

Fit Indices Versus Test Statistics

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Model evaluation is one of the most important aspects of structural equation modeling (SEM). Many model fit indices have been developed. It is not an exaggeration to say that nearly every publication using the SEM methodology has reported at least one fit index. Most fit indices are defined through test statistics. Studies and interpretation of fit indices commonly assume that the test statistics follow either a central chi-square distribution or a noncentral chi-square distribution. Because few statistics in practice follow a chi-square distribution, we study properties of the commonly used fit indices when dropping the chi-square distribution assumptions. The study identifies two sensible statistics for evaluating fit indices involving degrees of freedom. We also propose linearly approximating the distribution of a fit index/statistic by a known distribution or the distribution of the same fit index/statistic under a set of different conditions. The conditions include the sample size, the distribution of the data as well as the base-statistic. Results indicate that, for commonly used fit indices evaluated at sensible statistics, both the slope and the intercept in the linear relationship change substantially when conditions change. A fit index that changes the least might be due to an artificial factor. Thus, the value of a fit index is not just a measure of model fit but also of other uncontrollable factors. A discussion with conclusions is given on how to properly use fit indices.

In social and behavioral sciences, interesting attributes such as *stress*, *social support*, *socio-economic status* cannot be observed directly. They are measured by multiple indicators that are subject to measurement errors. By segregating measurement errors from the true scores of attributes, structural equation modeling (SEM), especially its special case of covariance structure analysis, provides a methodology for modeling the latent variables directly. Although there are many

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aspects to modeling, such as parameter estimation, model testing, and evaluating the size and significance of specific parameters, overall model evaluation is the most critical part in SEM. There is a huge body of literature on model evaluation that can be roughly classified into two categories: (a) overall-model-test statistics that judge whether a model fits the data exactly; (b) fit indices that evaluate the achievement of a model relative to a base model.

Fit indices and test statistics are often closely related. Actually, most interesting fit indices F s are defined through the so called chi-square statistics T s. The rationales behind these fit indices are often based on the properties of T . For example, under idealized conditions, T may approximately follow a central chi-square distribution under the null hypothesis and a noncentral chi-square distribution under an alternative hypothesis. In practice, data and model may not satisfy the idealized conditions and T may not follow (noncentral) chi-square distributions. Then, the rationales motivating these fit indices do not hold. There are a variety of studies on the performance of statistics; there also exist many studies on the performance of fit-indices. However, these two classes of studies are not well connected. For example, most of the studies on fit indices use just simulation with the normal theory based likelihood ratio statistic. There are also a few exceptions (e.g., Anderson, 1996; Hu & Bentler, 1998; Marsh, Hau, & Wen, 2004; Wang, Fan, & Willson, 1996; Zhang, 2004) but no study focused on the relationship between fit-indices and test statistics. This article will formally explore the relationship of the two. We are especially interested in conditions that affect the distributions of the commonly used fit indices. The purpose is to identify statistics that are most appropriate for calculating fit indices, to use fit indices more wisely and to evaluate models more scientifically.

We will use both analytical and empirical approaches to study various properties of fit indices. Our study will try to answer the following questions.

1. As point estimators, what are the population counterparts of the commonly used fit indices?
2. How the population counterparts related to model misspecifications?
3. Do we ever know the distribution of a fit index with real or even simulated data?
4. Are cutoff values such as 0.05 or 0.95 related to the distributions of the fit indices?
5. Are measures of model fit/misfit defined properly when the base-statistic does not follow a chi-square distribution? If not, can we have more sensible measures?
6. Whether confidence intervals for fit indices as printed in standard software cover the model fit/misfit with the desired probability?
7. How to reliably evaluate the power or sensitivity of fit indices?
8. Can we ever get an unbiased estimator of the population model fit/misfit as commonly defined?

Some of the questions have positive answers, some have negatives, and some may need further study. We will provide insightful discussions when definite answers are not available.

Although mean structure is an important part of SEM, in this article we will focus on covariance structure models due to their wide applications. In the next section, in order to facilitate the understanding of the development in later sections, we will give a brief review of the existing statistics and their properties, as well as fit indices and their rationales. We will discuss properties of fit indices under idealized conditions in the section entitled “Mean Values of Fit Indices Under Idealized Conditions.” Of course, idealized conditions do not hold in practice. In the section entitled “Approximating the Distribution of T Using a Linear Transformation,” we will introduce a linear transformation on the distribution of T to understand the difference between idealization and realization. With the help of the linear transformation, we will discuss the properties of fit indices in the section entitled “Properties of Fit Indices When T Does Not Follow a Chi-Square Distribution.” In the section entitled “Matching Fit Indices with Statistics,” we will match fit indices and statistics based on existing literature. An ad hoc correction to some existing statistics will also be given. The corrected statistics are definitionally more appropriate to define most fit indices. In the section entitled “Stability of Fit Indices When Conditions Change,” we discuss the sensitivity of fit indices to changes in other conditions besides model misspecification. Power issues related to fit indices will be discussed in the section entitled “The Power of a Fit Index.” In the Discussion section, we will discuss several critical issues related to measures of model fit and test statistics. We conclude the article by providing recommendations and pointing out remaining issues for further research.

SOME PROPERTIES OF STATISTICS AND RATIONALES FOR COMMONLY USED FIT INDICES

Let \mathbf{x} represent the underlying p -variate population from which a sample $\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N$ with $N = n + 1$ is drawn. We will first review properties of three classes of statistics. Then we discuss the rationales behind several commonly used fit-indices. This section will provide basic background information for later sections, where we discuss connections between fit indices and the existing statistics.

Statistics

The first class of statistics includes the normal theory likelihood ratio statistic and its rescaled version; the second one involves asymptotically distribution free statistics. These two classes are based on modeling the sample covariance matrix \mathbf{S} by a proposed model structure $\Sigma(\boldsymbol{\theta})$. The third class is based on robust proce-

dures which treat each observation \mathbf{x}_i individually instead of using the summary statistic \mathbf{S} .

The most widely utilized test statistic in SEM is the classical likelihood ratio statistic T_{ML} , based on the normal distribution assumption of the data. When data are truly normally distributed and the model structure is correctly specified, T_{ML} approaches a chi-square distribution χ_{df}^2 as the sample size N increases. Under certain conditions, this statistic asymptotically follows χ_{df}^2 even when data are not normally distributed (Amemiya & Anderson, 1990; Browne & Shapiro, 1988; Kano, 1992; Mooijaart & Bentler, 1991; Satorra, 1992; Satorra & Bentler, 1990; Yuan & Bentler, 1999b). Such a property is commonly called asymptotic robustness. However, procedures do not exist for verifying the conditions for asymptotic robustness. It seems foolish to blindly trust that T_{ML} will asymptotically follow χ_{df}^2 when data exhibit nonnormality. When data possess heavier tails than that of a multivariate normal distribution, the statistic T_{ML} is typically stochastically greater than the chi-square variate $\xi \sim \chi_{df}^2$. When the fourth-order moments of \mathbf{x} are all finite, the statistic T_{ML} can be decomposed into a linear combination of independent $\xi_j \sim \chi_1^2$. That is,

$$T_{ML} = \sum_{j=1}^{df} \kappa_j \xi_j + o_p(1),$$

where the κ_j s depend on the fourth-order moments of \mathbf{x} as well as the model structure, and $o_p(1)$ is a term that approaches zero in probability as sample size N increases. When \mathbf{x} follows elliptical or pseudo elliptical distributions with a common kurtosis κ , $\kappa_1 = \kappa_2 = \dots = \kappa_{df} = \kappa$. Then (see Browne, 1984; Shapiro & Browne, 1987; Yuan & Bentler, 1999b)

$$T_{ML} = \kappa \xi + o_p(1). \quad (1)$$

When a consistent estimator $\hat{\kappa}$ of κ is available, one can divide T_{ML} by $\hat{\kappa}$ so that the resulting statistic still asymptotically approaches χ_{df}^2 . Satorra and Bentler (1988) proposed $\hat{\kappa} = (\hat{\kappa}_1 + \dots + \hat{\kappa}_{df})/df$ and the resulting statistic

$$T_R = \hat{\kappa}^{-1} T_{ML},$$

is often referred to as the Satorra-Bentler rescaled statistic. Like T_{ML} , T_R can also follow a chi-square distribution when certain asymptotic robustness conditions are satisfied (Kano, 1992; Yuan & Bentler, 1999b). Simulation studies indicate that T_R performs quite robustly under a variety of conditions (Chou, Bentler, & Satorra, 1991; Curran, West, & Finch, 1996; Hu, Bentler, & Kano, 1992). However, data

generation in some of the studies is not clearly stated and may satisfy the asymptotic robustness condition for T_R (see Yuan & Bentler, 1999b). In general, T_R does not approach a chi-square distribution. Instead, it approaches a variate η with $E(\eta) = df$. It is likely that the distribution shape of η is far from that of a chi-square. In such cases, T_R will not behave like a chi-square. It can also lead to inappropriate conclusions when referring T_R to a chi-square distribution.

With typical nonnormal data in the social and behavioral sciences (Micceri, 1989), the ideal is to have a statistic that approximately follows a chi-square distribution regardless of the underlying distribution of the data. One of the original proposals in this direction was made by Browne (1984). His statistic is commonly called the asymptotically distribution free (ADF) statistic T_{ADF} , due to its asymptotically following χ_{df}^2 as long as \mathbf{x} has finite fourth-order moments. The ADF property is desirable. However, the distribution of T_{ADF} can be far from that of χ_{df}^2 for typical sample sizes encountered in practice (Hu et al., 1992). Specifically, the mean and variance of T_{ADF} are much greater than those of χ_{df}^2 . Most correctly specified models are rejected if using $T_{ADF} \sim \chi_{df}^2$. In an effort to find statistics that perform better in rejection rate with smaller N s, Yuan and Bentler (1997b) proposed a corrected statistic

$$T_{CADF} = T_{ADF}/(1 + T_{ADF}/n).$$

Like T_{ADF} , T_{CADF} asymptotically follows χ_{df}^2 as long as \mathbf{x} has finite fourth-order moments, thus, it is asymptotically distribution free. The mean of T_{CADF} approximately equals df for all sample sizes across various distributions (Yuan & Bentler, 1997b). However, at small sample sizes T_{CADF} over-corrects the behavior of T_{ADF} due to its rejection rate with correct models being smaller than the nominal level. Furthermore, T_{CADF} also carries the drawback of the ADF estimation method with nonconvergences at smaller sample sizes. In addition to T_{ADF} , Browne (1984) also proposed a residual-based ADF statistic T_{RADF} in which the estimator just needs to be consistent. However, T_{RADF} behaves almost the same as T_{ADF} , rejecting most correct models at smaller sample sizes. Parallel to T_{CADF} , Yuan and Bentler (1998b) proposed T_{CRADF} whose performance is almost the same as T_{CADF} , with its empirical mean approximately equal to df and under-rejecting the correct model for small sample sizes (Bentler & Yuan, 1999; Yuan & Bentler, 1998b).

The third class of statistics is obtained from robust procedures. It is well-known that the sample covariance matrix \mathbf{S} is very sensitive to influential observations and is biased for $\Sigma_0 = \text{Cov}(\mathbf{x})$ when data contain outliers (see Yuan & Bentler, 1998c). In such a situation, removing these outliers followed by modeling \mathbf{S} will lead to proper analysis of the covariance structure model. However, in a given data set, determining which cases are outliers may be difficult. The heavy tails, often indicated by larger marginal kurtoses or Mardia's (1970) multivariate kurtosis, might

be due to the heavy tails in the distribution of \mathbf{x} . When a data set possesses moderate heavy tails, \mathbf{S} will be an inefficient estimator of its population counterpart Σ_0 (Tyler, 1983). When the heavy tails are severe, the population 4th-order moments do not exist. Then modeling \mathbf{S} and referring T_{ML} , T_R , T_{ADF} , T_{RADF} , T_{CADF} or T_{CRADF} to χ_{df}^2 is meaningless. In such cases, it is better to model a robust covariance matrix \mathbf{S}_r . Since the empirical experimentations by Huba and Harlow (1987), various technical procedures have been developed for modeling robust covariance matrices (Yuan & Bentler, 1998a, 1998c, 2000; Yuan, Chan, & Bentler, 2000). They differ in how to control the weight attached to each case (see Yuan & Bentler, 1998a, 1998c; Yuan & Hayashi, 2003). Three types of statistics are proposed related to modeling \mathbf{S}_r . One is to model \mathbf{S}_r using the rescaled statistic T_R (Yuan & Bentler, 1998c); another is to just use T_{ML} treating \mathbf{S}_r as \mathbf{S} (Yuan et al., 2000; Yuan & Hayashi, 2003); the third is an ADF type statistic using the inverse of a robust version of the fourth-order moment matrix as the weight matrix (Yuan & Bentler, 1998a, 2000). The most preferred is T_{ML} coupling with a proper weight-control scheme (Yuan, Bentler, & Chan, 2004; Yuan & Hayashi, 2003). A brief outline as well as an introduction to a SAS IML program for robustifying a sample is provided in the appendix.

Note that many studies on T_{ML} , T_R , T_{ADF} and T_{CADF} in the first two classes only reported the rejection rates and their empirical means and standard deviations. Few studies examined their overall distributions (Curran, Bollen, Paxton, Kirby, & Chen, 2002; Yuan & Bentler, 1998b). By controlling the weights in some robust procedures, Yuan and Hayashi (2003) found that the distribution of T_{ML} applied to \mathbf{S}_r can be well described by χ_{df}^2 . But the distributions of T_R and T_{ADF} could not be well approximated by χ_{df}^2 even when data were fairly normally distributed.

In addition to the above three classes of statistics, other statistics such as the normal theory generalized least squares statistic T_{GLS} and the heterogeneous kurtosis statistic T_{HK} (Kano, Berkane, & Bentler, 1990) are also available in standard software (Bentler, 1995). These are related to the normal theory based statistics of the first class. Yuan and Bentler (1998b, 1999a) also proposed two F -statistics T_F and T_{FR} based on T_{ADF} and T_{RADF} , respectively. They are asymptotically distribution free and thus belong to the second class. We will not specifically deal with these statistics in this article although preliminary work indicates that the latter especially might be very valuable in testing the significance of a model.

Fit Indices

Fit indices can be classified into two categories, those that are defined explicitly through the overall test statistic T versus those that are not involving the statistic T directly. For example, the standardized root-mean-square residual (SRMR) is

not defined through T but through residuals at the convergence of a model fitting procedure. Actually, when a model is approximately correct, all the overall statistics can be approximately regarded as a sum of weighted squares of residuals (Shapiro, 1985), and the weights are optimized according to the chosen estimation procedure. Thus, those defined through a T better utilize the residuals than SRMR. Fit indices that are explicitly defined through T also fall into three types (e.g., Hu & Bentler, 1998; Marsh, Balla, & McDonald, 1988): (a) Indices that do not need the involved statistic to follow any known distribution; (b) indices that assume the statistic T to satisfy $E(T|H_0) = df$, which is automatically satisfied when $T \sim \chi_{df}^2$; and (c) indices that assume $(T|H_0) \sim \chi_{df}^2$ and $(T|H_1) \sim \chi_{df}^2(\delta)$, where H_0 represents the null hypothesis, H_1 represents the alternative hypothesis and δ is the noncentrality parameter (NCP). Many fit indices are available in standard software (e.g., EQS, LISREL, SAS CALIS). The commonly reported ones are CFI, GFI, NFI, NNFI, and RMSEA, according to the review of McDonald and Ho (2002). We will mainly discuss the commonly used ones although our analysis and discussion also equally apply to other fit indices (e.g., Bollen, 1986, 1989; Hoelter, 1983; Steiger, 1989).

Let T_M and T_I be the chosen statistic T evaluated at the substantive model $\Sigma_M = \Sigma(\theta)$ and the independence model $\Sigma_I = \text{diag}(\sigma_{11}, \sigma_{22}, \dots, \sigma_{pp})$, respectively. The normed fit index (Bentler & Bonett, 1980) is defined as

$$\text{NFI} = 1 - \frac{T_M}{T_I}.$$

As discussed by Hu and Bentler (1998), the T in NFI does not necessarily need to follow a particular distribution. Another widely used fit index is the nonnormed fit index (Bentler & Bonett, 1980; Tucker & Lewis, 1973)

$$\text{NNFI} = 1 - \frac{(T_M / df_M - 1)}{(T_I / df_I - 1)},$$

where df_M and df_I are the degrees of freedom in model Σ_M and Σ_I , respectively. The difference between NFI and NNFI is that T is replaced by $T/df - 1$, which is motivated by $(T|H_0) \sim \chi_{df}^2$ (see Bentler, 1995). When Σ_M is correctly specified, $E(T_M) = df_M$ and $\text{NNFI} \approx 1$. A more popular fit index is the comparative fit index (Bentler, 1990)

$$\text{CFI} = 1 - \frac{\max[(T_M - df_M), 0]}{\max[(T_I - df_I), (T_M - df_M)]}.$$

CFI is always within the range of [0,1]. Essentially equivalent to CFI but not necessarily within the range of [0,1] is the relative noncentrality index

$$\text{RNI} = 1 - \frac{T_M - df_M}{T_I - df_I},$$

independently proposed by Bentler (1990) and McDonald and Marsh (1990). Both CFI and RNI are motivated by $(T | H_1) \sim \chi_{df}^2(\delta)$ so that they measure the reduction of the NCP by Σ_M relative to that by Σ_I (see Bentler, 1990, 1995). Another commonly used index is the goodness of fit index, its general form was given in Bentler (1983, Equation 3.5) as

$$\text{GFI} = 1 - \frac{\mathbf{e}'\mathbf{W}\mathbf{e}}{\mathbf{s}'\mathbf{W}\mathbf{s}},$$

where $\mathbf{e} = \mathbf{s} - \boldsymbol{\sigma}(\hat{\boldsymbol{\theta}})$ with \mathbf{s} and $\boldsymbol{\sigma}$ being vectors containing the nonduplicated elements of \mathbf{S} and $\boldsymbol{\Sigma}$, respectively; the weight matrix \mathbf{W} is subject to the estimation method. We can rewrite

$$\text{GFI} = 1 - \frac{T_M}{T_0},$$

where $T_M = \mathbf{ne}'\mathbf{W}\mathbf{e}$ is the normal theory based iteratively reweighted least squares (IRLS) statistic T_{IRLS} when the normal theory based \mathbf{W} is used (see Bentler, 1995, p. 216) and $T_0 = \mathbf{s}'\mathbf{W}\mathbf{s}$ can also be regarded as a T_{IRLS} for testing $\boldsymbol{\Sigma}_0 = \mathbf{0}$. When the ADF weight matrix \mathbf{W} is used, T_M and T_0 share similar meanings (Tanaka & Huba, 1985).

The last fit index we will discuss is the root-mean-square error of approximation (Browne & Cudeck, 1993; Steiger & Lind, 1980)

$$\text{RMSEA} = \sqrt{\max[(T_M - df_M)/(n \times df_M), 0]}.$$

The implicit assumption in RMSEA is that $(T | H_1) \sim \chi_{df}^2(\delta)$ and δ equals n times the “model misfit” so that it estimates the model misfit per degree of freedom. Notice that RMSEA only involves T_M . The literature on fit indices classifies it as an absolute fit index while those involving T_I are relative fit indices. There are also many other fit indices (see Hu & Bentler, 1998) in addition to the above six, but they are less frequently used.

MEAN VALUES OF FIT INDICES UNDER IDEALIZED CONDITIONS

A fit index F may aim to estimate a quantity measuring the model fit/misfit at the population level. F is always an unbiased estimate of $E(F)$ when the expectation exists. We need to consider how $E(F)$ is related to model fit/misfit. Analytically, this can only be studied when the involved statistics follow known distributions. Suppose

$$T_M \sim \chi_{df_M}^2(\delta_M) \text{ and } T_I \sim \chi_{df_I}^2(\delta_I). \tag{2}$$

Then

$$E(T_M) = df_M + \delta_M \text{ and } E(T_I) = df_I + \delta_I, \tag{3}$$

and

$$\hat{\delta}_M = T_M - df_M \text{ and } \hat{\delta}_I = T_I - df_I,$$

are the unbiased estimators of δ_M and δ_I , respectively. Most fit indices are functions of $\hat{\delta}_M$ and $\hat{\delta}_I$. It is interesting to see how the mean of a fit index is related to δ_M and δ_I . Suppose the fit index F is given by

$$F = g(\hat{\delta}_M, \hat{\delta}_I).$$

Because g is typically a nonlinear function, we might use a Taylor expansion to facilitate the study. Denote \ddot{g}_{ij} as the second derivatives of g with respect to the parameters indicated in the subscripts. We have

$$E[g(\hat{\delta}_M, \hat{\delta}_I)] = g(\delta_M, \delta_I) + E[\ddot{g}_{11}(\bar{\delta}_M, \bar{\delta}_I)(\hat{\delta}_M - \delta_M)^2 / 2] + E[\ddot{g}_{22}(\bar{\delta}_M, \bar{\delta}_I)(\hat{\delta}_I - \delta_I)^2 / 2] + E[\ddot{g}_{12}(\bar{\delta}_M, \bar{\delta}_I)(\hat{\delta}_M - \delta_M)(\hat{\delta}_I - \delta_I)], \tag{4}$$

where $\bar{\delta}_M$ is a number between δ_M and $\hat{\delta}_M$, and $\bar{\delta}_I$ is a number between δ_I and $\hat{\delta}_I$. Most incremental fit indices can be expressed as $g(t_1, t_2) = 1 - (t_1 - c_1)/(t_2 - c_2)$, where c_1 and c_2 are constants. Then $\ddot{g}_{11} = 0, \ddot{g}_{22} = -2(t_1 - c_1)/(t_2 - c_2)^3$ can be either negative or positive, $\ddot{g}_{12} = 1/(t_2 - c_2)^2 > 0$. It follows from Equation 4 that $E[g(\hat{\delta}_M, \hat{\delta}_I)]$ can be greater than $g(\delta_M, \delta_I)$ or smaller than $g(\delta_M, \delta_I)$,

depending on the last two terms in Equation 4. It is unlikely for $E[g(\hat{\delta}_M, \hat{\delta}_I)]$ to equal $g(\delta_M, \delta_I)$.

For indices that are only based on $\hat{\delta}_M$, there exists

$$E[g(\hat{\delta}_M)] = g(\delta_M) + E\left[\ddot{g}(\bar{\delta}_M)(\hat{\delta}_M - \delta_M)^2 / 2\right]. \tag{5}$$

When g is a strictly concave function, $\ddot{g}(\bar{\delta}_M) < 0$ and it follows from Equation 5 that $E[g(\hat{\delta}_M)] < g(\delta_M)$. When g is a convex function, the opposite is true. For the index Mc proposed in McDonald (1989) for example, we have $g(t) = \exp[-t/(2n)]$, $\ddot{g}(t) = 0.25n^{-2} \exp[-t/(2n)]$, and

$$E(Mc) > \exp[-\delta_M/(2n)].$$

The difference between $E(Mc)$ and $\exp[-\delta_M/(2n)]$ is proportional to the variance of T_M . The widely used RMSEA is a composite function $g(\hat{\delta}_M) = g_1[g_2(\hat{\delta}_M)]$, where $g_1(t) = \sqrt{t}$ is strictly concave with $\ddot{g}_1(t) = -0.25t^{-3/2} < 0$; and $g_2(t) = \max[t/(n \times df_M), 0]$ is convex but not strictly. For bad models, T_M may never be below df_M so that $g_2(t)$ becomes an identity function, then

$$E(\text{RMSEA}) < \sqrt{\delta_M / (n \times df_M)}.$$

For correct or nearly correct models, $E(\text{RMSEA})$ might be greater than $\sqrt{\delta_M / (n \times df_M)}$ due to the convexity of $g_2(t)$ (see Raykov, 2000).

Note that the assumptions in Equation 2 are unlikely to be true in practice. The above analysis implies that we do not know the mean values of the commonly used fit indices even under idealized conditions. Cutoff values such as 0.05 or 0.95 have little to do with the mean value of the fit indices or the NCPs of the idealized chi-squares. Of course, the chosen overall test statistic T maybe far from following a chi-square distribution. When a noncentral chi-square does not exist, then NCP is irrelevant. The rest of the article will discuss different aspects of F and T .

APPROXIMATING THE DISTRIBUTION OF T USING A LINEAR TRANSFORMATION

As reviewed in the previous section, Equations 2 and 3 are widely used in the context of fit indices (Bentler, 1990; Browne & Cudeck, 1993; Kim, 2003; MacCallum, Browne, & Sugawara, 1996; McDonald, 1989; McDonald & Marsh, 1990; Ogasawara, 2001; Steiger & Lind, 1980). Results in Equation 2 might be

justified by the asymptotic result (see e.g., Bentler & Dijkstra, 1985; Browne, 1984; Satorra, 1989; Shapiro, 1983)

$$T = nD[\mathbf{S}, \boldsymbol{\Sigma}(\hat{\boldsymbol{\theta}})] \xrightarrow{\mathcal{L}} \chi_{df}^2(\delta), \tag{6}$$

where $\hat{\boldsymbol{\theta}}$ is the parameter estimator, $\delta = n\tau$ and

$$\tau = D[\boldsymbol{\Sigma}_0, \boldsymbol{\Sigma}(\boldsymbol{\theta}^*)] = \min_{\boldsymbol{\theta}} D[\boldsymbol{\Sigma}_0, \boldsymbol{\Sigma}(\boldsymbol{\theta})], \tag{7}$$

is the “model misfit” measured by a discrepancy function $D(\cdot, \cdot)$. Actually, Equation 6 is not true¹ with a given $\boldsymbol{\Sigma}_0$ (see Olsson, Foss, & Breivik, 2004; Yuan & Bentler, in press; Yuan & Chan, in press). The proof of Equation 6 needs the assumption

$$E(\mathbf{S}) = \boldsymbol{\Sigma}_0 = \boldsymbol{\Sigma}_{0,n} = \boldsymbol{\Sigma}(\boldsymbol{\theta}^*) + \frac{\boldsymbol{\Delta}}{\sqrt{n}}, \tag{8}$$

where $\boldsymbol{\Delta}$ is a constant matrix. Note that \mathbf{S} contains sampling error ($\mathbf{S} - \boldsymbol{\Sigma}_0$) whose magnitude is of order $1/\sqrt{n}$. One might understand Equation 8 by thinking that the systematic error $\boldsymbol{\Delta}/\sqrt{n}$ approximately equals the sampling error. However, $\boldsymbol{\Sigma}_0$ is the population covariance matrix that should not depend on n while Equation 8 implies that the amount of misspecification in $\boldsymbol{\Sigma}(\boldsymbol{\theta})$ decreases as n increases. Such a condition has an obvious fault but it provides the mathematical convenience to allow one to show that Equation 6 holds with δ being a constant that does not depend on n . With fixed $\boldsymbol{\Sigma}_0$ and n , what the asymptotic statistical theory really tells us is that, when $\boldsymbol{\Sigma}_0$ is sufficiently close to $\boldsymbol{\Sigma}(\boldsymbol{\theta})$, the distribution of T can be approximated by $\chi_{df}^2(\delta)$. But it says nothing about the goodness of the approximation.

Let’s look at the T_{ML} with normally distributed data generated by the following confirmatory factor model (CFM)

$$\mathbf{x} = \boldsymbol{\mu} + \boldsymbol{\Lambda}\mathbf{f} + \mathbf{e} \quad \text{with} \quad \text{Cov}(\mathbf{x}) = \boldsymbol{\Lambda}\boldsymbol{\Phi}\boldsymbol{\Lambda}' + \boldsymbol{\Psi},$$

$$\boldsymbol{\Lambda} = \begin{pmatrix} \boldsymbol{\lambda} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \boldsymbol{\lambda} & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \boldsymbol{\lambda} \end{pmatrix}, \quad \boldsymbol{\Phi} = \begin{pmatrix} 1.0 & .30 & .40 \\ .30 & 1.0 & .50 \\ .40 & .50 & 1.0 \end{pmatrix}, \tag{9}$$

where $\boldsymbol{\lambda}' = (.70, .70, .75, .80, .80)$ and $\boldsymbol{\Psi}$ is a diagonal matrix chosen so that $\boldsymbol{\Sigma}_0$ is a correlation matrix. Suppose one fits the sample covariance matrix \mathbf{S} by the independence model $\boldsymbol{\Sigma}_I$ using the normal theory maximum likelihood (ML). Then $\tau = 6.859$. With 500 replications, the top panel of Figure 1 plots the quantiles of the sta-

¹Curran et al.’s (2002) conclusion that Equation 6 approximately holds in some of their simulation conditions is based on 500 or few converged samples out of 650 replications. A different conclusion might be reached when all samples converge.

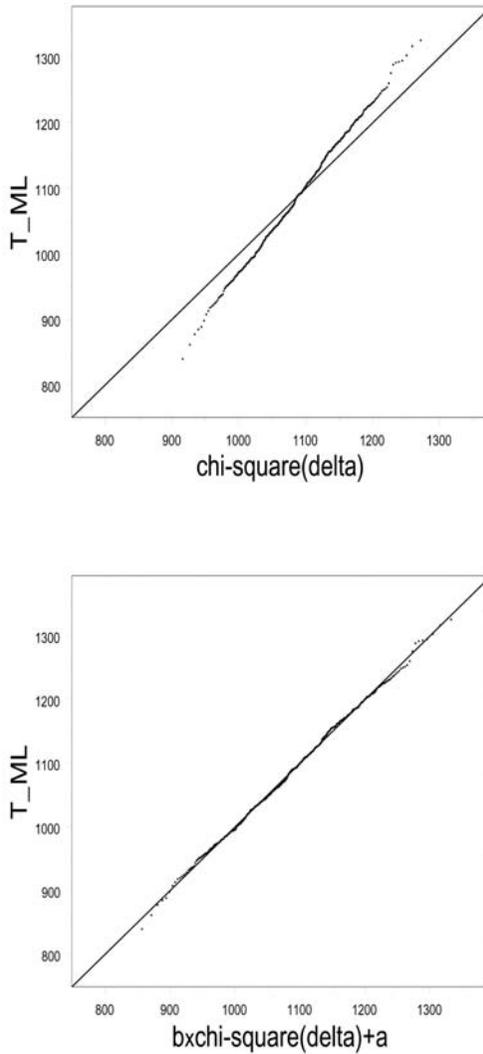


FIGURE 1 QQ plots of T_{ML} versus $\chi^2_{105}(\delta)$ and T_{ML} versus $b\chi^2_{105}(\delta) + a$ with $a = -368.000$ and $b = 1.337$, $N = 150$ and 500 replications.

tistic T_{ML} against the quantiles of $\chi^2_{105}(\delta)$ (QQ plot) at $N = 150$. It is obvious that the upper tail of the distribution of T_{ML} is much longer than that of $\chi^2_{105}(\delta)$ and the lower tail is much shorter than that of $\chi^2_{105}(\delta)$.

Motivated by Equations 1 and 3, we might describe the distribution of T by

$$T = b\chi^2_{df}(\delta) + a, \tag{10}$$

where the intercept a and slope b can be estimated by regression of the quantiles of T against those of $\chi_{df}^2(\delta)$. The bottom panel of Figure 1 plots the quantiles of T_{ML} against those of $b\chi_{105}^2(\delta) + a$ with $a = -368.000$ and $b = 1.337$ being estimated based on the 500 replications. It is obvious that Equation 10 describes the distribution of T_{ML} very well. Actually, judged by a visual inspection, the two sets of quantiles in the bottom panel of Figure 1 match as close as a QQ plot (not presented here) of a sample of simulated chi-squares against its population quantiles. When each of the samples is fitted by the 3-factor model as that generated the sample in Equation 9, then H_0 holds with $df = 87$. Figure 2 contains the QQ plots of T_{ML} for the correct model with 500 simulated normal samples of size $N = 150$. It is obvious that T_{ML} is stochastically greater than χ_{87}^2 and is described by $b\chi_{87}^2 + a$ very well. Our limited evidence also implies that $b\chi_{df}^2(\delta) + a$ can also describe the distributions of the other statistics quite well. Notice that a and b depend on the involved statistic, the sample size, the distribution of the data, as well as the model itself.

Transformation Equation 10 can be extended to

$$(T_2|C_2) = b(T_1|C_1) + a, \quad (11)$$

where T_2 and T_1 can be two different statistics or the same statistic evaluated at two different conditions C_1 and C_2 . The conditions include the sample size, the distribution of \mathbf{x} , and the model specification. Equation 11 is quite useful in describing the distributional change of T when changing conditions. Of course, we would wish to have $b = 1$, $a = 0$ regardless of T_1 , C_1 and T_2 , C_2 . However, a and b change with the sample size and the distribution of \mathbf{x} even when T_1 and T_2 are the same statistic. To provide more information on a and b under different conditions, we let T_1 and T_2 be the same statistic while C_1 and C_2 have different sample sizes or distributions of \mathbf{x} . Using the same CFM as in Equation 9, the distribution of \mathbf{x} in C_2 changes from elliptically distributed data to lognormal factors and errors, as further clarified in Table 1. The parameters are estimated by the ML. The statistics presented in Table 2 are T_{ML} , T_R , T_{RADF} , T_{CRADF} , as reviewed in the section entitled "Some Properties of Statistics and Rationales for Commonly Used Fit Indices." The statistics T_{AML} and T_{AR} are ad hoc corrections respectively to T_{ML} and T_R , which will be discussed further in a later section. The T_{IRLS} here is for the study of the properties of GFI. The T_0 corresponding to T_{RADF} is also for the properties of GFI, where \mathbf{W} is just the ADF weight matrix due to the model $\boldsymbol{\sigma} = \mathbf{0}$ having all its derivatives equal to zero (see Browne, 1984). With 500 replications, Table 2 contains the a and b at $N = 150$ while the distribution of \mathbf{x} changes; Table 3 contains the parallel results corresponding to $N = 500$. It is obvious that a and b change for all the statistics, especially at the misspecified independence model. A larger sample size does not help much even for the ADF type statistics. Table 4 contains similar result when T_1 and T_2 are the same statistic and C_1 and C_2 have the same distribution but different sample sizes. Compared to the distributional variation condition, sample size has a rela-

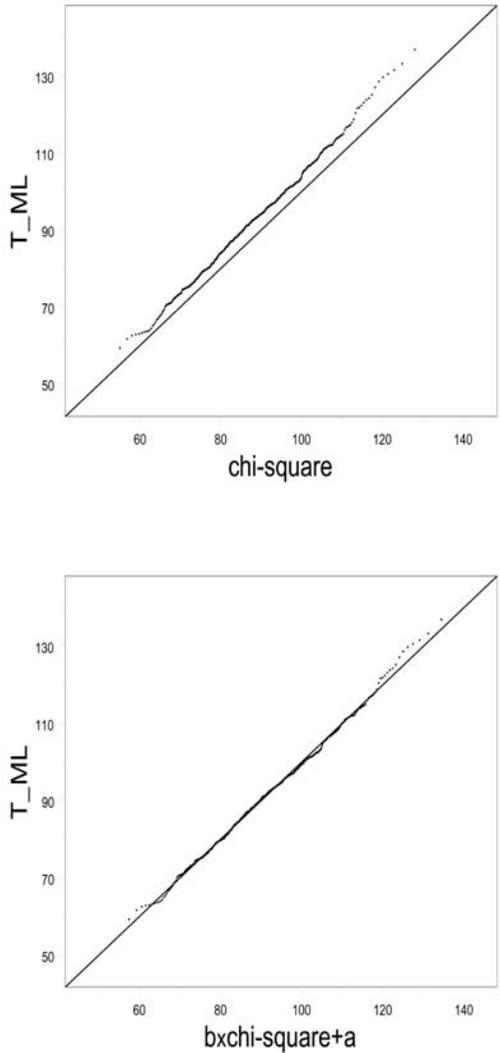


FIGURE 2 QQ plots of T_{ML} versus χ_{87}^2 and T_{ML} versus $b\chi_{87}^2 + a$ with $a = -0.555$ and $b = 1.054$, $N = 150$ and 500 replications.

tively smaller effect on b . But there is still a substantial difference among the a s, especially when the model is misspecified.

Notice that, for all the simulation conditions in Table 1, T_R asymptotically follows χ_{87}^2 for the correctly specified model. Actually, Equation 1 holds with $\kappa = 3$ for all the non-normal distribution conditions in Table 1. A greater κ will have a

TABLE 1
Distribution Conditions of \mathbf{x}

Normal \mathbf{x}	$\mathbf{x} = \boldsymbol{\mu} + \Lambda\mathbf{f} + \mathbf{e}, \mathbf{f} \sim N(\mathbf{0}, \Phi), \mathbf{e} \sim N(\mathbf{0}, \Psi)$
Elliptical \mathbf{x}	$\mathbf{x} = \boldsymbol{\mu} + (\Lambda\mathbf{f} + \mathbf{e})/r, \mathbf{f} \sim N(\mathbf{0}, \Phi), \mathbf{e} \sim N(\mathbf{0}, \Psi), r \sim \sqrt{\chi_5^2/3}$
Skew \mathbf{f} & normal \mathbf{e}	$\mathbf{x} = \boldsymbol{\mu} + (\Lambda\mathbf{f} + \mathbf{e})/r, \mathbf{f} \sim \text{Lognormal}(\mathbf{0}, \Psi), \mathbf{e} \sim N(\mathbf{0}, \Phi), r \sim \sqrt{\chi_5^2/3}$
Skew \mathbf{f} & \mathbf{e}	$\mathbf{x} = \boldsymbol{\mu} + (\Lambda\mathbf{f} + \mathbf{e})/r, \mathbf{f} \sim \text{Lognormal}(\mathbf{0}, \Phi), \mathbf{e} \sim \text{Lognormal}(\mathbf{0}, \Psi), r \sim \sqrt{\chi_5^2/3}$

Note. \mathbf{e}, \mathbf{f} , and r are independent; $\mathbf{f} \sim \text{Lognormal}(\mathbf{0}, \Phi)$ is obtained by $\mathbf{f} = \Phi^{1/2} \boldsymbol{\xi}$, where $\boldsymbol{\xi} = (\xi_1, \xi_2, \xi_3)'$ and ξ_j is to standardize $\exp(z)$ with $z \sim N(0, 1)$; and similarly for $\mathbf{e} \sim \text{Lognormal}(\mathbf{0}, \Psi)$.

TABLE 2
Intercept a and Slope b in $(T_2|C_2) = b(T_1|C_1) + a$, Where T_2 and T_1 Are the Same Statistic, C_1 and C_2 Have the Same Sample Size $N = 150$, C_1 Has Normally Distributed Data

T	C_2	Elliptical \mathbf{x}		Skew \mathbf{f} & Normal \mathbf{e}		Skew \mathbf{f} & \mathbf{e}	
		b	a	b	a	b	a
T_{ML}	T_M	3.166	-117.177	3.132	-113.210	3.347	-152.676
	T_I	1.492	-462.592	4.103	-3471.507	4.652	-3923.917
T_{AML}	T_M	3.166	-111.541	3.132	-107.765	3.347	-145.332
	T_I	1.492	-444.481	4.103	-3335.598	4.652	-3770.297
T_R	T_M	0.884	10.059	0.948	6.126	1.058	4.273
	T_I	1.472	-941.064	1.001	-725.069	1.231	-920.951
T_{AR}	T_M	0.884	9.575	0.948	5.831	1.058	4.067
	T_I	1.472	-904.222	1.001	-696.683	1.231	-884.896
T_{RADF}	T_M	0.799	38.717	0.825	31.263	0.710	54.620
	T_I	0.717	33.396	0.466	38.273	0.571	6.968
	T_0	0.317	235.940	0.312	201.524	0.239	-2.411
T_{CRADF}	T_M	0.884	9.361	0.911	6.666	0.806	15.779
	T_I	1.141	-23.544	1.477	-76.333	1.406	-63.747
T_{IRLS}	T_M	4.217	-201.608	4.107	-194.497	3.688	-186.975
	T_0	3.304	-2676.005	2.876	-2170.028	2.824	-2173.540

greater effect on a and b corresponding to T_{ML} in Tables 2 and 3. When Equation 1 does not hold or when κ changes from sample to sample, we would have more changes on a and b corresponding to T_{ML}, T_{AML}, T_R and T_{AR} . Of course, the a and b are also model dependent. When a different model other than that in Equation 9 is used, we will observe different patterns on a and b . Furthermore, a and b can only be estimated empirically. The estimators obtained from one sample may not be applied to a different sample for the purpose of correcting the performance of statistics. But Equations 10 and 11 do provide a simple procedure for comparing two distributions.

TABLE 3
 Intercept a and Slope b in $(T_2|C_2) = b(T_1|C_1) + a$, Where T_2 and T_1 Are the Same Statistic, C_1 and C_2 Have the Same Sample Size $N = 500$, C_1 Has Normally Distributed Data

T	C_2	<i>Elliptical x</i>		<i>Skew f & Normal e</i>		<i>Skew f & e</i>	
		b	a	b	a	b	a
T_{ML}	T_M	4.580	-207.985	4.621	-214.494	3.980	-176.997
	T_I	1.536	-1783.119	4.855	-13638.862	5.291	-14887.005
T_{AML}	T_M	4.580	-204.998	4.621	-211.413	3.980	-174.455
	T_I	1.586	-1762.275	4.855	-13479.424	5.291	-14712.976
T_R	T_M	0.863	10.916	0.855	11.533	0.976	10.939
	T_I	2.097	-5135.660	1.438	-3916.342	1.620	-4440.145
T_{AR}	T_M	0.863	10.759	0.855	11.368	0.976	10.782
	T_I	2.097	-5075.624	1.438	-3870.560	1.620	-4388.239
T_{RADF}	T_M	0.899	11.116	0.876	13.179	0.725	28.562
	T_I	0.755	-120.164	0.430	-136.305	0.476	-163.075
	T_0	0.289	-259.937	0.285	-293.929	0.162	-254.850
T_{CRADF}	T_M	0.898	9.265	0.881	10.447	0.732	23.084
	T_I	1.337	-165.491	1.463	-302.667	1.557	-326.911
T_{IRLS}	T_M	6.124	-371.420	6.309	-395.120	4.496	-258.577
	T_0	2.951	-7377.125	2.432	-5408.609	2.182	-4572.810

In summary, chi-square distributions are generally not achievable even when data are normally distributed. Distribution shapes of the commonly used statistics vary substantially when conditions such as the sample size, the distribution of \mathbf{x} , model size and model misspecification change. The results suggest that using (noncentral) chi-square distributions of T to describe the properties of a fit index F is inappropriate. In the following section we will discuss the properties of fit indices when T does not follow a chi-square distribution but can be approximated by Equations 10 or 11.

Note that all robust procedures are tailor-made. One has to properly choose the weight function in order to obtain the estimator or test statistic. For a given sample, one can find a proper weighting scheme so that $(T_{ML}|H_0)$ applied to the robustified sample approximately follows χ_{df}^2 (Yuan & Hayashi, 2003). When T_{ML} approximately follows the same distribution across two robustified samples, then we will approximately have $a = 0$ and $b = 1$. But even if $(T_{ML}|H_0)$ approximately follows χ_{df}^2 , this does not imply that $(T_{ML}|H_1)$ will approximately follow $\chi_{df}^2(\delta)$. We will further discuss whether $T \sim \chi_{df}^2(\delta)$ is achievable or necessary when interpreting fit indices involving T .

TABLE 4
 Intercept a and Slope b in $(T_2|C_2) = b(T_1|C_1) + a$, Where T_2 and T_1 Are the Same Statistic and C_1 and C_2 Have the Same Distribution, but C_1 Has $N = 150$ and C_2 Has $N = 500$

T	$C_1 \& C_2$	<i>Normal x</i>		<i>Elliptical x</i>		<i>Skew f & Normal e</i>		<i>Skew f & e</i>	
		b	a	b	a	b	a	b	a
T_{ML}	T_M	1.032	-5.963	1.599	-78.382	1.659	-92.813	1.299	-24.126
	T_I	1.830	1464.784	1.890	1333.560	2.077	1095.487	2.031	1101.599
T_{AML}	T_M	1.069	-5.877	1.656	-77.256	1.717	-91.480	1.345	-23.779
	T_I	1.883	1447.661	1.944	1317.970	2.136	1082.681	2.089	1088.722
T_R	T_M	1.008	-5.148	0.984	-3.406	0.910	1.476	0.932	1.712
	T_I	1.747	1358.086	2.514	37.271	2.502	-139.501	2.305	-124.598
T_{AR}	T_M	1.044	-5.074	1.019	-3.357	0.942	1.455	0.965	1.687
	T_I	1.797	1342.210	2.586	36.835	2.573	-137.871	2.371	-123.141
T_{RADF}	T_M	0.327	29.938	0.367	24.015	0.346	28.761	0.333	32.408
	T_I	0.300	526.297	0.311	270.135	0.277	79.972	0.250	85.748
	T_0	0.240	3827.699	0.220	793.140	0.221	750.715	0.164	363.810
T_{CRADF}	T_M	1.576	-54.287	1.585	-53.106	1.515	-46.771	1.416	-37.834
	T_I	2.437	-3.535	2.866	-103.960	2.399	-121.866	2.693	-159.363
T_{IRLS}	T_M	1.094	-1.971	1.824	-103.984	2.035	-141.218	1.402	-27.787
	T_0	2.589	735.523	2.342	942.115	2.328	954.490	1.913	1484.340

PROPERTIES OF FIT INDICES WHEN T DOES NOT FOLLOW A CHI-SQUARE DISTRIBUTION

To facilitate the discussion we use

$$T = bt + a, \tag{12}$$

where t might have a known distribution as in Equation 10 or might be the distribution of a given statistic under the normal sampling scheme with a fixed sample size, as in Equation 11. We denote the a and b at the independence model as a_I, b_I and at the substantive model as a_M and b_M , respectively. We will mainly discuss the stability of fit indices when the underlying distribution of \mathbf{x} changes. Due to the uncritical use of T_{ML} in the practice of SEM, we will pay special attention to the properties of fit indices associated with T_{ML} . We cannot predict a_I, a_M or b_I, b_M in general. But in the special case when Σ_I is approximately correct, we might approximately have $a_I = a_M = 0$ and $b_I = b_M = b$ within the class of elliptical distributions, which will be specifically mentioned when a nice property holds.

The normed fit index NFI can be written as

$$\text{NFI} = 1 - \frac{b_M t_M + a_M}{b_I t_I + a_I},$$

which is invariant to $b_I = b_M = b$ and $a_I = a_M = 0$. When model Σ_I is approximately correct and $T = T_{ML}$, there may approximately exist $b_I = b_M = b$ and $a_I = a_M = 0$ within the class of elliptical distributions. However, when the off diagonal of Σ_0 cannot be ignored, $b_I \neq b_M$ and a_I or a_M will not equal zero either. The distribution of NFI will change when \mathbf{x} changes distributions or when Σ_M changes. Notice that the change happens not just in the mean value $E(\text{NFI})$ but also the distributional form of NFI. When n tends towards infinity,

$$\text{NFI} = 1 - \frac{\min_{\boldsymbol{\theta}} D[\mathbf{S}, \Sigma_M(\boldsymbol{\theta})]}{\min_{\boldsymbol{\theta}} D[\mathbf{S}, \Sigma_I(\boldsymbol{\theta})]} \rightarrow 1 - \frac{\min_{\boldsymbol{\theta}} D[\Sigma_0, \Sigma_M(\boldsymbol{\theta})]}{\min_{\boldsymbol{\theta}} D[\Sigma_0, \Sigma_I(\boldsymbol{\theta})]}, \quad (13)$$

which measures the reduction (explanation) of all the covariances by Σ_M relative to Σ_I . However, the limit in Equation 13 still depends on the measure $D(\cdot, \cdot)$ of distance (see La Du & Tanaka, 1989, 1995; Sugawara & MacCallum, 1993; Tanaka, 1987) unless the model Σ_M is correctly specified (Yuan & Chan, in press). In practice, N is always finite,

$$E(\text{NFI}) \neq 1 - \frac{\min_{\boldsymbol{\theta}} D[\Sigma_0, \Sigma_M(\boldsymbol{\theta})]}{\min_{\boldsymbol{\theta}} D[\Sigma_0, \Sigma_I(\boldsymbol{\theta})]},$$

in general. Furthermore, the speed of convergence to the limit in Equation 13 might be very slow. So interpreting NFI according to its limit is not appropriate.

Very similar to NFI is GFI. Using the notation from the previous section, we can write GFI as

$$\text{GFI} = 1 - \frac{b_M t_M + a_M}{b_0 t_0 + a_0}.$$

When the model is correctly specified, $n\mathbf{e}'\mathbf{W}\mathbf{e}$ asymptotically follows a chi-square distribution, but $n\mathbf{s}'\mathbf{W}\mathbf{s}$ will not approximately follow a noncentral chi-square distribution. Based on the result from the previous section, GFI changes its value as well as its distribution when N , the distribution of \mathbf{x} , as well as the model specification and the magnitude of Σ_0 change. As n tends to infinity, the limit of GFI measures the relative reduction of all the variances-covariances by model Σ_M relative

to a model with all elements equal to zeros. Similar to NFI, the limit of GFI depends on the measure $D(\cdot, \cdot)$ which further depends on the weight \mathbf{W} . The speed of convergence can be very slow. Interpreting GFI as a reduction of all the variances and covariances in the population may not be appropriate.

Using Equation 12, the fit index NNFI can be rewritten as

$$\text{NNFI} = 1 - \frac{b_M t_M / df_M + a_M / df_M - 1}{b_I t_I / df_I + a_I / df_I - 1}.$$

When T_M and T_I are based on T_{ML} and Σ_I is approximately correct, then $b_I \approx b_M = b$ and $a_I \approx a_M \approx 0$ within the class of elliptical distributions. Thus, $\text{NNFI} \approx (t_I / df_I - t_M / df_M)(t_I / df_I - 1/b)$ is a decreasing function of b . However, the model may not be approximately correct in practice, and hence the value of NNFI as well as its distribution will be affected by the distribution of \mathbf{x} , the sample size N , the estimation method, as well as model misspecifications. Of course, using a noncentral chi-square distribution to interpret the value of NNFI is generally inappropriate.

For RNI, the nonnormed version of CFI, we have

$$\text{RNI} = 1 - \frac{b_M t_M + a_M - df_M}{b_I t_I + a_I - df_I}.$$

In practice, CFI and RNI will be affected by sample size, the distribution of \mathbf{x} and the measure $D(\cdot, \cdot)$ used in constructing the fit indices. When T cannot be described by a noncentral chi-square distribution, interpreting CFI or RNI as a reduction of NCP may not be appropriate.

Using Equation 12 we can rewrite RMSEA as

$$\text{RMSEA} = \sqrt{(b_M t_M + a_M - df_M) / (n \times df_M)}.$$

It is obvious that RMSEA is an increasing function of a_M and b_M , which will change with the distribution of \mathbf{x} , the sample size, the model misspecification as well as the discrepancy function $D(\cdot, \cdot)$. The confidence interval for RMSEA, as printed in standard software, makes sense only when $b_M = 1$ and $a_M = 0$, which will not hold in any realistic situation. Notice that the larger the n , the smaller the effect of a_M ; but the effect of b_M will not diminish even when n is huge.

In conclusion, when the t in Equation 12 is anchored at a chi-square distribution or at the distribution of a statistic under fixed conditions, fit indices F defined through T are functions of a and b , which change across conditions. Desir-

able properties of F are difficult to achieve even with the idealized elliptically distributed data.

MATCHING FIT INDICES WITH STATISTICS

For fit indices involving $T - df$, the rationale is $E(T|H_0) = df$ so that $\hat{\delta} = T - df$ is an unbiased estimate of the part of $E(T)$ due to model misspecifications while df is the part due to random sampling that is not related to model misspecification. When $E(T|H_0) \neq df$, the meaning of $T - df$ is not clear. One cannot use the “degrees of freedom” to justify the use of $\hat{\delta} = T - df$. For example, when $E(T|H_0) = 1.5df$, then $\hat{\delta} = T - df$ does not make more sense than $\tilde{\delta} = T - c$ with c being an arbitrary constant. Fortunately, there do exist a few statistics that approximately satisfy $E(T|H_0) = df$. Based on the simulation results of Hu et al. (1992), Chou et al. (1991), Curran et al. (1996), Yuan and Bentler (1998b), Fouladi (2000) and others, the statistic T_R approximately have a mean equal to df when n is relatively large. This is also implied by the asymptotic theory, as reviewed in the second section of this article. For small n , results in Bentler and Yuan (1999) indicate that $E(T_R)$ is substantially greater than df even when \mathbf{x} follows a multivariate normal distribution. The results in Bentler and Yuan (1999) and Yuan and Bentler (1998b) indicate that $(T_{CRADF}|H_0)$ approximately has a mean equal to df regardless of the distribution of \mathbf{x} and N . For convenience, some of the empirical results in Bentler and Yuan (1999) and Yuan and Bentler (1998b) are reproduced in Table 5, where similar data generation schemes as in Table 1 were used.

TABLE 5
Empirical Means^a of T_R , T_{AR} , and T_{CRADF} ($df = 87$)

		Sample Size N									
		90	100	110	120	150	200	300	500	1000	5000
Normal \mathbf{x}	T_R	96.33	95.70	94.95	94.16	90.89	89.75	88.24	87.72	87.91	87.29
	T_{AR}	88.57	88.77	88.71	88.49	86.52	86.52	86.12	86.46	87.28	87.16
	T_{CRADF}	87.42	88.90	89.90	89.66	90.36	90.50	90.22	89.81	88.31	87.44
Elliptical \mathbf{x}	T_R	96.52	95.76	94.57	93.80	90.47	88.48	87.19	86.11	86.02	86.67
	T_{AR}	88.75	88.83	88.35	88.15	86.12	85.29	85.10	84.87	85.40	86.55
	T_{CRADF}	87.50	88.55	88.97	88.83	89.28	90.11	89.99	89.91	89.09	87.84
Normal \mathbf{f} & skew \mathbf{e}	T_R	98.80	98.30	96.66	95.75	92.12	90.64	88.93	88.19	88.24	87.19
	T_{AR}	90.84	91.18	90.30	89.98	87.69	87.38	86.80	86.92	87.61	87.07
	T_{CRADF}	87.39	88.21	88.76	89.07	88.57	89.04	88.39	88.06	88.81	88.08
Skew \mathbf{f} & \mathbf{e}	T_R	99.68	97.36	96.34	95.40	91.77	90.77	88.92	87.52	86.68	87.61
	T_{AR}	91.65	90.31	90.01	89.65	87.36	87.50	86.79	86.26	86.06	87.48
	T_{CRADF}	87.45	88.44	88.85	89.43	88.52	88.87	87.97	88.17	88.36	88.50

^aAll the three statistics asymptotically follow χ^2_{87} .

Notice that the empirical mean of T_{ML} is greater than df even when data are normally distributed, especially when N is small (see Bentler & Yuan, 1999). The well-known Bartlett correction is to make $E(T_{ML})$ more close to df . We might apply such a correction to the statistic T_R when data are not normally distributed. The Bartlett correction for exploratory factor model (EFM) with normal data already exists (Lawley & Maxwell, 1971, pp. 35–36). Yuan, Marshall and Bentler (2002) applied it to the rescaled statistic in exploratory factor analysis. The Bartlett correction for general covariance structures also exists for T_{ML} when data are normally distributed (see Wakaki, Eguchi, & Fujikoshi, 1990). However, the correction is quite complicated even for rather simple models. For nonnormal data, this correction is no longer the Bartlett correction. Here we propose an ad hoc correction to the statistic T_R . Consider the Bartlett correction with m factor in EFM,

$$T_{BML} = [N - 1 - (2p + 5)/6 - 2m/3] D_{ML} [\mathbf{S}, \boldsymbol{\Sigma}(\hat{\boldsymbol{\theta}})], \quad (14)$$

where the total number of free parameters is $p(m + 1) - m(m - 1)/2$. Notice that the CFM and EFM are identical when there is a single factor, then the coefficient in Equation 14 is $N - 5/3 - (2p + 5)/6$. In formulating a SEM model, unidimensional (univocal) measurements are recommended and typically used in practice (Anderson & Gerbing, 1988). When all the measurements are unidimensional, the number of factor loadings in the SEM model is equal to that of the 1-factor model. The factors can be correlated or predicted by each other. With m factors, there are $m(m - 1)/2$ additional parameters when the structural model is saturated. Considering that there may exist double or multiple loadings and the structural model may not be saturated, we propose the following ad hoc correction to T_R ,

$$T_{AR} = n^{-1} [N - 5/3 - (2p + 5)/6 - (m - 1)/3] T_R. \quad (15)$$

Table 5 contrasts the empirical means of T_{AR} and T_R , based on the empirical results in Yuan and Bentler (1998b) and Bentler and Yuan (1999). Although many of the empirical means of T_{AR} are still greater than $df = 87$ when n is small, they are much nearer to the df than the corresponding ones of T_R . A parallel T_{AML} follows when applying the correction factor in Equation 15 to T_{ML} .

The statistic T_{CADF} also has empirical means very close to df for various distribution and sample size conditions (Yuan & Bentler, 1997b). However, it is easier to get a converged solution with minimizing D_{ML} than with D_{ADF} . The statistic T_{AML} can also be used when data are normally distributed. But $E(T_{AML})$ is unpredictable when the distribution of \mathbf{x} is unknown as with practical data. A nonsignificant Mardia's multivariate kurtosis does not guarantee that $E(T_{ML}) \approx df$, as showed in Example 3 of Yuan and Hayashi (2003). So it is wise to avoid T_{ML} or even T_{AML} be-

fore an effective way of checking normality is performed. The means of T_{ADF} and T_{RADF} are much greater than df unless N is huge (see Hu et al., 1992; Yuan & Bentler, 1997b, 1998b), thus, they are not recommended to be used with fit indices involving df . Although both T_{CRADF} and T_R are now available in EQS (Bentler, 2004), T_{CRADF} can only be applied when $n > df$. Similarly, T_R makes sense only when $n > df$ (see Bentler & Yuan, 1999) although it can still be numerically calculated when $n < df$.

Yuan and Hayashi (2003) showed that the statistic T_{ML} applied to a proper robustified sample can closely follow χ^2_{df} . When T approximately follows χ^2_{df} , the empirical mean of T also approximately equals df . Also, applying T_{ML} with a robust procedure does not need the specific condition $n > df$. Of course, one may not be able to apply the robust procedures to any data. We will further discuss this in the appendix.

In this section we argue that fit indices involving $T - df$ do not make much sense when $E(T) \neq df$. Because T_{AR} and T_{CRADF} approximately satisfy $E(T) = df$, they should be used in defining fit indices involving df instead of T_{ML} or T_{ADF} . Robust procedures with T_{ML} are preferred when data contain heavy tails or when $n < df$, but one needs to check that $E(T_{ML}) = df$ using resampling or simulation (see Yuan & Hayashi, 2003).

STABILITY OF FIT INDICES WHEN CONDITIONS CHANGE

Notice that the distributions of T_{AR} and T_{CRADF} still change substantially even though their empirical means approximately equal df . So we still need to study how the distributions of fit indices change when changing conditions. In this section, we mainly study the effect of the distribution of \mathbf{x} with different sample sizes. This is partially because people seldom check the distribution of the data when using fit indices in practical data analysis. Previous studies on fit indices mainly focused on T s based on $\mathbf{x} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ with different sample sizes or estimation methods (Anderson & Gerbing, 1984; Fan, Thompson, & Wang, 1999; La Du & Tanaka, 1989, 1995; Marsh et al., 1988, 2004), but not the statistic T_{AR} or T_{CRADF} . We will resort to Equation 12 and rewrite it as $F = bf + a$ for the current purpose, where f represents a fit index when $\mathbf{x} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ and F represents a fit index when \mathbf{x} follows a different distribution. Because it does not make sense to apply T_{ML} or T_{RADF} to fit indices involving df when data are nonnormally distributed or when sample size is not huge, we only study the performance of these fit indices when evaluated by T_{AR} and T_{CRADF} . Specifically, we will study the effect of distributional change of \mathbf{x} on NFI, NNFI, GFI, CFI, and RMSEA. Because NFI does not involve df , we will report its distribution when evaluated using T_{ML} , T_{AR} , T_{RADF} , and

T_{CRADF} . Similarly, we will evaluate GFI when T_M and T_0 are T_{IRLS} or T_{RADF} , as previously discussed. Parallel to Tables 2 to 4 and with the same CFM as in Equation 9, we choose three distribution conditions for \mathbf{x} . Table 6 contains the intercept a and slope b for the above designs when $N = 150$ and 500 . It is obvious that most of the fit indices change distributions substantially when \mathbf{x} changes its distribution. Sample size also has a big effect on a and b . Among these, RMSEA changes its overall distribution the least. Notice that both T_{AR} and T_{CRADF} change distributions substantially in Tables 2 to 4. RMSEA is a function of $\hat{\delta}$ and totally decided by T . The greater stability of RMSEA is due to the effect of a square root, while no other fit index studied here uses such a transformation. The same square root effect may attenuate its sensitivity when the model Σ_M changes. Hu and Bentler (1998) studied sensitivity of fit indices, but not when they are evaluated by T_{AR} and T_{CRADF} . Further study in this direction is needed.

TABLE 6
Intercept a and Slope b With $F = bf + a$, Where f Is the Distribution of F for Normal Data

F	T	N	Elliptical \mathbf{x}		Skew \mathbf{f} & Normal \mathbf{e}		Skew \mathbf{f} & \mathbf{e}	
			b	a	b	a	b	a
NFI	T_{ML}	150	2.510	-1.443	3.902	-2.734	3.306	-2.152
		500	4.220	-3.160	5.766	-4.667	4.756	-3.673
	T_{AR}	150	3.676	-2.504	9.244	-7.715	9.795	-8.207
		500	5.780	-4.668	20.864	-19.394	21.538	-20.048
	T_{RADF}	150	1.257	-0.265	1.466	-0.576	1.453	-0.502
		500	1.692	-0.688	3.383	-2.442	3.312	-2.355
	T_{CRADF}	150	0.949	-0.009	0.901	-0.057	0.891	-0.027
		500	1.200	-0.215	1.816	-0.892	1.738	-0.815
GFI	T_{IRLS}	150	3.244	-2.137	3.246	-2.137	3.166	-2.046
		500	5.354	-4.274	5.482	-4.398	4.409	-3.344
	T_{RADF}	150	2.337	-1.351	2.288	-1.307	3.342	-2.367
		500	3.963	-2.968	4.012	-3.019	7.657	-6.680
CFI	T_{AR}	150	2.256	-1.251	9.524	-8.489	12.132	-11.103
		500	2.571	-1.567	11.760	-10.737	14.082	-13.060
	T_{CRADF}	150	1.122	-0.111	1.616	-0.663	1.423	-0.407
		500	1.346	-0.346	3.618	-2.626	3.001	-2.002
NNFI	T_{AR}	150	2.322	-1.313	280.605	-278.981	15.448	-14.396
		500	2.948	-1.939	21.691	-20.635	18.770	-17.729
	T_{CRADF}	150	1.044	-0.018	61.079	-47.357	32.034	-23.782
		500	1.301	-0.305	5.680	-4.590	3.439	-2.413
RMSEA	T_{AR}	150	0.932	-0.002	0.916	-0.002	1.017	0.006
		500	0.914	-0.003	0.871	-0.003	1.015	-0.001
	T_{CRADF}	150	0.930	-0.001	0.937	-0.001	0.864	-0.001
		500	0.964	0.000	0.937	0.000	0.861	-0.000

The results of this section tell us that fit indices change their distributions substantially when conditions change. Commonly used cutoff values or confidence intervals for fit indices do not reflect these changes and thus provide questionable values regarding model fit/misfit.

THE POWER OF A FIT INDEX

There exist two seemingly unrelated aspects of power based on fit indices. One is the ability for a fit index to distinguish a good model from a bad model. The other is to first use fit indices F to specify null and alternative hypotheses and then to analyze the power of T under these two hypothesis. We will discuss these two aspects separately.

Commonly used cutoff values such as 0.05 or 0.95 for fit indices do not directly relate the fit indices to any null hypothesis. So one cannot treat the cutoff value as the traditional critical value and use the corresponding rejection rate as power. Even when one specifies a null hypothesis, say the model is correctly specified or the distance between the model $\Sigma(\theta)$ and the population Σ_0 is less than a small number, then 0.05 or 0.95 may have nothing to do with the upper 5% quantiles of the distribution of the fit index under the specified null hypothesis. As we have seen from the previous sections, all fit indices change distributions when N , \mathbf{x} and $D(\cdot, \cdot)$ change; we cannot relate the means of fit indices to the NCP even under the idealized conditions in Equation 2. In practice, we generally do not know the distribution of fit indices even when Σ_M is correctly specified, and ignoring their relationship to model misspecifications.

With the above mentioned difficulties, a more sensible approach to study the sensitivity or power of fit indices is by simulation or bootstrap, as in Hu and Bentler (1998, 1999), Gerbing and Anderson (1993), Bollen and Stine (1993), Yung and Bentler (1996), Muthén and Muthén (2002), and Yuan and Hayashi (2003). In such a study, the N , $D(\cdot, \cdot)$, and the distribution of \mathbf{x} are all given. We still need to specify interesting null H_0 and alternative H_1 hypotheses. The null and alternative hypotheses should include both Σ_0 and Σ_M . Then the distributions of a fit index F under H_0 and H_1 can be estimated through simulations. The cutoff value can be obtained from a proper quantile of the estimated distribution ($\widehat{F} | H_0$). The power of F at the given conditions will be the proportion of ($\widehat{F} | H_1$) that fall into the rejection region decided by the cutoff value. We need to emphasize that simulation or bootstrap are tailor-made approaches and may not be generalizable when conditions change. As we have seen, in addition to model misspecifications, the distribution of F changes with the sample size N , the distribution of \mathbf{x} as well as the chosen statistic T . The most sensitive F under one set of conditions may no longer be most sensitive under a different set of conditions.

When studying power of T one has to specify H_0 and H_1 (e.g., Satorra & Saris, 1985). Traditionally, H_0 represents a correct model and H_1 represents an interesting alternative model. Because substantive models are at most approximately correct, MacCallum et al. (1996), MacCallum and Hong (1997), and Kim (2003) proposed to let H_0 represent a misspecified model with misspecification measured by RMSEA, CFI, GFI, and other fit indices. They used $(T | H_0) \sim \chi_{df}^2(\delta_0)$ and $(T | H_1) \sim \chi_{df}^2(\delta_1)$ to calculate the critical value and power. Such a proposal would allow researchers to test interesting but not necessarily perfect models when both $(T | H_0) \sim \chi_{df}^2(\delta_0)$ and $(T | H_1) \sim \chi_{df}^2(\delta_1)$ are attainable. When they are not attainable, then the critical value obtained using $\chi_{df}^2(\delta_0)$ or power based on $\chi_{df}^2(\delta_1)$ may have little relationship with the model misspecification. As with distributions of fit indices, distributions of the statistics depend on sample size, the distribution of \mathbf{x} as well as the choice of statistic T , in addition to model misspecification. Again, a more sensible approach to power might be by simulation or bootstrap (see Yuan & Hayashi, 2003).

In conclusion, power or sensitivity of fit indices and test statistics might need to be evaluated by simulation. In such a study, one has to control both type of errors² using simulated critical values rather than referring F to 0.05 or 0.95 or referring T to $\chi_{df}^2(\delta)$. However, the findings by simulation in one set of conditions may not be generalizable to a different set of conditions.

DISCUSSION

The analysis and empirical results in the previous sections should help to provide a clearer overall picture of the relationship between fit indices and test statistics. We will further discuss some issues related to their relationship, which may lead to a better understanding of the properties of the fit indices.

As indicators for model fit, fit indices should relate to model misspecification. The commonly used measure of model misspecification is given by the τ in Equation 7. However, the definition there only makes a clear sense when $D = D_{ML}$ corresponding to T_{ML} at the population level. At the sample level, $D_{ML}[\mathbf{S}, \Sigma(\hat{\theta})]$ can still be calculated but it depends on N and the distribution of \mathbf{x} . Unfortunately, current literature does not provide us an analytical formula relating $D_{ML}[\mathbf{S}, \Sigma(\hat{\theta})]$ to τ and other conditions such as N and the underlying distribution of \mathbf{x} . For example, at a given N we do not know how good the approximation in Equation 1 is even when \mathbf{x} follows an elliptical distribution and the model is correctly specified. What we know is that $D_{ML}[\mathbf{S}, \Sigma(\hat{\theta})]$ is consistent for τ , as well as that $E\{D_{ML}[\mathbf{S}, \Sigma(\hat{\theta})]\}$ is generally increasing with τ and the population kurtosis κ when \mathbf{x} follows an elliptical

²Yuan and Bentler (1997b), Fouladi (2000), and Curren et al. (1996) studied the power of various statistics but none of these studies control type I errors properly.

cal distribution. But such knowledge is not enough to accurately estimate τ , establish reliable cutoff values, or compute meaningful confidence intervals for τ .

When using $T_R, T_{AR}, T_{ADF}, T_{CADF}, T_{RADF}, T_{CRADF}$ for model evaluation, the measure corresponding to τ should be

$$\tau_{\Gamma} = \min_{\theta} D^{(\Gamma)} [\Sigma_0, \Sigma(\theta)],$$

where Γ is the fourth-order population covariance matrix approximately equal to $NCov(s)$, $D^{(\Gamma)}$ is the population counterpart of the discrepancy function corresponding to each T . In Monte Carlo studies, Γ can be obtained if one generates random numbers following the procedure of Yuan and Bentler (1997a). One generally does not know Γ with real data. The problem is that all fit indices and test statistics are related to Γ at the population level, including $E\{D_{ML}[\mathbf{S}, \Sigma(\hat{\theta})]\}$. But Γ itself is difficult to estimate due to its large dimension. The idea in robust procedures is to transform the sample so that the corresponding Γ is comparable to that in a normal population with the same Σ_0 . The sensitivity or power of a fit index is closely related to Γ . It is well-known that, with given Σ_0 and $\Sigma(\theta)$, the greater the Γ (in the sense of positive definiteness) the smaller the power of a test statistic (e.g., Shapiro & Browne, 1987; Tyler, 1983; Yuan & Hayashi, 2003; Yuan et al., 2004). Fit indices that only involve T_M will inherit such a property. It is unclear whether the sensitivity of a fit index involving T_I will also be inversely affected by a large Γ matrix.

In practice, we deal with τ or τ_{Γ} through T . When $T \sim \chi_{df}^2(\delta)$ holds, $\hat{\delta} = T - df$ contains valuable information for model misspecification. The confidence intervals for δ and monotonic functions of δ also make probabilistic sense, although we may not be able to relate δ to τ or to the expectation of a fit index $E(F)$ explicitly. We want to reemphasize that the result in Equations 6 and 7 is not correct when the population covariance matrix Σ_0 is fixed.

When $E(T|H_0) = df$ is true but $T \sim \chi_{df}^2(\delta)$ does not hold, $\hat{\delta} = T - df$ also contains valuable information for model misspecification under the condition that $E(T|H_1)$ increases as the model deteriorates. We can interpret $\hat{\delta}$ as the systematic part of T that is beyond the effect of sampling errors corresponding to the correct model. But a confidence interval for δ based on $T \sim \chi_{df}^2(\delta)$ does not make sense.

When $E(T|H_0) \neq df$, we might modify $\hat{\delta} = T - df$ to

$$\hat{\delta}_D = T - E(\widehat{T|H_0}),$$

where the subscript D is used to denote the involved discrepancy measure in formulating T . The population counterpart of $\hat{\delta}_D$ is $\delta_D = E(T|H_1) - E(T|H_0)$, which is the systematic part of T that is beyond the effect of sampling errors cor-

responding to the correct model. It is obvious that $\delta_D = 0$ when $\Sigma(\theta)$ is correctly specified, regardless of N , D or \mathbf{x} . The drawback of $\hat{\delta}_D$ is that $E(\widehat{T} | H_0)$ is not as easy to obtain as df . Instead, one has to use a resampling based procedure or other alternatives to obtain a $E(\widehat{T} | H_0)$. See Yuan and Marshall (2004) for further comparison of δ_D with δ and ways to obtain confidence intervals for δ_D .

With the distribution of \mathbf{x} generally unknown and N uncontrollable, $E(T|H_0) \approx df$ is not difficult to achieve. When N is not too small, $(T|H_0)$ approximately following χ_{df}^2 is also achievable by choosing a proper T . But $(T|H_1)$ approximately following $\chi_{df}^2(\delta)$ is generally not achievable. Actually, we cannot analytically relate $E(F)$ to δ even when $(T | H_1) \sim \chi_{df}^2(\delta)$. For fit indices that are monotonic functions of T without referring to a base model, a confidence interval for δ can be obtained by assuming $T \sim \chi_{df}^2(\delta)$, but not a confidence interval for τ or τ_Γ because the key equation

$$\tau = \delta/n,$$

still depends on unrealistic conditions (Yuan & Bentler, in press; Yuan & Marshall, 2004). For fit indices $g(\hat{\delta}_M, \hat{\delta}_I)$ that involve both T_M and T_I , assumptions in Equation 2 do not allow one to obtain even a confidence interval for $g(\delta_M, \delta_I)$. So, the noncentral chi-square assumption is neither realistic nor necessary in making inference for τ or τ_Γ . For NFI, the sample size N is cancelled by the ratio. We will have a consistent estimator for $\text{NFI}_0 = 1 - \min_{\theta} D[\Sigma_0, \Sigma(\theta)] / \min_{\theta} D[\Sigma_0, \Sigma_I(\theta)]$. Similar conclusion can be reached for other fit indices when N is cancelled. Of course, consistency is a very minimal requirement, which does not tell us how far it is from the population quantity for a finite N .

We might need to distinguish the role of a T as a statistic to evaluate whether a model is correctly specified from its role in formulating a fit index. As a test statistic, T is well behaved when $(T|H_0)$ approximately overlaps with χ_{df}^2 at the 95% quantile and the greater the $(T|H_1)$ the better the T . We do not really need for T to follow chi-square distributions.

CONCLUSION AND RECOMMENDATION

We wanted to identify some statistics or estimation methods coupled with fit indices that are relatively stable when sample sizes and sampling distributions change. We may also want a fit index F to perform stably when changing the base-statistic T . Since the choice of T is controllable, it is not too bad when F varies across different T s. The problem is that most fit indices are not stable across the uncontrollable conditions N and the distribution of \mathbf{x} . RMSEA is relatively most stable among the commonly used fit indices. But it is totally decided by T and none of the T s are stable when changing N or \mathbf{x} . Thus, the stability of RMSEA may be just due to the ar-

tificial effect of the square root, which will attenuate its sensitivity across model misspecifications.

Although our analysis of fit indices still leaves many uncertainties, some conclusions can nonetheless be reached.

1. Given the population covariance matrix and the model structure, the mean value as well as the distribution of fit indices change with the sample size, the distribution of the data as well as the chosen statistic. Fit indices also reflect these variables in addition to reflecting model fit. Thus, cutoff values for fit indices, confidence intervals for model fit/misfit, and power analysis based on fit indices are open to question.

2. Statistics T_{AR} and T_{CRADF} have means approximately equal to df and are recommended for calculating fit indices involving df . Other statistics should not be used unless certain conditions such as normally distributed data and a large enough sample size are realized.

3. The asymptotic distribution of T_{AR} is generally unknown. The distribution of T_{CRADF} may be far from χ_{df}^2 when the sample size is not large enough. Assuming they follow noncentral chi-square distributions is even more farfetched. Confidence intervals for model fit/misfit are not justified even when T_{AR} or T_{CRADF} are used in the evaluation.

4. The statistic T_{ML} applied to a robustified sample is recommended when the majority of the data are symmetrically distributed. This procedure is better coupled with the bootstrap procedure so that one can check that $(T_{ML}|H_0)$ approximately follows χ_{df}^2 . The noncentral chi-square distribution is neither realistic nor necessary for evaluating model fit in the bootstrap procedure.

5. Our conclusions are temporary, as further study is needed to understand how each fit index is related to model misfit when evaluated by the more reasonable statistics T_{AR} , T_{CRADF} , or T_{ML} based on a robust procedure.

We believe our analysis and discussion have effectively addressed the questions raised at the beginning of the article. Although most of our results are on the negative side of current practice, fit indices are still meaningful as relative measures of model fit/misfit. Specifically, for NFI, NNFI, CFI, and GFI there exists $E(F|C_1) > E(F|C_2)$ when C_1 contains the model corresponding to a smaller τ or τ_{Γ} than C_2 while all the other conditions such as N , the distribution of \mathbf{x} , and $D(\cdot, \cdot)$ in C_1 and C_2 are the same; the opposite inequality sign holds for RMSEA. But such a relative value of fit indices should be distinguished from relative fit indices that utilize a base model. The relative value should not be interpreted as cutoff values being relative, for example, some use 0.95 while others use 0.90 to decide the adequacy of a model. We want to emphasize that the purpose of the article is not to abandon cutoff values but to point out the misunderstanding of fit indices and their

cutoff values. We hope the result of the article can lead to better efforts to establish a scientific norm on the application of fit indices.

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APPENDIX

EQS 6.0 (Bentler, 2004) has the case-robust procedure based on Yuan and Bentler (1998c), which contains two tuning parameters in controlling the case weights. Here we give a brief introduction to the robust transformation proposed by Yuan et al. (2000), where Huber-type weights (see Tyler, 1983) that only contain one tuning parameter are used.

Let ρ be the percentage of influential cases one wants to control, and r be a constant decided by ρ through $P(\chi_p^2 > r^2) = \rho$. Denote the Mahalanobis distance as

$$d = \left[(\mathbf{x} - \boldsymbol{\mu})' \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right]^{1/2}.$$

The Huber-type weights are given by

$$u_1(d) = \begin{cases} 1, & \text{if } d \leq r \\ r/d, & \text{if } d > r \end{cases}$$

and $u_2(d^2) = [u_1(d)]^2/\varphi$, where φ is a constant decided by ρ through $E[\chi_{\rho}^2 u_2(\chi_{\rho}^2)] = p$. The purpose of φ is to make the resulting covariance matrix estimator unbiased when $\mathbf{x} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma})$. Robust mean vector and covariance matrix can be obtained by iteratively solving

$$\boldsymbol{\mu} = \sum_{i=1}^N u_1(d_i) \mathbf{x}_i / \sum_{i=1}^N u_1(d_i), \tag{16}$$

$$\boldsymbol{\Sigma} = \sum_{i=1}^N u_2(d_i^2) (\mathbf{x}_i - \boldsymbol{\mu})(\mathbf{x}_i - \boldsymbol{\mu})' / N, \tag{17}$$

where d_i is the M -distance based on the i^{th} case. Notice that the only tuning parameter in solving Equations 16 and 17 is ρ . It is obvious that the greater the d_i the smaller the weights $u_{1i} = u_1(d_i)$ and $u_{2i} = u_2(d_i^2)$. Denote the solution to Equations 16 and 17 as $\hat{\boldsymbol{\mu}}$ and $\hat{\boldsymbol{\Sigma}}$. Yuan et al. (2000) proposed the transformation

$$\mathbf{x}_i^{(\rho)} = \sqrt{u_{2i}} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}). \tag{18}$$

Yuan and Hayashi (2003) suggested applying the bootstrap procedure to $\mathbf{x}_i^{(\rho)}$ and verifying the distribution of $(\mathcal{I}H_0)$ across the bootstrap samples against the distribution of χ_{df}^2 . When data contain heavy tails, by adjusting ρ , they showed that the empirical distribution of $(T_{ML}|H_0)$ can match χ_{df}^2 very well for different real data sets.

It is obvious that $\mathbf{S}_r = \hat{\boldsymbol{\Sigma}}$ is the sample covariance matrix of $\mathbf{x}_i^{(\rho)}$. In this setup, the population covariance matrix $\boldsymbol{\Sigma}_r$ corresponding to \mathbf{S}_r may not equal $\boldsymbol{\Sigma}_0$. When the data are elliptically distributed, analyzing \mathbf{S}_r and \mathbf{S} leads to the same substantive conclusion. In practice, data might contain outliers which will make a true symmetric distribution skewed at the sample level. In such a situation, analyzing \mathbf{S}_r is preferred. If one believes that the true distribution of \mathbf{x} is skewed, then the results corresponding to $\boldsymbol{\Sigma}_r$ may not be substantively equivalent to those corresponding to $\boldsymbol{\Sigma}_0$. Hampel, Ronchetti, Rousseeuw, and Stehel's (1986, p. 401) discussion implies that analyzing \mathbf{S}_r might still be preferred even when \mathbf{x} has a skew distribution.

The SAS IML program at www.nd.edu/~kyuan/courses/sem/RTRANS.SAS performs the iterative procedure in Equations 16 and 17. The transformed sample $\mathbf{x}_i^{(\rho)}$ in Equation 18 is printed out at the end of the program. When applying this program, one needs to modify the program lines 2 and 3 so that a proper ASCII file is correctly read. The only other thing one needs to change is the tuning parameter ρ (rho) in the main subroutine. The program also includes Mardia's multivariate skewness and kurtosis for \mathbf{x}_i and $\mathbf{x}_i^{(\rho)}$. Yuan et al. (2000) suggested using the standardized Mardia's kurtosis < 1.96 to select ρ . Examples in Yuan and Hayashi (2003) implies that even when Mardia's kurtosis is not significantly different from that of a multivariate normal distribution, $(T_{ML}|H_0)$ may not approximately follow χ_{df}^2 .