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# Preliminary Testing for Normality: Is This a Good Practice?

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## *Invited Article:* **Preliminary Testing for Normality: Is This a Good Practice?**

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Normality is a distributional requirement of classical test statistics. In order for the test statistic to provide valid results leading to sound and reliable conclusions this requirement must be satisfied. In the not too distant past, it was claimed that violations of normality would not likely jeopardize scientific findings (See [Hsu & Feldt, 1969;](#page-17-0) [Lunney,](#page-17-1)  [1970\)](#page-17-1). Recent revelations suggest otherwise (See e.g., [Micceri, 1989;](#page-17-2) [Keselman, Huberty,](#page-17-3)  [Lix et al., 1998;](#page-17-3) [Erceg-Hurn, Wilcox, & Keselman, 2013;](#page-16-0) [Wilcox and Keselman, 2003;](#page-18-0) [Wilcox, 2012a,](#page-18-1) [b\)](#page-18-2). Unfortunately the data obtained in psychological investigations rarely, if ever, meet the requirement of normally distributed data [\(Micceri, 1989;](#page-17-2) [Wilcox, 2012a,](#page-18-1)  [b\)](#page-18-2). Consequently, it could be the case that the results from many of the investigations conducted in psychology provide invalid results. Accordingly, authors recommend that researchers attempt to assess the validity of assuming data are normal in form prior to conducting a test of significance [\(Erceg-Hurn, et al., 2013;](#page-16-0) [Keselman, et al., 1998\)](#page-17-3). Present evidence suggests that a popular fit-statistic, the Kolmogorov-Smirnov test does a poor job of evaluating whether data are normal. Our investigation based on this statistic and other fit-statistics provides a more favorable picture of preliminary testing for normality.

*Keywords:* Assessing normality, fit statistics, g-and-h non-normal skewed and kurtotic data, contaminated mixed-normal distributions; outlying value(s), Likert scales

## **Introduction**

Psychological researchers are often reminded that the validity of their statistical tests and the conclusions derived from these tests depends to a great extent on whether the derivational assumptions of the test procedures have been satisfied (e.g., See [Keselman, Huberty, Lix et al., 1998;](#page-17-3) [Wilcox, 2012a,](#page-18-1) [b;](#page-18-2) [Wilcox &](#page-18-0) 

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[Keselman, 2003\)](#page-18-0). Consequently, though not a common practice, researchers are still reminded about assessing derivational assumptions (See [Erceg-Hurn, Wilcox,](#page-16-0)  [& Keselman, 2013;](#page-16-0) [Kirk, 2013;](#page-17-4) [Schoder, Himmelmann & Wilhelm, 2006;](#page-18-3) [Wilcox](#page-18-0)  [& Keselman, 2003\)](#page-18-0). Almost all inferential methods require that in the population(s) the data is (are) normally distributed (as well as other requirements not relevant to this paper). Violation of the normality assumption can have a deleterious effect on the Type I error rate of test statistics (See [Wilcox, 2012a,](#page-18-1) [b;](#page-18-2) [Wilcox & Keselman, 2003\)](#page-18-0). Although the Type I error rate is widely viewed as being relatively unaffected by non-normality, [Bradley \(1980\)](#page-16-1) has pointed out conditions in which this is not true. This finding is also evident in the findings of recent studies and published texts (e. g., See [Hempel, Ronchetti, & Rousseeuw,](#page-16-2)  [1986;](#page-16-2) [Huber & Ronchetti, 2009;](#page-17-5) [Maronna, Martin, & Yohai, 2006;](#page-17-6) [Micceri, 1989;](#page-17-2) [Schoder, et al., 2006;](#page-18-3) [Staudte & Sheather, 1990;](#page-18-4) [Wilcox, 2012a,](#page-18-1) [b;](#page-18-2) [Wilcox &](#page-18-0)  [Keselman, 2003\)](#page-18-0).

Applied researchers can examine plots of their data and/or perform tests to assess the assumption, i. e., normality. Evaluating graphs (e.g., box-plots, stemand-leaf, box and whisker, QQ plots) of ones data to assess whether data are normally distributed can be problematic since the determination relies on a subjective assessment [\(Wilk & Gnanadesikan, 1968\)](#page-18-5). Thus, this practice is oftentimes not typically used when assessing the shape of the distribution of data (See [Schoder, et al., 2006\)](#page-18-3). Researchers tend to prefer exact methods based on formal tests for normality such as the Kolmogorov-Smirnov (K-S) goodness-of-fit statistic (See [Muller & Fetterman, 2002,](#page-17-7) Chapter 7). Furthermore, researchers commonly use the result from a goodness-of-fit test to determine whether the normality of classical test procedures is satisfied thus providing legitimacy to the use of a classical test statistic. Consequently, preliminary testing for normality or any distributional shape is quite important in the whole inferential process and has been discussed in various contexts (See e.g., [Cardoso de Oliveira & Ferreira,](#page-16-3)  [2010;](#page-16-3) [Doornik & Hansen, 2008;](#page-16-4) [Sürücü, 2006\)](#page-18-6). However, if the assumption of normality does not appear to be satisfied, researchers use this information to select alternative procedures such as nonparametric methods. Thus, it is important to know how well a preliminary test for normality, e. g., the K-S test, works in detecting non-normal data.

Unfortunately, according to [Schoder, Himmelman, and Wilhelm \(2006\)](#page-18-3) "The Kolmogorov-Smirnov test performs badly on data with single outliers, 10% outliers, and skewed data at sample sizes  $\langle 100 \rangle$  (p. 757) These authors investigated the performance of the K-S test for four types of non-normal data (e.g., normal distribution with a single outlier, normal distribution with 10%

outliers, skewed lognormal distribution with varying skewness, and an ordinal 5 point Likert scale with varying multinomial probabilities) and varying sample size in a pretest-posttest design. The assessment for normality was conducted at a 5% significance level. Unfortunately, the results tabled by [Schoder et al.](#page-18-3) do not support the use of the K-S test as a preliminary test to assess normality of the data.

Because it is strongly believed that validity assumptions such as normality should be verified before adopting a classical test of significance that assumes the data in the population is normal in shape, it important to replicate the findings reported by [Schoder, Himmelmann, and Wilhelm \(2006\)](#page-18-3) and extend their study in important ways. (For a contrary view previously noted in this journal, see [Sawilowsky, 2002,](#page-18-3) p. 466-467). Other goodness-of-fit statistics are available (see, e.g., [Muller & Fetterman, 2002,](#page-17-7) Chapter 7). Accordingly, a simulation study was conducted investigating three goodness-of-fit statistics, varying the degree of nonnormality with other distributional shapes not investigated by [Schoder,](#page-18-3)  [Himmelmann, and Wilhelm \(2006\),](#page-18-3) using sample sizes more likely to be encountered in psychological and educational research.

## **Method**

Specifically, in this study the following are manipulated: (1) the procedure used to assess shape of distribution [K-S, Cramer-von Mises (CvM), Anderson-Darling (A-D)] fit-statistics (available through the SAS system), (2) the shapes of distributions (26 cases—14 g-and-h distributions, 8 contaminated normal mixture models, and 4 multinomial models), (3) the sample sizes (20, 40, and 80), depending on distribution, and (4) the level of significance for the fit-statistics (i.e.,  $\alpha = .05, .10, .15$  and .20).

Most statistical packages (e.g., the SAS system) provide numerous fit statistics. Accordingly, it is possible that other tests other than the K-S can adequately assess whether normality, or other distributions, exists in the data. The SAS system was used to implement the K-S, CvM, and A-D fit-statistics. The choices of non-normal distributions are modifications from [Schoder,](#page-18-3)  [Himmelmann and Wilhelm \(2006\)](#page-18-3) and [Zimmerman \(1998\).](#page-18-7) [Schoder, et al. \(2006\)](#page-18-3) investigated a normal distribution with a single outlier, a normal distribution with 10% outliers, skewed lognormal distributions with varying skewness, and an ordinal 5-point Likert scale with varying multinomial probabilities (common they state in dermatological investigations). Many non-normal distributions were investigated via g-and-h distributions (See [Headrick, Kowalchuk, & Sheng, 2008;](#page-16-5) [Hoaglin, 1983;](#page-16-6) [1985;](#page-17-8) [Kowalchuk & Headrick, 2010;](#page-17-9) [Tukey, 1960\)](#page-18-8). These

distributions with their values for skewness and kurtosis are enumerated in [Table](#page-4-0)  [1.](#page-4-0) A substantial number of values of g and h were chosen to cover as broad a spectrum of non-normal distributions that could occur in psychological and behavioral science experiments (e.g., See [Keselman, Huberty, Lix et al., 1998;](#page-17-3) [Micceri, 1989;](#page-17-2) [Wilcox, 2012a,](#page-18-1) [b\)](#page-18-2).

<b>Distribution</b>	<b>Skewness</b>	Kurtosis	<b>Distribution</b>	<b>Skewness</b>	<b>Kurtosis</b>
$q=0, h=.05$	0.00	0.82	$g = .4, h = 0$	1.32	3.26
$q=0, h=.075$	0.00	1.49	$q = .6, h = 0$	2.26	10.27
$q=0, h=.1$	0.00	2.51	$g=1, h=0$	6.19	110.94
$q=0, h=.125$	0.00	4.16	$q = 0.2$ , h= 0.1	1.08	5.50
$q=0, h=.15$	0.00	7.17	$g = .4, h = .1$	2.45	20.30
$q=0, h=.2$	0.00	33.22	$q = .6, h = .1$	4.69	89.80
$q = 2, h = 0$	0.61	0.68	$q = .8, h = .1$	9.27	603.61

<span id="page-4-0"></span>**Table 1.** g-and h-distributions examined in the simulation study with their corresponding measures of skewness and kurtosis

<span id="page-4-1"></span>



The SAS system was used on a Sun Fire X4600 M2 x64 server: 8 x AMD Opteron Model 8220 processor (2.8GHz-dual-core) to generate g- and-h data, by modifying standard normal variates  $Z \sim N(0,1)$  to non-normal variates by specifying values of g and h in the following quantile functions:

$$
q(Z) = q_{g,h}(Z) = \frac{\exp(gZ) - 1}{g} \exp\left(\frac{hZ^2}{2}\right),
$$
 (1)

$$
q(Z) = q_{g,0}(Z) = \frac{\exp(gZ) - 1}{g},
$$
\n(2)

$$
q(Z) = q_{0,h(Z)} = Z \exp\left(\frac{hZ^2}{2}\right) \tag{3}
$$

<span id="page-5-1"></span><span id="page-5-0"></span>Equations [\(2\)](#page-5-0) and [\(3\)](#page-5-1) generate lognormal and symmetric h distributions, respectively. As [Kowalchuk and Headrick \(2010\)](#page-17-9) noted "The parameter  $\pm g$ controls the skew of a distribution in terms of both direction and magnitude. The parameter h controls the tail weight or elongation of a distribution and is positively related with kurtosis." (p. 63). As well, Type I error rates were investigated when data were obtained from a normal distribution  $[g = h = 0]$ , the standard normal distribution (skewness and kurtosis  $= 0$ ).

A number of different contaminated mixed-normal distributions were examined, such as those reported in [Zimmerman \(1998\).](#page-18-7) Contaminated mixednormal distributions have one or more outlying values that deviate from the central mean of the distribution by some amount measured in standard deviation units. For example, Zimmerman examined a mixed normal distribution consisting of samples from *N*(0,1) with probability .95 and from *N*(0, 400) with probability .05. [Tukey \(1960\)](#page-18-8) suggested that outliers are a common occurrence in distributions and others have indicated that skewed distributions frequently depict psychological data (e.g., reaction time data). Accordingly, eight contaminated mixed normal distributions were examined that had one, two, or four outlying values which were five or ten standard deviations from the mean value. These distributions are enumerated in [Table 2.](#page-4-1)

Finally, like [Schoder, Himmelmann and Wilhelm \(2006\),](#page-18-3) a 5-point Likert scale was simulated; such data is frequently gathered in psychological (e.g., from clinical, personality, and social psychological investigations) and other behavioral science investigations. The same conditions investigated by [Schoder et al. \(2006\)](#page-18-3) were investigated. Specifically,

- 1) even distribution (*p*=.02 for each category 0-4);
- 2) symmetric distribution  $(p_0 = 0.1, p_1 = 0.2, p_2 = 0.4, p_3 = 0.2, p_4 = 0.1)$ ; 3) moderately skewed distribution
	- $(p_0 = 0.5, p_1 = 0.3, p_2 = 0.15, p_3 = 0.04, p_4 = 0.01)$ ; and
- 4) heavily skewed distribution (  $p_0 = 0.7$ ,  $p_1 = 0.2$ ,  $p_2 = 0.06$ ,  $p_3 = 0.03$ ,  $p_4 = 0.01$ ).

Thus, for the 5-point Likert scale data there were 4 multinomial distributions that were simulated (See [Table 3\)](#page-6-0).

<span id="page-6-0"></span>**Table 3.** Multinomial distributions based upon Schoder, Himmelmann, and Wilhelm's (2006) probabilities simulated as a five-point Likert Scale

	Even	<b>Symmetric</b>	<b>Moderately</b> <b>Skewed</b>	<b>Heavily Skewed</b>	
n	$(p_0, p_1, p_2, p_3, p_4)$	$(p_0, p_1, p_2, p_3, p_4)$	$(p_0, p_1, p_2, p_3, p_4)$	$(p_0, p_1, p_2, p_3, p_4)$	
20	(.2, .2, .2, .2, .2)	(.1, .2, .4, .2, .1)	(.5, .3, .15, .04, .01)	(.7, .2, .06, .03, .01)	
40	(.2, .2, .2, .2, .2)	(.1, .2, .4, .2, .1)	(.5, .3, .15, .04, .01)	(.7, .2, .06, .03, .01)	
80	(.2, .2, .2, .2, .2)	(.1, .2, .4, .2, .1)	(.5, .3, .15, .04, .01)	(.7, .2, .06, .03, .01)	

The same number of sample size conditions as [Schoder, Himmelmann, and](#page-18-3)  [Wilhelm \(2006\)](#page-18-3) were not investigated, but a reasonable range of values (i.e.,  $n =$ 20,40,80) were includled, depending on the condition investigated. Specifically,

- (i) for the 14 g- and h- distributions, and 2 contaminated normal distributions,  $.95N(0,1) + .05N(0, k)$ ,  $k=25$ , 100, sample sizes of 20, 40 and 80 were chosen.
- (ii) For 2 contaminated normal distributions,  $.9N(0,1) + .1N(0,k)$ ,  $k=25$ , 100, sample sizes of 20 and 40 were chosen.
- (iii) For 2 contaminated normal distributions,  $.975N(01) + .025N(0,k)$ ,  $k=25$ , 100, sample sizes of 40 and 80 were chosen.
- (iv) For 2 contaminated normal distributions,  $.9875N(0,1) + .0125N(0,k)$ ,  $k=25$ , 100, sample size of 80 was chosen.

Lastly, because in preliminary testing it would be quite important to guard against a Type II error (falsely accepting the null hypothesis that the data are normal in form), we selected significance levels of .10 , .15, and .20, in addition to the standard .05. Each condition in the investigation was replicated 5,000 times.

	<b>Distribution</b>	<b>Skewness</b>	<b>Kurtosis</b>	$\alpha = .05$	$\alpha = .10$	$\alpha = .15$
Kolmogorov- <b>Smirnov</b>	Normal*	0.00	0.00	0.0524	0.1082	0.1530
	$g=0, h=.05$	0.00	0.82	0.0726	0.1304	0.1834
	$g=0, h=.075$	0.00	1.49	0.0870	0.1540	0.2094
	$g=0, h=.1$	0.00	2.51	0.1066	0.1838	0.2392
	$g=0, h=.125$	0.00	4.16	0.1320	0.2156	0.2726
	$g=0, h=.15$	0.00	7.17	0.1626	0.2502	0.3100
	$g=0, h=.2$	0.00	33.22	0.2296	0.3194	0.3834
	$g = 0.2$ , $h = 0$	0.61	0.68	0.1030	0.1678	0.2286
	$g = .4, h = 0$	1.32	3.26	0.2436	0.3506	0.4262
	$g = .6, h = 0$	2.26	10.27	0.4450	0.5662	0.6416
	$g=1, h=0$	6.19	110.94	0.7852	0.8648	0.9008
	$g = 2, h = 1$	1.08	5.50	0.1662	0.2530	0.3100
	$g = 4, h = 1$	2.45	20.30	0.3218	0.4204	0.4842
	$g = .6, h = .1$	4.69	89.80	0.5018	0.6026	0.6642
	$g = .8, h = .1$	9.27	603.61	0.6698	0.7602	0.8096
	Normal*	0.00	0.00	0.0494	0.1036	0.1490
	$g=0, h=.05$	0.00	0.82	0.0752	0.1368	0.1952
<b>Cramer-von</b> <b>Mises</b>	$g=0, h=.075$	0.00	1.49	0.0970	0.1658	0.2260
	$g=0, h=.1$	0.00	2.51	0.1286	0.1996	0.2632
	$g=0, h=.125$	0.00	4.16	0.1608	0.2426	0.3038
	$g=0, h=.15$	0.00	7.17	0.1990	0.2842	0.3400
	$g=0, h=.2$	0.00	33.22	0.2756	0.3580	0.4232
	$g = 2, h = 0$	0.61	0.68	0.1100	0.1814	0.2444
	$g = .4, h = 0$	1.32	3.26	0.3082	0.4064	0.4872
	$g = .6, h = 0$	2.26	10.27	0.5570	0.6590	0.7204
	$g=1, h=0$	6.19	110.94	0.8822	0.9268	0.9484
	$g = 2, h = 1$	1.08	5.50	0.1990	0.2826	0.3454
	$g = 4, h = 1$	2.45	20.30	0.3808	0.4730	0.5370
	$g = .6, h = .1$	4.69	89.80	0.5922	0.6728	0.7216
	$g = 0.8, h = 0.1$	9.27	603.61	0.7594	0.8552	0.8626
	Normal*	0.00	0.00	0.0494	0.1036	0.1490
Anderson- <b>Darling</b>	$g=0, h=.05$	0.00	0.82	0.0810	0.1456	0.2040
	$g=0, h=.075$	0.00	1.49	0.1090	0.1816	0.2378
	$g=0, h=.1$	0.00	2.51	0.1444	0.2162	0.2766
	$g=0, h=.125$	0.00	4.16	0.1784	0.2582	0.3200
	$g=0, h=.15$	0.00	7.17	0.2182	0.2992	0.3590
	$g=0, h=.2$	0.00	33.22	0.2924	0.3798	0.4386
	$g = 0.2$ , $h = 0$	0.61	0.68	0.1222	0.1966	0.2584
	$g = .4, h = 0$	1.32	3.26	0.3388	0.4456	0.5258
	$g = .6, h = 0$	2.26	10.27	0.6012	0.6988	0.7528
	$g=1, h=0$	6.19	110.94	0.9086	0.9448	0.9602
	$g = 2, h = 1$	1.08	5.50	0.2190	0.2984	0.3610
	$g = 4, h = 1$	2.45	20.30	0.4084	0.4968	0.5590
	$g = .6, h = .1$	4.69	89.80	0.6168	0.6972	0.7444
	$g = .8, h = .1$	9.27	603.61	0.7876	0.8474	0.8766

<span id="page-7-0"></span>**Table 4.** Power rates for the goodness-of-fit test on normality  $(n = 20)$ .

\*Type 1 error rates



<span id="page-8-0"></span>**Table 5.** Power rates for the goodness-of-fit test on normality  $(n = 40)$ .

\*Type 1 error rates



<span id="page-9-0"></span>**Table 6.** Power rates for the goodness-of-fit test on normality  $(n = 80)$ .

\*Type 1 error rates



<span id="page-10-0"></span>**Table 7**. Number of times the g- and h- non-normal power values are equal to or greater than .80 for the fit-statistics (K-S, CvM, and A-D)

Note: --- and \*: PROC UNIVARIATE in SAS does not provide exact p-values for K-S at  $α = .20$ 

## **Results**

#### **g- and h- Non-normal Distributions**

[Table 4](#page-7-0) presents Type I error and power rates for the K-S, CvM, and A-D fitstatistics when sample size was 20. A number of conclusions can be drawn from this table. First, Type I error was controlled for each level of significance. Second, for the non-normal alternatives investigated, the K-S was typically the least powerful procedure, followed by CvM, and the A-D is typically most powerful. Also evident from the data is that for kurtotic data, none of the procedures displayed reasonable power (i.e., >.80). Although for skewed and kurtotic data the fit-statistics were only reasonably powerful for extreme departures from normality. As expected, power to detect non-normal distributions increased with more liberal

levels of significance; we excluded the  $\alpha = 0.20$  values from the tables since the values are naturally larger than those reported for the other significance levels examined.

For moderate sample size case (See [Table 5\)](#page-8-0) the same pattern of results held; however, the fit-statistics had more power to detect non-normal data when sample size was 40. Finally, the same pattern of results occurred for our largest investigated sample size of 80 (See [Table 6\)](#page-9-0). And as expected the power to detect non-normal data increased with the increase in sample size.

To summarize the findings for the g-and-h non-normal distributions examined in this study we provide in [Table 7](#page-10-0) a count of the number of times the power values were equal to or greater than .80 across the simulated conditions. Over the significance levels that can be used with the K-S test (i.e.,  $\alpha$  = .05,.10, and .15) the A-D procedure was most powerful to detect non-normal distributions, followed closely by CvM and then by K-S. Clearly the A-D is most sensitive of the three. Also most evident is that the power to detect non-normal distributions is affected by the level of significance as would be expected. Also evident is that contrary to the warning given by [Schoder, Himmelmann, and](#page-18-3)  [Wilhelm \(2006\)](#page-18-3) researchers can detect non-normal distributions with sample sizes less than 100 (80 in our case).

#### **Contaminated Mixed-Normal Distributions**

The power rates for the contaminated normal distributions for the three fitstatistics, K-S, CvM, and A-D are contained in Tables [8,](#page-12-0) [9,](#page-12-1) and [10,](#page-13-0) respectively. As we found for the g- and- h non-normal data, the A-D fit-statistic was most powerful for detecting normal data with outlying values than both the CvM and K-S fit-statistics. And, as expected, power increased with sample size and level of significance. Indeed, to a large extent the reported power values are in reasonably close proximity to .80 for most of the contaminated normal distributions examined. Furthermore, again, as expected the power values were largest when the level of significance was  $> .05$ .

#### **Likert Non-normal data**

The final type of non-normal data that we investigated was data that is obtained when five-point Likert scales are used in measuring the dependent variable. Subjects in the investigations indicate their preference, liking, attitude, etc. on five point type scales (e.g., very unfavorable, unfavorable, neutral, pleasant, very pleasant). Such responses obviously cannot be normally distributed.



<span id="page-12-0"></span>**Table 8.** Power of the Kolmogorov-Smirnov goodness-of-fit test on normality of data for contaminated normal distributions

<span id="page-12-1"></span>**Table 9.** Power of the Cramer-von Mises goodness-of-fit test on normality of data for contaminated normal distributions





**Table 9, continued.** Power of the Cramer-von Mises goodness-of-fit test on normality of data for contaminated normal distributions

<span id="page-13-0"></span>**Table 10.** Power of the Anderson-Darling goodness-of-fit test on normality of data for contaminated normal distributions





<span id="page-14-0"></span>











\*PROC UNIVARIATE does not allow α = .20 for the Kolmogorov-Smirnov test.<br><sup>b</sup>The power values for non-tabled n = 40 values are all 1.000

[Table 11](#page-14-0) provides power rates for the three fit-statistics for detecting nonnormality arising from using a Likert scale for assessing the dependent variable. Preliminary findings indicated that power values were 100% for sample sizes greater than 20 in the vast majority of cases. Thus, it was decided to include a smaller sample size case (i.e.,  $n = 10$ ) to examine power values for a relatively modest number of subjects. The findings are quite positive; that is, in just about every case examined, the power to detect non-normality is  $> 0.80$ . Indeed, out of the 106 tabled values 83 are greater in value than .80. Once again, the A-D statistic provides the best power values, followed by CvM, and then by K-S.

## **Discussion**

Applied researchers use statistical tests to assess whether or not the effect of an experimental manipulation is significant. Unfortunately, the results of many of these investigations are suspect as they often involve the use of statistical procedures with questionable validity. In these cases, the reported effects may be misleading or, in many cases, wrong. Clearly, such erroneous decisions can have serious negative consequences for both the advancement of knowledge in a given field as well as the effective translation of research results into practice. The intent of this paper was to examine whether one can effectively test whether one's data confirms to the validity assumption of normality—a requirement for most classical test statistics. Prior research suggested that one could not use the Kolmogorov-Smirnov goodness-of-fit test to effectively test whether data were normally distributed or not (See e.g., [Schoder, Himmelmann, and Wilhelm, 2006\)](#page-18-3).

We looked into this negative finding by also investigating other fit statistics, the Cramer von Mises and Anderson-Darling tests (See [Muller & Fetterman, 2002](#page-17-7) Chapter 7), varying the skewness and kurtosis values of numerous g-and hdistributions, examining a number of contaminated mixed-normal distributions and examining results when the dependent variable was obtained from nonnormal five-point Likert data. We also manipulated sample sizes  $(n = 20, 40, 80)$ and the level of significance for the test of normality  $\alpha = .05, .10, .15$  and .20).

Of the three fit-statistics we found that the Anderson-Darling procedure was most effective in detecting non-normality being superior to both the Kolmogorov-Smirnov and Cramer-von Mises tests. We also determined that one could reasonably detect non-normality with reasonable sample sizes  $(n = 10,20,40)$ , unlike what was reported by [Schoder, Himmelmann, and Wilhelm \(2006\).](#page-18-3) Lastly, and importantly, since in this context one would want to increase the power to detect effects and concomitantly reduce the probability of falsely accepting the null hypothesis that data are normally distributed, we suggest that preliminary testing be performed with significance levels larger than .05, say  $\alpha = .15$  or  $\alpha = .20$ .

We conclude by reminding researchers that if normality is not present in the data current analytic practices allow researchers to test hypotheses say about mean equality in multiple group designs with software that does not require that data be normally distributed (See e. g., SAS's Glimmix procedure). Or, researchers can choose to replace classical test statistics and their least squares estimators for the mean and variance with robust test statistics with robust estimators (i.e., trimmed means and Winsorized variances (See e.g., [Wilcox,](#page-18-1)  [2012a,](#page-18-1) [b;](#page-18-2) [Wilcox & Keselman, 2003\)](#page-18-0), procedures that have been found to be robust to non-normality [e.g., Erceg-Hurn, Wilcox,  $\&$  Keselman (2013); [Keselman, Algina, Lix, Wilcox, & Deering \(2008a,](#page-17-10) [b\)](#page-17-11)].

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