diag}(\pi) \bar{L}'
\end{pmatrix}$$

(11)

In short, calculating the elements of $\Sigma$ involves calculation of the first, second, third, and fourth order marginal probabilities. Because of full-independence, the model-implied probabilities for the third and fourth order margins can be obtained as products of univariate proportions.
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